MAT 5030 Chapter 2: R Environment

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Working Directory

```
setwd("c:/Users/Kazuhiko/WSU/Teach/MAT5030-16W/Ch2")
  # (Useful if you read data from a file or
  # output results to a file)
```

After this line, you only have to specify file names (e.g., "File01.txt") to access to the files in the working directory.

```
getwd() # show the current working directory
```

In OSX from the menu bar, you can choose "Misc" to set, get or reset the working directly.

Terminating R

Type "q()" or "quit()" to quit R (at least the Windows version and Mac OSX version ask if you really want to terminate).

Existing Objects

Save Workspace

- By GUI:
 - Windows: From the menu bar, "File" → "Save Workspace".
 - Mac OSX: From the menu bar, "Workspace" → "Save Workspace File".
- By Script: Type "save.image("XXXX.RData")".

Note: The command saves all objects and functions you have created, but it does not save the scripts you entered (cf. "savehistory").

Load Workspace

- By GUI:
 - Windows: From the menu bar, "File" → "Load Workspace".
 - Mac OSX: From the menu bar, "Workspace" → "Load Workspace File".
- By Script: Type "load("XXXX.RData")".

Output to a file

You may want to output large results into a text file.

```
> A <- 1:100
> sink("Output01.txt") # output into the file
> A^2 # output results to "Output01.txt"
> sink() # output on screen
> A[1:10]
 [1] 1 2 3 4 5 6 7 8 9 10
> sink("Output01.txt") # overwrite results (A^2 will be eliminated)
> A^3
> sink("Output01.txt",append=T) # append new results
> 2*A
```

Input from a file

You may want to load your data, functions and environment from a file with one command "source".

Suppose the source file "S1.txt" is as follows:

```
A <- 1:10
B <- diag(c(4,2))
```

```
> source("S1.txt") # load the code in "S1.txt"
> A
  [1] 1 2 3 4 5 6 7 8 9 10
> source("S1.txt", echo=T) # this command shows the code onscreen
> A <- 1:10
> B <- diag(c(4,2))</pre>
```

History (Windows only)

The letters you entered in R are called history.

Note:

"loadhistory" only loads history, and does not run the codes in the file.

A Summary of File Management

Import:

- Data: 'read.table' etc.
- Run a code: 'source'
- See a code: 'loadhistory'
- State of the R session (R objects): 'load', "Load Workspace".

Export:

- Data: 'sink', 'write.table' etc.
- Save a code (as text): 'savehistory' (Windows only).
- State of the R session (R objects): 'save.image', "Save Workspace".

Library

R has many add-on packages (sometimes called *libraries*) which include procedures for specific statistical analysis.

To use a package,

- install the package (first time only):
 - ▶ By GUI: In the menu bar, "packages" → "Install packages", then follow the instruction. R downloads the package via the Internet or using a local file.
 - By script: Type "options(CRAN="http://cran.r-project.org")" then "install.packages("Rcmdr")" ("Rcmdr" is the name of the package).
- Ioad the package (everytime you started R):
 - By GUI: In the menu bar, "packages" → "Load packages".
 - By script: Type "library(Rcmdr)".

Example: library e1071

The base R package does not have functions to calculate skewness and kurtosis. Several packages have such functions.

```
> X <- rnorm(100) # 100 random numbers following standard normal
> skewness(X) # no such a fucntion in the base package
Error: could not find function "skewness"
```

- > install.packages("e1071") # install the package from the Internet
 >library(e1071) # load the package in your R session
- > skewness(X) # skewness of X
 [1] 0.1657817

skewness =
$$\frac{1}{n} \sum_{i=1}^{100} (x_i - \bar{x})^3 / s^3$$

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To remove package, type

```
> detach("package:Rcmdr")
```

To see what packages are loaded, use "search":

Dataframe: "attach" and "detach"

The "attach" makes it possible to use variables in a dataframe without specifying the dataframe name.

```
> D
 Chase Citi Amex
   3000 4000 5000
2 20000 10000 8000
> D$Chase
     3000 20000
         # R does not recognize where is "Chase".
Error: object "Chase" not found
> attach(D)
> Chase
[1] 3000 20000
> detach(D) # cancel "attach(D)"
```

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Dataframe: "subset" and "transform"

```
> D
         Chase Citi Amex
Andy
          3000
               4000 5000
Rob
         20000 10000 8000
Chris
        4000
               9000 6400
         4000 10000 1500
Dan
> D2 <- subset(D, Amex > 6000) # Make a subset of D
> D2
     Chase Citi Amex
Bob
     20000
           10000 8000
           9000 6400
Chris 4000
> transform(D2, Total = Chase + Citi + Amex) # add a column: Total
     Chase Citi Amex Total
Bob
     20000 10000 8000 38000
Chris 4000
           9000 6400 19400
```

An alternative way to add a column:

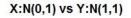
There are two tools to customize graphics:

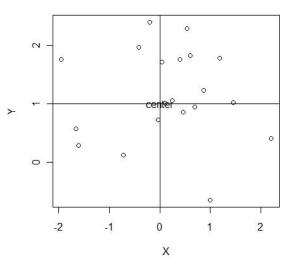
- Overlay simple objects such as axes, texts, legends one by one.
- Change the graphic parameters by using the "par" command.

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Example 1: One by one

```
> plot(X,Y, axes = F) # plot only, no axes
> axis(1, at = c(-2,-1,0,1,2)) # label for X-axis
> axis(2, at = c(-1, 0,1,2,3)) # label for Y-axis
> abline(h = 1, v = 0) # axes (Y=1 and X=0)
> box() # draw an outer frame
> text(0,1,"center") # put the text "center" at (0,1)
> title(main = "X:N(0,1) vs Y:N(1,1)") # title
```

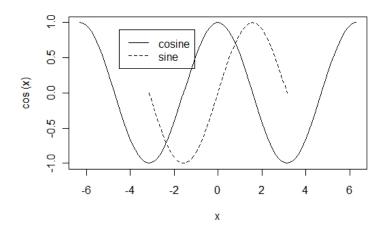




Example 2: Graphic parameters

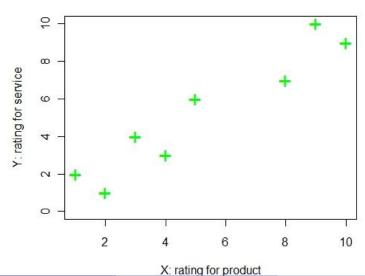
```
> plot(cos, -2*pi, 2*pi, lty=1) # cos curve on [-2pi, 2pi]
> plot(sin, -pi, pi, lty=2, add=T) # sin curve on [-pi,pi]
> # lty: line type, "add=T": overlay
> legend(-4.5,0.9, c("cosine","sine"), lty=1:2)
> # add legend at (-4.5, 0.9)
```

Note: "add=T" does not apply for more complicated plots such as "plot(X,Y)".



Example 3: Graphic parameters

Relationship between X and Y



Graphic parameters: Example 4

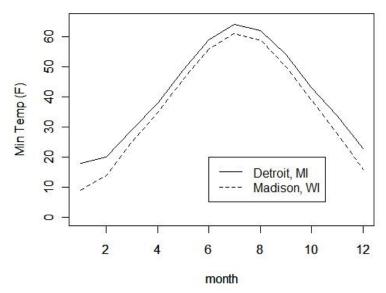
```
TempMI < c(18.20.29.38.49.59.64.62.54.43.34.23)
TempWI < c(9,14,25,35,46,56,61,59,50,39,28,16)
plot(TempMI, ylim=c(0,65), type="1", lty=1,
        xlab="month", ylab="Min Temp (F)")
par(new=T) # overlay figures
plot(TempWI, ylim=c(0,65), type="1", lty=2,
        xlab="month", ylab="Min Temp (F)")
par(new=F) # finish to overlay figures
legend(6, 20, c("Detroit, MI", "Madison, WI"), lty=1:2) # legend at (6,20)
```

Note:

The same range for y-axis (and x-axis) should be used to overlay two plots. To this end point, you may write:

```
> Ylim <- range(TempMI, TempWI) # range = c(Min,Max)
> Ylim
[1] 9 64
```

Then you write "ylim=Ylim" in all "plot()".



Loop: "while" and "repeat"

Suppose we want to find the smallest n such that $2^n > N$, given a positive integer N. How to code?

- Code for N = 123456789.
- Code for general N.

Example 1: while

```
> N <- 123456789
> n <- 1
> while (2^n <= N){ n <- n + 1} # increase n by 1 while "2^n <= N"
> n
[1] 27
> 2^27 # verify the result
[1] 134217728
```

Example 2: repeat & if ~ break

```
> N <- 123456789
> n <- 1
> repeat{ (n <- n + 1)  # repeat until...
+ if (2^n > N) break  # 2^n > N
+ }
> n
[1] 27
```

The "if" statements

Example 1:

When an integer n is given, print "congratulations!" on screen if and only if n is divisible by 9.

```
> n <- 17
> if (n %% 9 == 0){print("congratulations!")}
>
> n <- 18
> if (n %% 9 == 0){print("congratulations!")}
[1] "congratulations!"
```

Example 2:

When a number \boldsymbol{X} is given, let \boldsymbol{Y} be 0 if \boldsymbol{X} is negative and 1 if \boldsymbol{x} is non-negative.

```
> X <- 0.3
> if (X < 0) Y <- 0 else Y <- 1
> Y
[1] 1

> X <- -0.5
> if (X < 0) Y <- 0 else Y <- 1
> Y
[1] 0
```

Example 3: When a number **X** is given, let **Y** be 1 and **Z** be 2 if **X** is between -10 and 10, and **Y** be 0 and **Z** be -2 otherwise.

```
> X < -12
> {
          if ((X >= -10) & (X <= 10)){
+
                  Y <- 1
                  Z < -2
                   else
+
                  \{Y < -0\}
+
                  Z < -2
+
+ }
> c(Y,Z)
Γ1 0 -2
```

Defining a function

We want to develop a **function** "binarylength" which returns n when we input N.

```
> binarylength <- function(N){  # define function "binarylength"
+ n <- 1
+  while (2^n <= N){ n <- n + 1}
+  n
+ }
> binarylength(123456789)
[1] 27
```

Note: "+" appears in R when a function is defined using more than 1 line.

We can create a function with multiple inputs and/or outputs. Suppose we want to create a function to find n such that $q^n > N$ when positive integers $q \ge 2$ and N are given. We want to output n and q^n .

We can set default values for a function.

Loop: "for"

Suppose we want to create a vector $(x_2 - x_1, x_3 - x_2, \dots, x_n - x_{n-1})$ when $V = (x_1, \dots, x_n)$ is given.

Unlike the examples for while and repeat, We always have to iterate calculation for a fixed number of times (i.e., n - 1 times).

Example 1: Code using "for"

```
> dV <- function(V) {
+ n <- length(V)  # n = dimension of V
+ W <- numeric(n-1) # (n-1)-dim vector
+ for (i in 1:(n-1)) {  # loop n is from 1 to (n-1)
+ W[i] <- V[i+1] - V[i]  # define i-th element of W
+ }
+ W
+ }
> X <- c(1,3,6,10,15)
> dV(X)
[1] 2 3 4 5
```

Example 2: Code without using "for"

Loop functions are slower than matrix algebra in R. (This is common among statistical languages, and uncommon among fundamental languages like C). Avoid loops if possible.

A function to overlay plot

It is cumbersome to overlay two X-Y plots. Create a function to do it at once.

Input:

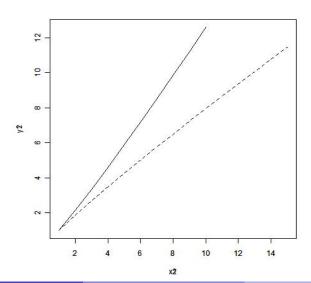
- x1: x-coordinates for the 1st data
- y1: y-coordinates for the 1st data
- x2: x-coordinates for the 2nd data
- y2" y-coordinates for the 2nd data

Output:

Save the figure as "G1.jpg".

Function:

```
jpgout2 \leftarrow function(x1, y1, x2, y2, ...)
        Xlim <- range(x1, x2)</pre>
        Ylim <- range(y1, y2)
        ipeg("G1.jpg")
        plot(x1, y1, xlim= Xlim, ylim= Ylim, type = "l", lty=1, ...)
        par(new=T)
        plot(x2, y2, xlim= Xlim, ylim= Ylim, type = "l", lty=2, ...)
        par(new=F)
        dev.off()
Sample:
> X1 < -1.10
> Y1 < - X1^1.1 \# Y = X^1.1 \text{ on } [1,10]
> X2 <- 1:15
> Y2 <- X2^0.9 \# Y = X^0.9 \text{ on } [1,15]
> jpgout2(X1, Y1, X2, Y2)
windows
      2
```



Text files

Text file: "Credit.txt"

Chase	Citi	Amex
3000	4000	5000
20000	10000	8000
4000	9000	6400
4000	10000	1500
5000	3000	5500

To save a spread sheet as a text file, you can cut and paste data in a text editor or save as a tab delimited text file.

Clipboard

You can copy the data into the clipboard ("Ctrl + C" in Windows; "Command + C" in Mac OSX), then input it into R by:

```
Windows:
```

```
> read.table("clipboard", header=T)
Mac OSX:
> read.table(pipe("pbpaste"), header=T)
Warning message:
In read.table(pipe("pbpaste"), header = T) :
   incomplete final line found by readTableHeader on 'pbpaste'
```

MS Excel

It is recommended to load data as a text or csv file, but there are a few options to load data from Microsoft Excel.

library(xlsx):

- Install and load the package "xlsx".
 - The package "xlsxjars" is also required.
 - Java is also required. The R asks you to install the Java when you type 'library(xlsx)'.
- Use the 'read.xlsx' or 'read.xlsx2' function to import the data. The latter is faster for the data with many rows.

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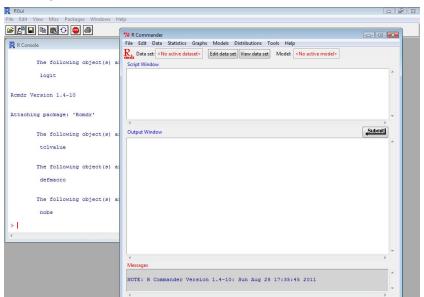
MS Excel: Sample Code

```
> read.xlsx("Data1.xls",1, header=T) # 1: 1st sheet
 A B C
1 6 13 7
2 8 8 10
3 7 10 9
4 8 9
       9
> read.xlsx2("Data1.xls",1, header=T)
> read.xlsx2("Data1.xlsx",sheetName="GHDay", header=T)
       Weight Height Color
    7.4
           23 dark
    8.9 26 light
 6.5 19 dark
    7.2 24 light
    5.9 18 dark
```

library(Rcmdr):

- Install and load the pacakage "Rcmdr" (R commander).
- From the menu bar in R commander, choose "Data" → "Import Data"
 → "from Excel, ...".

Note: The R commander is a graphical user interface similar to more basic statistical softwares such as Minitab. You can do various statistical analysis using this interface.



Some other formats

Some other formats such as SAS, SPSS, Minitab data files can be loaded by the "foreign" package.

The R commander also has a interface to load such datasets.