

ISU

MAT 356 R Tutorial, Spring 2004



0. R Basics

0.1. What is R?

R is a software package especially suitable for data analysis and graphical representation. Functions and results of analysis are all stored as objects, allowing easy function modification and model building. R provides the language, tool, and environment in one convenient package.

It is very flexible and highly customizable. Excellent graphical tools make R an ideal environment for EDA (Exploratory Data Analysis). Since most high level functions are written in R language itself, you can learn the language by studying the function code.

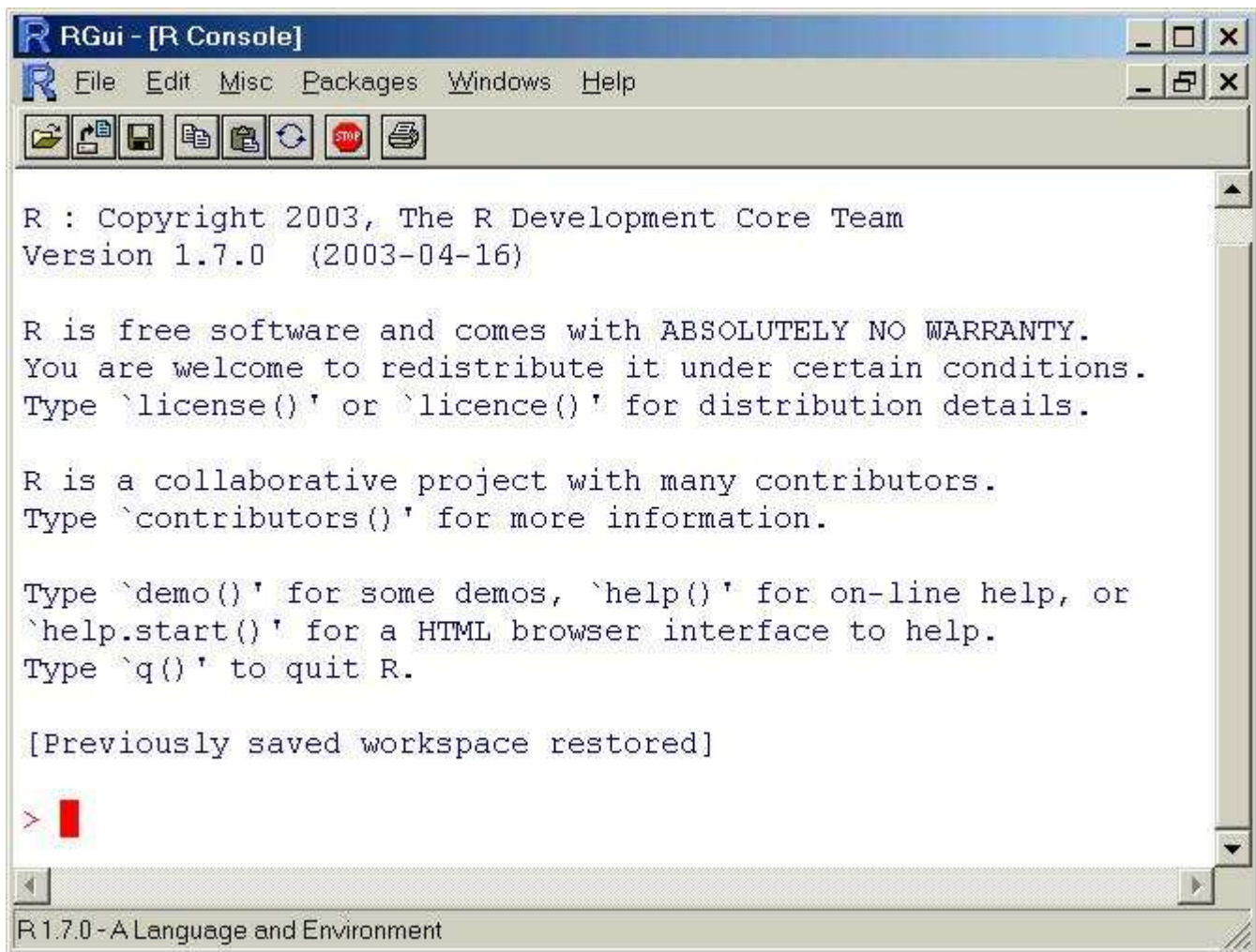
On the other hand, R has a few weaknesses. For example, R is not particularly efficient in handling large data sets. Also, it is rather slow in executing a large number of for loops, compared to compiler languages such as C/C++. Learning curve is somewhat steep compared to "point and click" software.

0.2 Where do I get R?

There are versions for Unix, Windows, and Macintosh. All of them are free, and Windows version is downloadable at:

<http://cran.us.r-project.org/bin/windows>

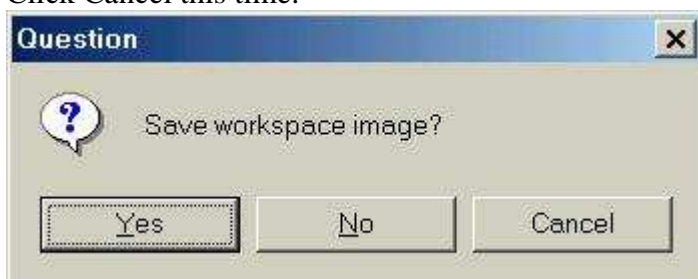
and follow the download instructions.



0.3 Invoking R

If properly installed, usually R has a shortcut icon on the desktop screen and/or you can find it under Start|Programs|R menu. If not, search and run the executable file rgui.exe by double clicking from the search result window.

To quit R, type `q()` at the R prompt (`>`) and press Enter key. A dialog box will ask whether to save the objects you have created during the session so that they will become available next time you invoke R. Click Cancel this time.



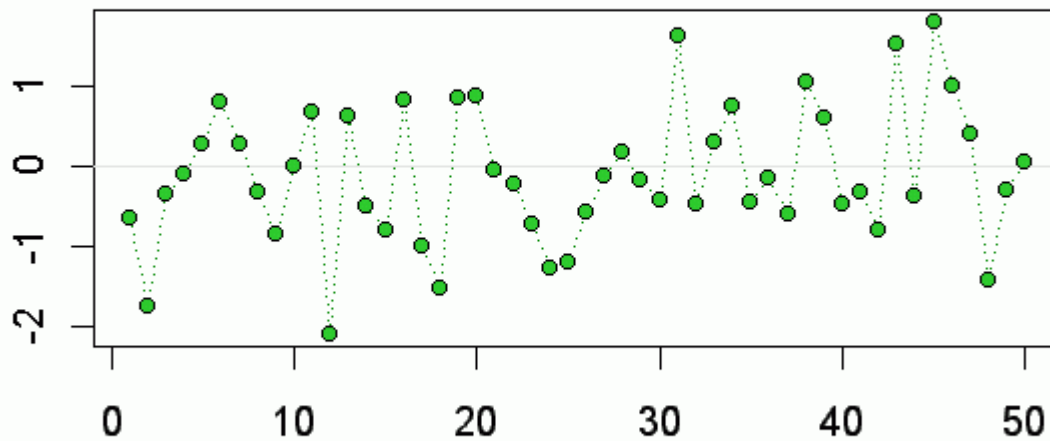
Commands you entered can be easily recalled and modified. Just by hitting the arrow keys in the keyboard, you can navigate through the recently entered commands.

```
> objects() # list the names of all objects
> rm(data1) #remove the object named data1 from the current environment
```

1. Graphics: a few examples

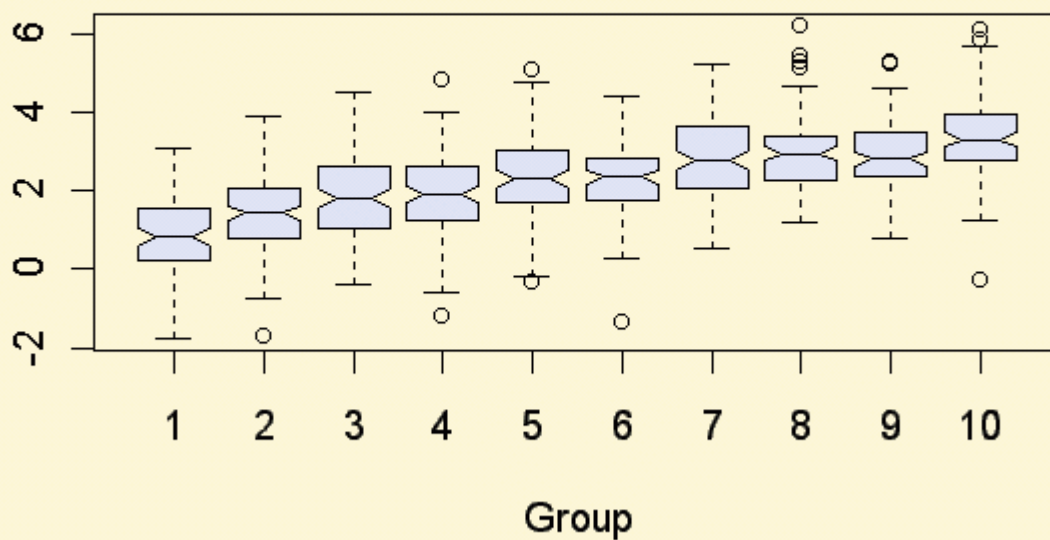
In addition to standard plots such as histogram, bar charts, pie charts and so forth, R provides an impressive array of graphical tools. The following series of plots shows a few of the extensive graphical capabilities of R.

Simple Use of Color In a Plot

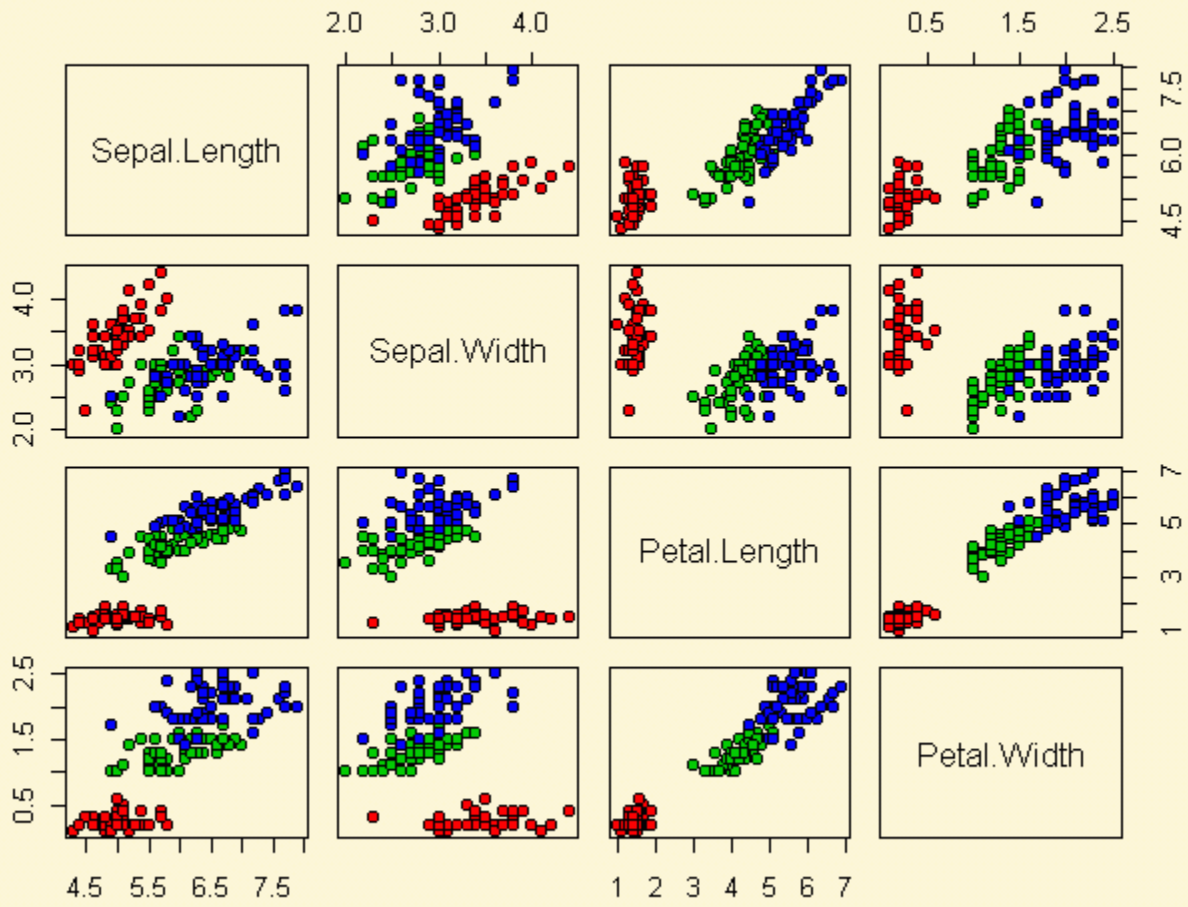


Just a Whisper of a Label

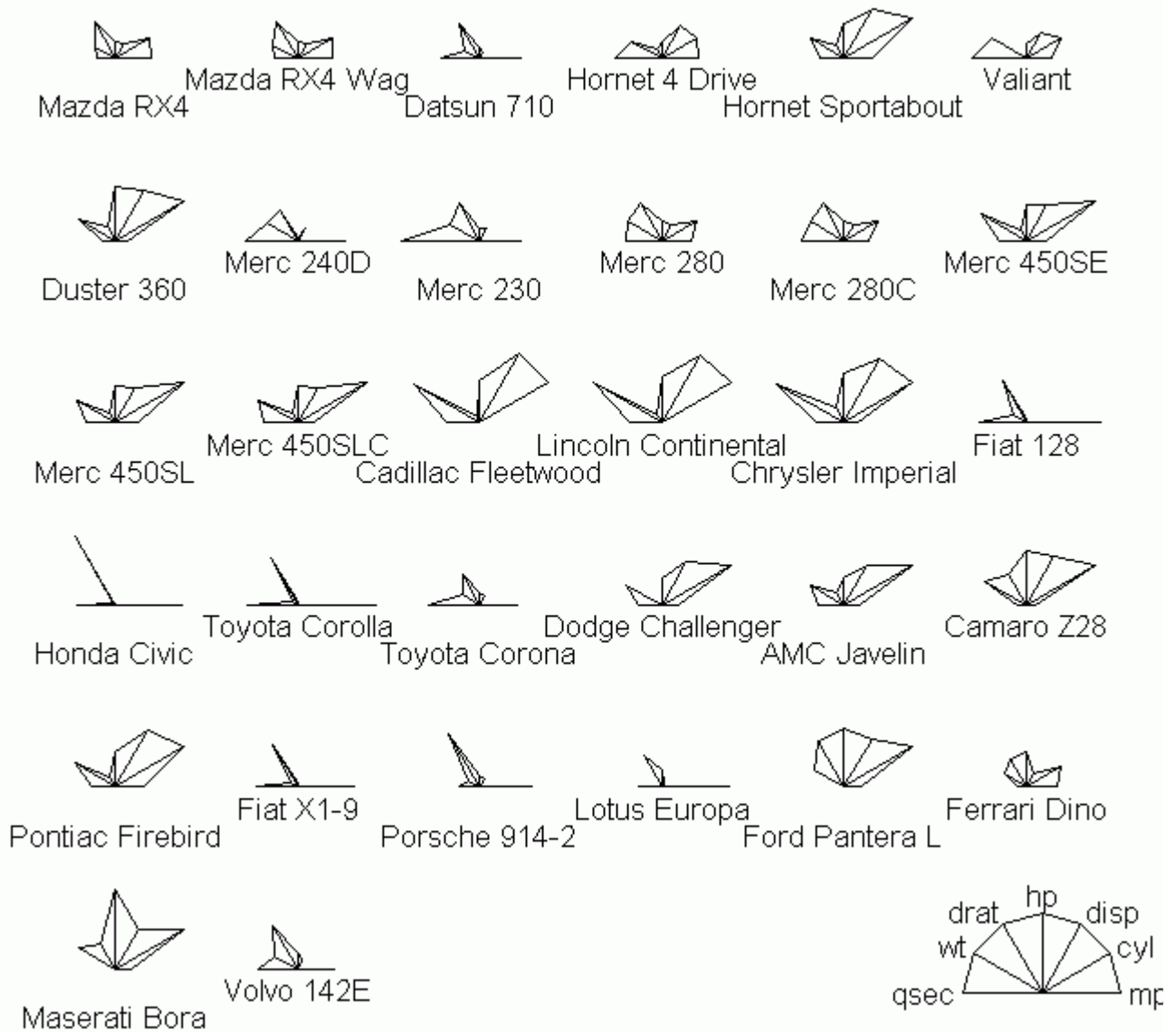
Notched Boxplots



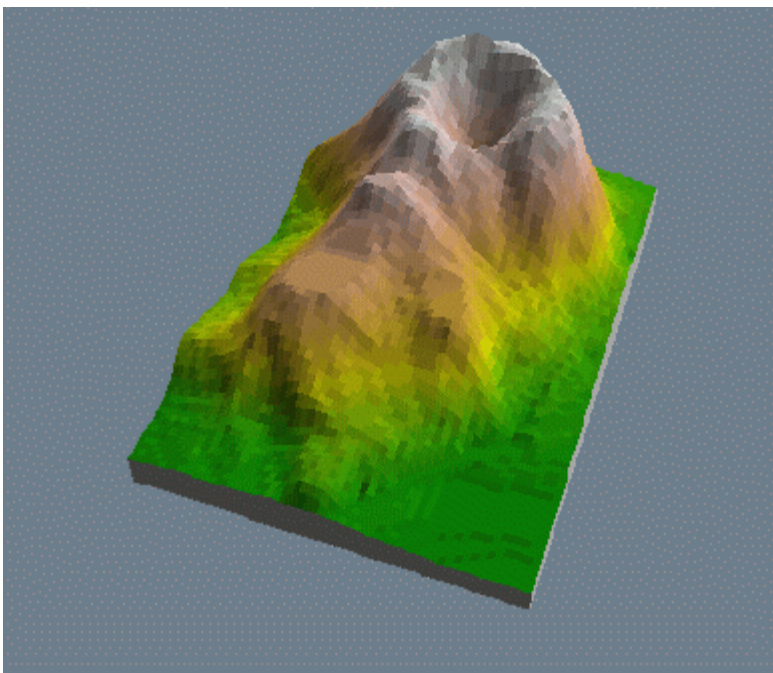
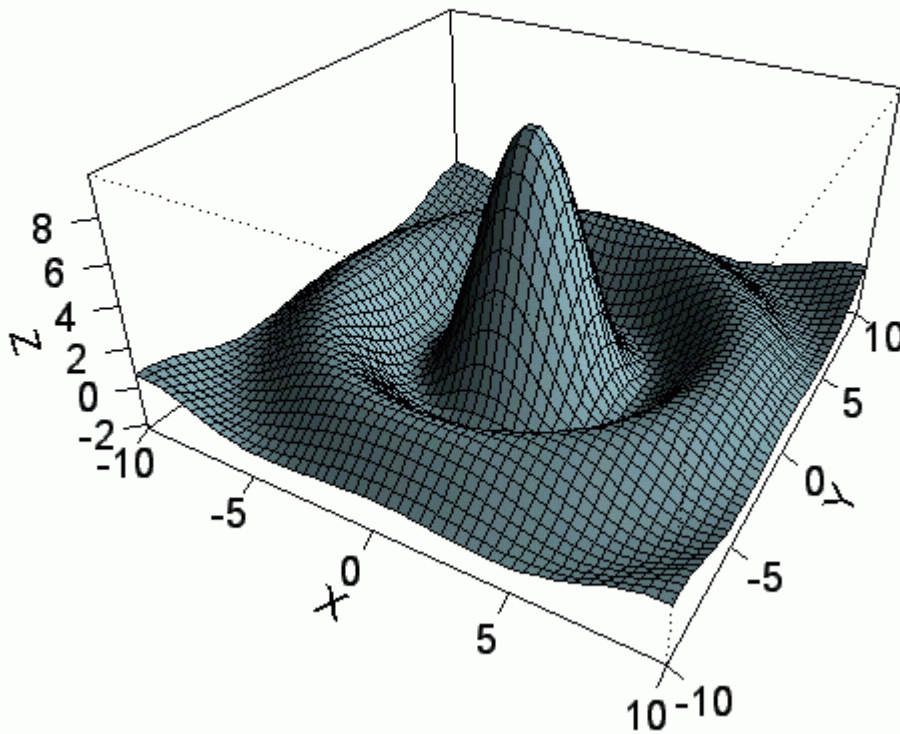
Edgar Anderson's Iris Data



Motor Trend Cars : stars(*, full = F)

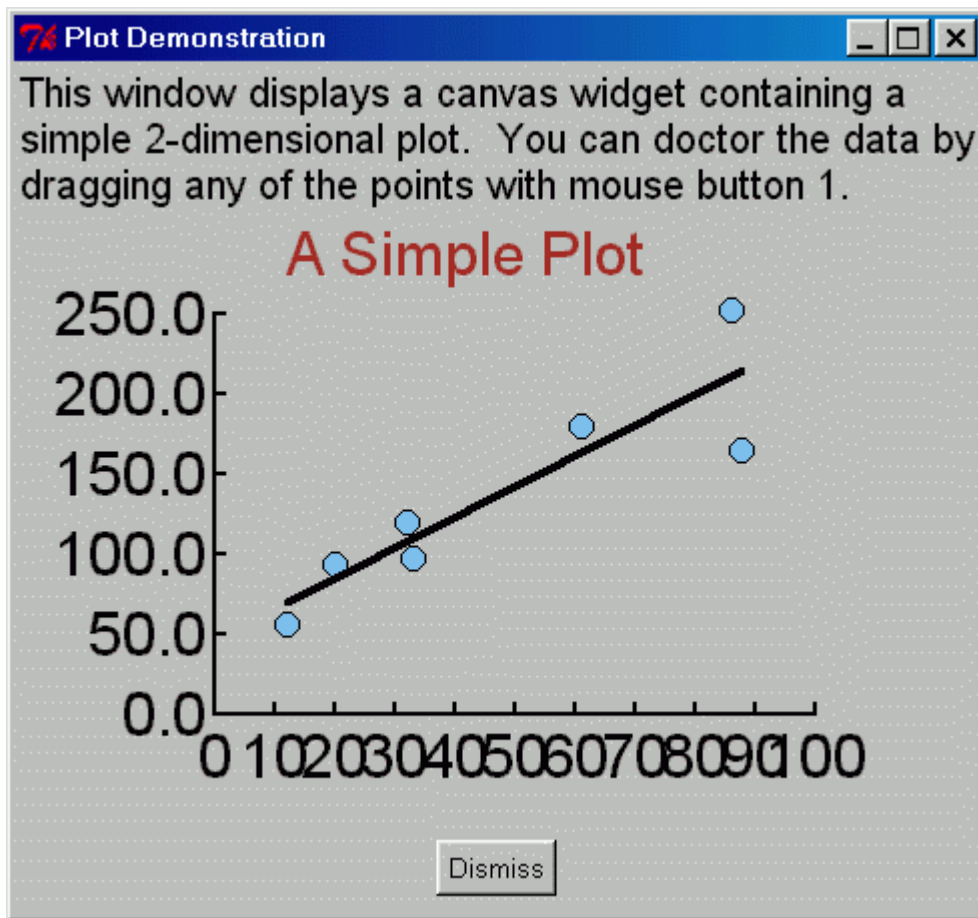


$$z = \text{Sinc}(\sqrt{x^2 + y^2})$$



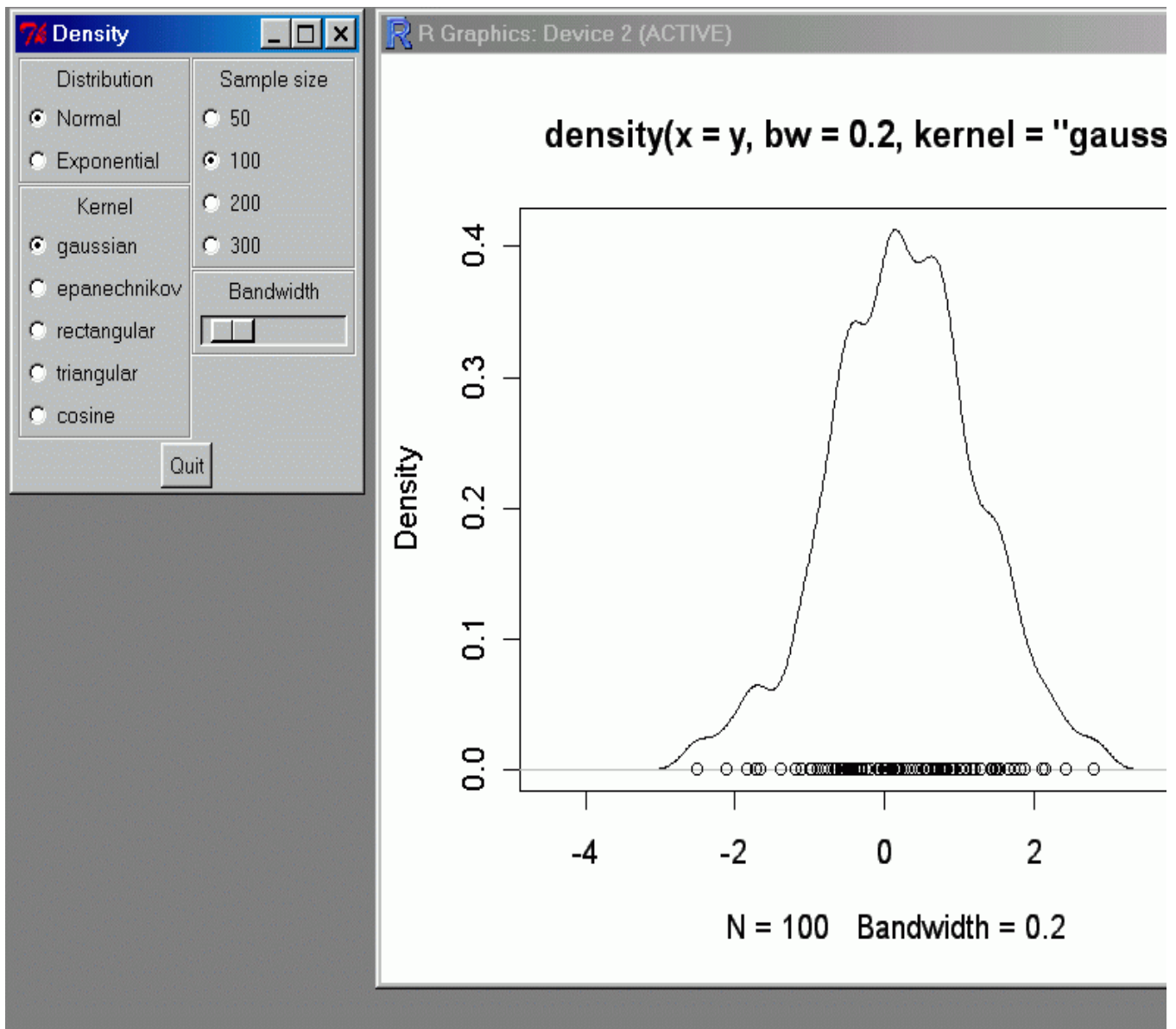
Interactive graphics can serve as a great learning tool. Students can quickly grasp the role of outliers and influential points in a simple linear regression by the following example.

```
> library(tcltk)
> demo(tkcanvas)
```



Effect of kernel choice, sample size and bandwidth can be conveniently illustrated by the following demonstration:

```
> library(tcltk)
> demo(tkdensity)
```



2. Basic Operations

2.1 Computation

First of all, R can be used as an ordinary calculator. There are a few examples:

```
> 2 + 3 * 5      # Note the order of operations.
> log (10)      # Natural logarithm with base e=2.718282
> 4^2           # 4 raised to the second power
> 3/2           # Division
> sqrt (16)     # Square root
> abs (3-7)     # Absolute value of 3-7
> pi            # The mysterious number
> exp(2)        # exponential function
> 15 %/% 4      # This is the integer divide operation
> # This is a comment line
```

Assignment operator (`<-`) stores the value (object) on the right side of (`<-`) expression in the left side. Once assigned, the object can be used just as an ordinary component of the computation. To find out what the object looks like, simply type its name. Note that R is case sensitive, e.g., object names `abc`, `ABC`, `Abc` are all different.

```
> x<- log(2.843432) *pi
> x
[1] 3.283001
> sqrt(x)
[1] 1.811905
> floor(x)          # largest integer less than or equal to x (Gauss number)
[1] 3
> ceiling(x)       # smallest integer greater than or equal to x
[1] 4
```

R can handle complex numbers, too.

```
> x<-3+2i
> Re(x)          # Real part of the complex number x
[1] 3
> Im(x)          # Imaginary part of x
[1] 2
> y<- -1+1i
> x+y
[1] 2+3i
> x*y
[1] -5+1i
```

Important note: since there are many built-in functions in R, make sure that the new object names you assign are not already used by the system. A simple way of checking this is to type in the name you want to use. If the system returns an error message telling you that such object is not found, it is safe to use the name. For example, `c` (for concatenate) is a built-in function used to combine elements so NEVER assign an object to `c`!

2.2 Vector

R handles vector objects quite easily and intuitively.

```
> x<-c(1,3,2,10,5)    #create a vector x with 5 components
> x
[1] 1 3 2 10 5
> y<-1:5              #create a vector of consecutive integers
> y
[1] 1 2 3 4 5
> y+2                 #scalar addition
[1] 3 4 5 6 7
> 2*y                 #scalar multiplication
[1] 2 4 6 8 10
> y^2                 #raise each component to the second power
[1] 1 4 9 16 25
> 2^y                 #raise 2 to the first through fifth power
```

```
[1] 2 4 8 16 32
> y #y itself has not been unchanged
[1] 1 2 3 4 5
> y<-y*2
> y #it is now changed
[1] 2 4 6 8 10
```

More examples of vector arithmetic:

```
> x<-c(1,3,2,10,5); y<-1:5 #two or more statements are separated by semicolons
> x+y
[1] 2 5 5 14 10
> x*y
[1] 1 6 6 40 25
> x/y
[1] 1.0000000 1.5000000 0.6666667 2.5000000 1.0000000
> x^y
[1] 1 9 8 10000 3125

> sum(x) #sum of elements in x
[1] 21
> cumsum(x) #cumulative sum vector
[1] 1 4 6 16 21
> diff(x) # first difference
[1] 2 -1 8 -5
> diff(x,2) #second difference
[1] 1 7 3
> max(x) #maximum
[1] 10
> min(x) #minimum
[1] 1
```

Sorting can be done using `sort()` command:

```
> x
[1] 1 3 2 10 5
> sort(x) # increasing order
[1] 1 2 3 5 10
> sort(x, decreasing=T) # decreasing order
[1] 10 5 3 2 1
```

Component extraction is a very important part of vector calculation.

```
> x
[1] 1 3 2 10 5
> length(x) # number of elements in x
[1] 5
> x[3] # the third element of x
[1] 2
> x[3:5] # the third to fifth element of x, inclusive
[1] 2 10 5
> x[-2] # all except the second element
```

```
[1] 1 2 10 5
> x[x>3]          # list of elements in x greater than 3
[1] 10 5
```

Logical vector can be handy:

```
> x>3
[1] FALSE FALSE FALSE TRUE TRUE
> as.numeric(x>3) # as.numeric() function coerces logical components to numeric
[1] 0 0 0 1 1
> sum(x>3)        # number of elements in x greater than 3
[1] 2
> (1:length(x))[x<=2] # indices of x whose components are less than or equal to 2
[1] 1 3
> z<-as.logical(c(1,0,0,1)) # numeric to logical vector conversion
> z
[1] TRUE FALSE FALSE TRUE
```

Character vector:

```
> colors<-c("green", "blue", "orange", "yellow", "red")
> colors
[1] "green" "blue" "orange" "yellow" "red"
```

Individual components can be named and referenced by their names.

```
> names(x)          # check if any names are attached to x
NULL
> names(x)<-colors  # assign the names using the character vector colors
> names(x)
[1] "green" "blue" "orange" "yellow" "red"
> x
  green  blue orange yellow  red
    1    3    2    10    5
> x["green"]       # component reference by its name
green
  1
> names(x)<-NULL    # names can be removed by assigning NULL
> x
[1] 1 3 2 10 5
```

`seq()` and `rep()` provide convenient ways to a construct vectors with a certain pattern.

```
> seq(10)
[1] 1 2 3 4 5 6 7 8 9 10
> seq(0,1,length=10)
[1] 0.0000000 0.1111111 0.2222222 0.3333333 0.4444444 0.5555556 0.6666667
[8] 0.7777778 0.8888889 1.0000000
> seq(0,1,by=0.1)
[1] 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0
> rep(1,3)
[1] 1 1 1
> c(rep(1,3),rep(2,2),rep(-1,4))
```

```
[1] 1 1 1 2 2 -1 -1 -1 -1
> rep("Small",3)
[1] "Small" "Small" "Small"
> c(rep("Small",3),rep("Medium",4))
[1] "Small" "Small" "Small" "Medium" "Medium" "Medium" "Medium"
> rep(c("Low","High"),3)
[1] "Low" "High" "Low" "High" "Low" "High"
```

2.3 Matrices

A matrix refers to a numeric array of rows and columns. One of the easiest ways to create a matrix is to combine vectors of equal length using `cbind()`, meaning "column bind":

```
> x
[1] 1 3 2 10 5
> y
[1] 1 2 3 4 5
> m1<-cbind(x,y);m1
      x y
[1,] 1 1
[2,] 3 2
[3,] 2 3
[4,] 10 4
[5,] 5 5
> t(m1)          # transpose of m1
  [,1] [,2] [,3] [,4] [,5]
x     1     3     2    10     5
y     1     2     3     4     5

> m1<-t(cbind(x,y))  # Or you can combine them and assign in one step
> dim(m1)            # 2 by 5 matrix
[1] 2 5
> m1<-rbind(x,y)     # rbind() is for row bind and equivalent to t(cbind()).
```

Of course you can directly list the elements and specify the matrix:

```
> m2<-matrix(c(1,3,2,5,-1,2,2,3,9),nrow=3);m2
  [,1] [,2] [,3]
[1,]  1   5   2
[2,]  3  -1   3
[3,]  2   2   9
```

Note that the elements are used to fill the first column, then the second column and so on. To fill row-wise, we specify `byrow=T` option:

```
> m2<-matrix(c(1,3,2,5,-1,2,2,3,9),ncol=3,byrow=T);m2
  [,1] [,2] [,3]
[1,]  1   3   2
[2,]  5  -1   2
[3,]  2   3   9
```

Extracting the component of a matrix involves one or two indices.

```
> m2
```

```

      [,1] [,2] [,3]
[1,]    1    3    2
[2,]    5   -1    2
[3,]    2    3    9
> m2[2,3]           #element of m2 at the second row, third column
[1] 2
> m2[2,]           #second row
[1] 5 -1 2
> m2[,3]           #third column
[1] 2 2 9
> m2[-1,]          #submatrix of m2 without the first row
      [,1] [,2] [,3]
[1,]    5   -1    2
[2,]    2    3    9
> m2[,-1]          #ditto, sans the first column
      [,1] [,2]
[1,]    3    2
[2,]   -1    2
[3,]    3    9
> m2[-1,-1]        #submatrix of m2 with the first row and column removed
      [,1] [,2]
[1,]   -1    2
[2,]    3    9

```

Matrix computation is usually done component-wise.

```

> m1<-matrix(1:4, ncol=2); m2<-matrix(c(10,20,30,40),ncol=2)
> 2*m1              # scalar multiplication
      [,1] [,2]
[1,]    2    6
[2,]    4    8
> m1+m2             # matrix addition
      [,1] [,2]
[1,]   11   33
[2,]   22   44
> m1*m2             # component-wise multiplication
      [,1] [,2]
[1,]   10   90
[2,]   40  160

```

Note that $m1*m2$ is NOT the usual matrix multiplication. To do the matrix multiplication, you should use `%*%` operator instead.

```

> m1 %*% m2
      [,1] [,2]
[1,]   70  150
[2,]  100  220

> solve(m1)         #inverse matrix of m1
      [,1] [,2]

```

```

[1,]  -2  1.5
[2,]   1 -0.5
> solve(m1)%*%m1      #check if it is so
      [,1] [,2]
[1,]    1    0
[2,]    0    1
> diag(3)             #diag() is used to construct a k by k identity matrix
      [,1] [,2] [,3]
[1,]    1    0    0
[2,]    0    1    0
[3,]    0    0    1
> diag(c(2,3,3))     #as well as other diagonal matrices
      [,1] [,2] [,3]
[1,]    2    0    0
[2,]    0    3    0
[3,]    0    0    3

```

Eigenvalues and eigenvectors of a matrix is handled by `eigen()` function:

```

> eigen(m2)
$values
[1] 53.722813 -3.722813

$vectors
      [,1]      [,2]
[1,] -0.5657675 -0.9093767
[2,] -0.8245648  0.4159736

```

2.4 Finding roots: a simple example

A built-in R function `uniroot()` can be called from a user defined function `root.fun()` to compute the root of a univariate function and plot the graph of the function at the same time.

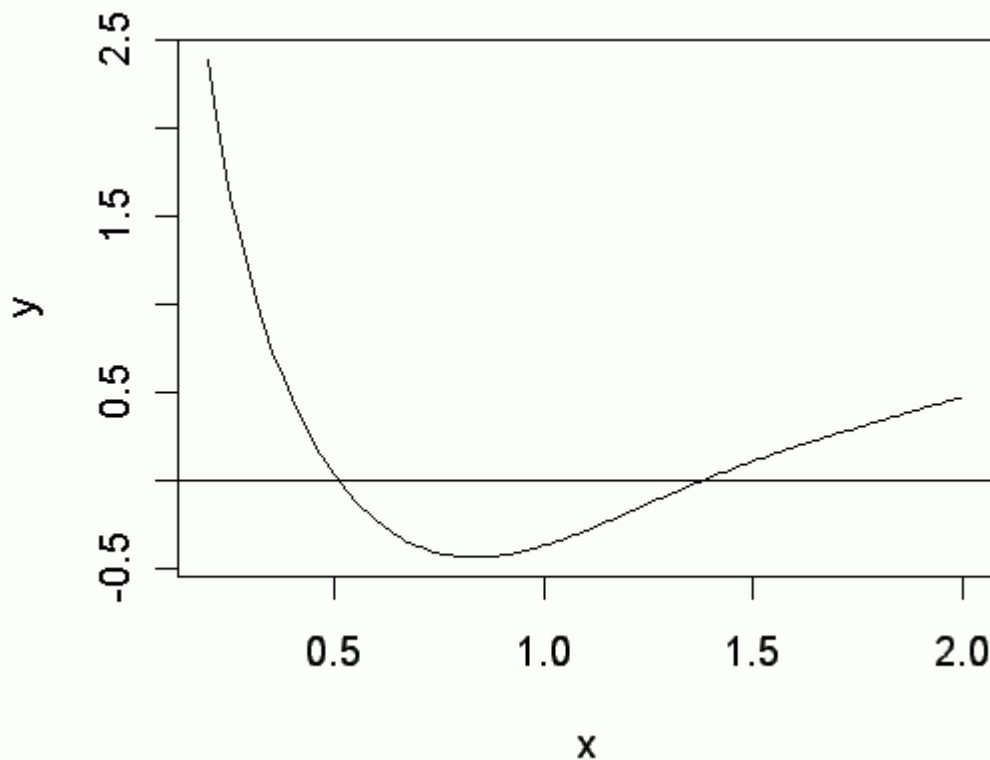
```

> y.fun<-function (x)
{y<-(log(x))^2-x*exp(-x^3)
}

> root.fun<- function ()
{
  x<-seq(0.2,2,0.01)
  y<-y.fun(x)
  win.graph()
  plot(x,y,type="l")
  abline(h=0)
  r1 <- uniroot(y.fun,lower=0.2,upper=1)$root
  r2 <- uniroot(y.fun,lower=1,upper=2)$root
  cat("Roots : ", round(r1,4), " ", round(r2,4), "\n")
}

> root.fun()

```



2.5 Data frame

Data frame is an array consisting of columns of various mode (numeric, character, etc). Small to moderate size data frame can be constructed by `data.frame()` function. For example, we illustrate how to construct a data frame from the car data*:

Make	Model	Cylinder	Weight	Mileage	Type
Honda	Civic	V4	2170	33	Sporty
Chevrolet	Beretta	V4	2655	26	Compact
Ford	Escort	V4	2345	33	Small
Eagle	Summit	V4	2560	33	Small
Volkswagen	Jetta	V4	2330	26	Small
Buick	Le Sabre	V6	3325	23	Large
Mitsubishi	Galant	V4	2745	25	Compact
Dodge	Grand Caravan	V6	3735	18	Van
Chrysler	New Yorker	V6	3450	22	Medium
Acura	Legend	V6	3265	20	Medium

*Source: adapted from a built-in data set `fuel.frame`.

```
> Make<-c("Honda","Chevrolet","Ford","Eagle","Volkswagen","Buick","Mitsbusihi",
+ "Dodge","Chrysler","Acura")
> Model<-c("Civic","Beretta","Escort","Summit","Jetta","Le Sabre","Galant",
+ "Grand Caravan","New Yorker","Legend")
```

Note that the plus sign (+) in the above commands are automatically inserted when the carriage return is pressed without completing the list. Save some typing by using `rep()` command. For example, `rep("V4", 5)` instructs R to repeat V4 five times.

```
> Cylinder<-c(rep("V4", 5), "V6", "V4", rep("V6", 3))
> Cylinder
 [1] "V4" "V4" "V4" "V4" "V4" "V6" "V4" "V6" "V6" "V6"
> Weight<-c(2170, 2655, 2345, 2560, 2330, 3325, 2745, 3735, 3450, 3265)
> Mileage<-c(33, 26, 33, 33, 26, 23, 25, 18, 22, 20)
> Type<-c("Sporty", "Compact", rep("Small", 3), "Large", "Compact", "Van", rep("Medium", 2))
```

Now `data.frame()` function combines the six vectors into a single data frame.

```
> Car<-data.frame(Make, Model, Cylinder, Weight, Mileage, Type)
> Car
      Make      Model Cylinder Weight Mileage  Type
1   Honda     Civic      V4    2170     33 Sporty
2 Chevrolet Beretta      V4    2655     26 Compact
3   Ford     Escort      V4    2345     33  Small
4   Eagle   Summit      V4    2560     33  Small
5 Volkswagen  Jetta      V4    2330     26  Small
6   Buick   Le Sabre      V6    3325     23  Large
7 Mitsubihi Galant      V4    2745     25 Compact
8   Dodge Grand Caravan      V6    3735     18   Van
9 Chrysler New Yorker      V6    3450     22 Medium
10  Acura   Legend      V6    3265     20 Medium

> names(Car)
 [1] "Make"      "Model"     "Cylinder"  "Weight"    "Mileage"   "Type"
```

Just as in matrix objects, partial information can be easily extracted from the data frame:

```
> Car[1,]
      Make Model Cylinder Weight Mileage  Type
1 Honda Civic      V4    2170     33 Sporty
```

In addition, individual columns can be referenced by their labels:

```
> Car$Mileage
 [1] 33 26 33 33 26 23 25 18 22 20
> Car[,5]           #equivalent expression, less informative
> mean(Car$Mileage) #average mileage of the 10 vehicles
 [1] 25.9
> min(Car$Weight)
 [1] 2170
```

`table()` command gives a frequency table:

```
> table(Car$Type)

Compact  Large  Medium  Small  Sporty  Van
      2      1      2      3      1      1
```

If the proportion is desired, type the following command instead:

```
> table(Car$Type)/10
```

```
Compact   Large   Medium   Small   Sporty   Van
    0.2     0.1     0.2     0.3     0.1     0.1
```

Note that the values were divided by 10 because there are that many vehicles in total. If you don't want to count them each time, the following does the trick:

```
> table(Car$Type)/length(Car$Type)
```

Cross tabulation is very easy, too:

```
> table(Car$Make, Car$Type)
```

```

           Compact Large Medium Small Sporty Van
Acura      0      0      1      0      0      0
Buick      0      1      0      0      0      0
Chevrolet  1      0      0      0      0      0
Chrysler   0      0      1      0      0      0
Dodge      0      0      0      0      0      1
Eagle     0      0      0      1      0      0
Ford       0      0      0      1      0      0
Honda      0      0      0      0      1      0
Mitsbusihi 1      0      0      0      0      0
Volkswagen 0      0      0      1      0      0
```

What if you want to arrange the data set by vehicle weight? `order()` gets the job done.

```
> i<-order(Car$Weight);i
```

```
[1] 1 5 3 4 2 7 10 6 9 8
```

```
> Car[i,]
```

```

      Make      Model Cylinder Weight Mileage  Type
1   Honda      Civic         V4  2170     33 Sporty
5 Volkswagen   Jetta         V4  2330     26  Small
3     Ford    Escort         V4  2345     33  Small
4     Eagle   Summit         V4  2560     33  Small
2 Chevrolet   Beretta         V4  2655     26 Compact
7 Mitsbusihi   Galant         V4  2745     25 Compact
10    Acura    Legend         V6  3265     20 Medium
6     Buick   Le Sabre         V6  3325     23  Large
9   Chrysler New Yorker         V6  3450     22 Medium
8     Dodge Grand Caravan         V6  3735     18   Van
```

2.6 Creating/editing data objects

```
> y
```

```
[1] 1 2 3 4 5
```

If you want to modify the data object, use `edit()` function and assign it to an object. For example, the following command opens notepad for editing. After editing is done, choose File | Save and Exit from Notepad.

```
> y<-edit(y)
```

If you prefer entering the data.frame in a spreadsheet style data editor, the following command invokes the built-in editor with an empty spreadsheet.

```
> data1<-edit(data.frame())
```

After entering a few data points, it looks like this:

	var1	var2	var3	var4	var5
1	1	aa	0.234		
2	2	bb	0.432		
3	3	cc	0.32		
4	4	dd	0.275		
5					
6					
7					
8					

You can also change the variable name by clicking once on the cell containing it. Doing so opens a dialog box:

variable name

type numeric character

When finished, click **X** in the upper right corner of the dialog box to return to the Data Editor window. Close the Data Editor to return to the R command window (R Console). Check the result by typing:

```
> data1
```

3. More on R Graphics

Not only R has fancy graphical tools, but also it has all sorts of useful commands that allow users to control almost every aspect of their graphical output to the finest details.

3.1 Histogram

We will use a data set [fuel.frame](#) which is based on makes of cars taken from the April 1990 issue of Consumer Reports.

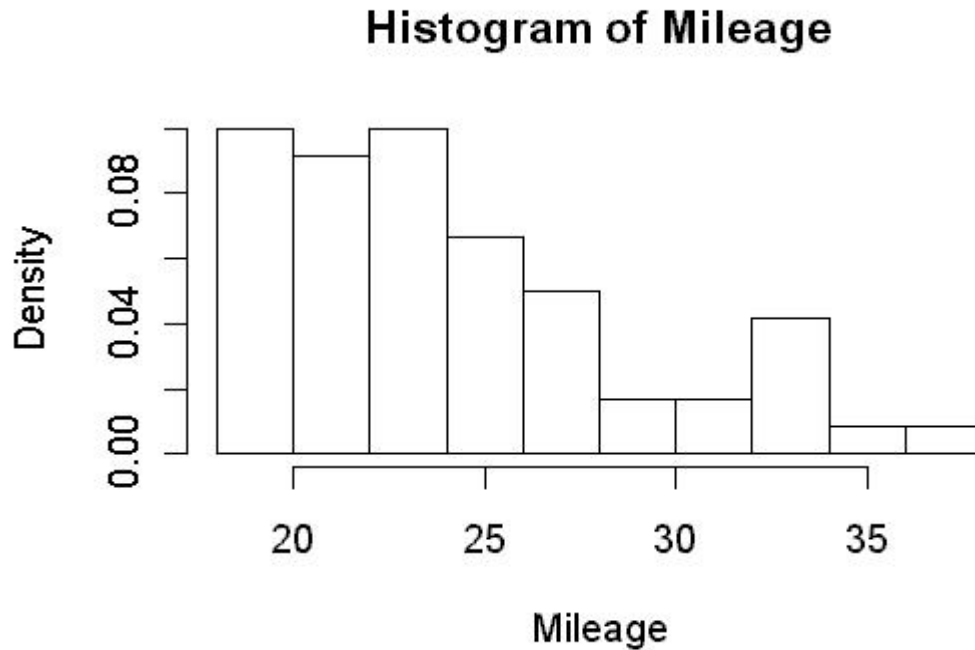
```
> fuel.frame<-read.table("c:/fuel-frame.txt", header=T, sep=",")
> names(fuel.frame)
[1] "row.names" "Weight"      "Disp."      "Mileage"    "Fuel"      "Type"
```

```
> attach(fuel.frame)
```

`attach()` allows to reference variables in `fuel.frame` without the cumbersome `fuel.frame$` prefix.

In general, graphic functions are very flexible and intuitive to use. For example, `hist()` produces a histogram, `boxplot()` does a boxplot, etc.

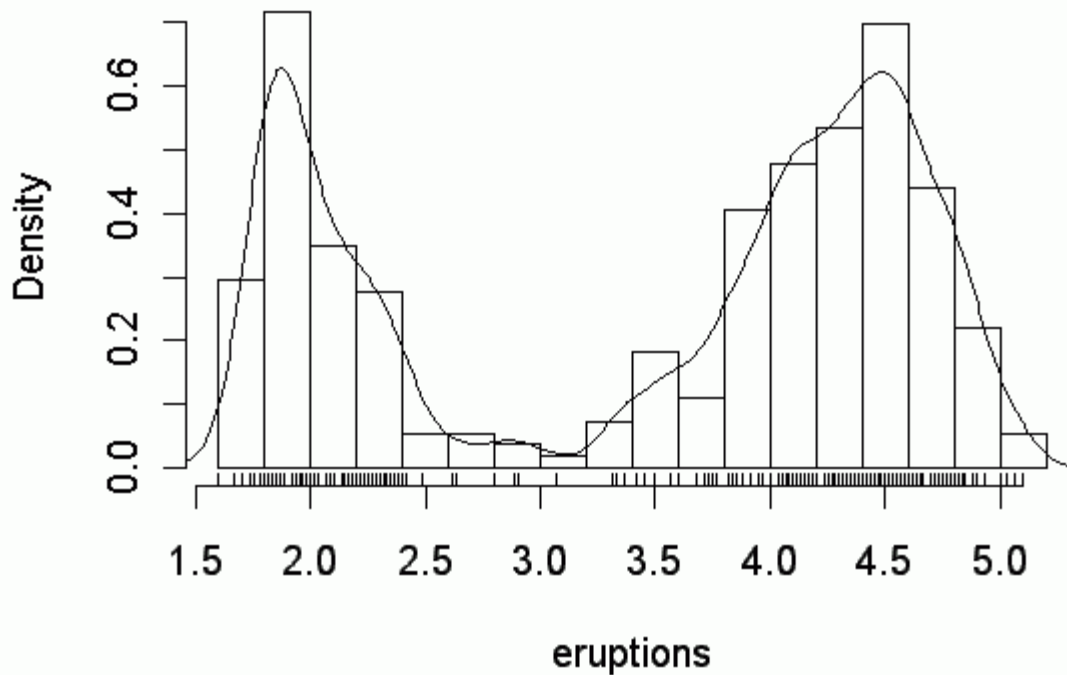
```
> hist(Mileage)
> hist(Mileage, freq=F)      # if probability instead of frequency is desired
```



Let us look at the Old Faithful geyser data, which is a built-in R data set.

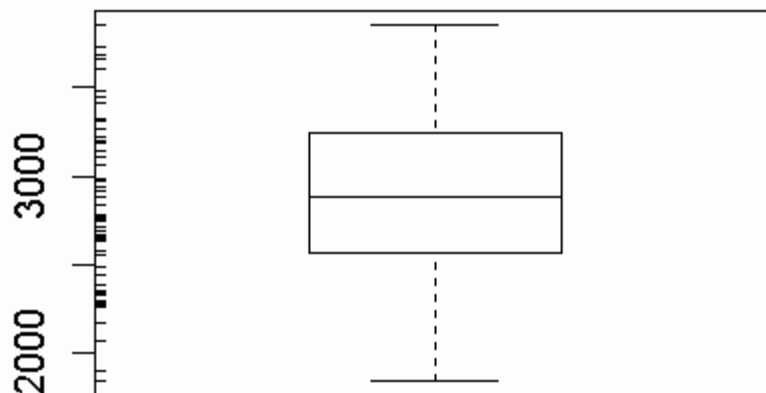
```
> data(faithful)
> attach(faithful)
> names(faithful)
[1] "eruptions" "waiting"
> hist(eruptions, seq(1.6, 5.2, 0.2), prob=T)
> lines(density(eruptions, bw=0.1))
> rug(eruptions, side=1)
```

Histogram of eruptions



3.2 Boxplot

```
> boxplot(Weight) # usual vertical boxplot
> boxplot(Weight, horizontal=T) # horizontal boxplot
> rug(Weight, side=2)
```



If you want to get the statistics involved in the boxplots, the following commands show them. In this example, `a$stats` gives the value of the lower end of the whisker, the first quartile (25th percentile), second quartile (median=50th percentile), third quartile (75th percentile), and the upper end of the whisker.

```
> a<-boxplot(Weight, plot=F)
```

```

> a$stats
      [,1]
[1,] 1845.0
[2,] 2567.5
[3,] 2885.0
[4,] 3242.5
[5,] 3855.0
> a      #gives additional information
> fivenum(Weight)      #directly obtain the five number summary
[1] 1845.0 2567.5 2885.0 3242.5 3855.0

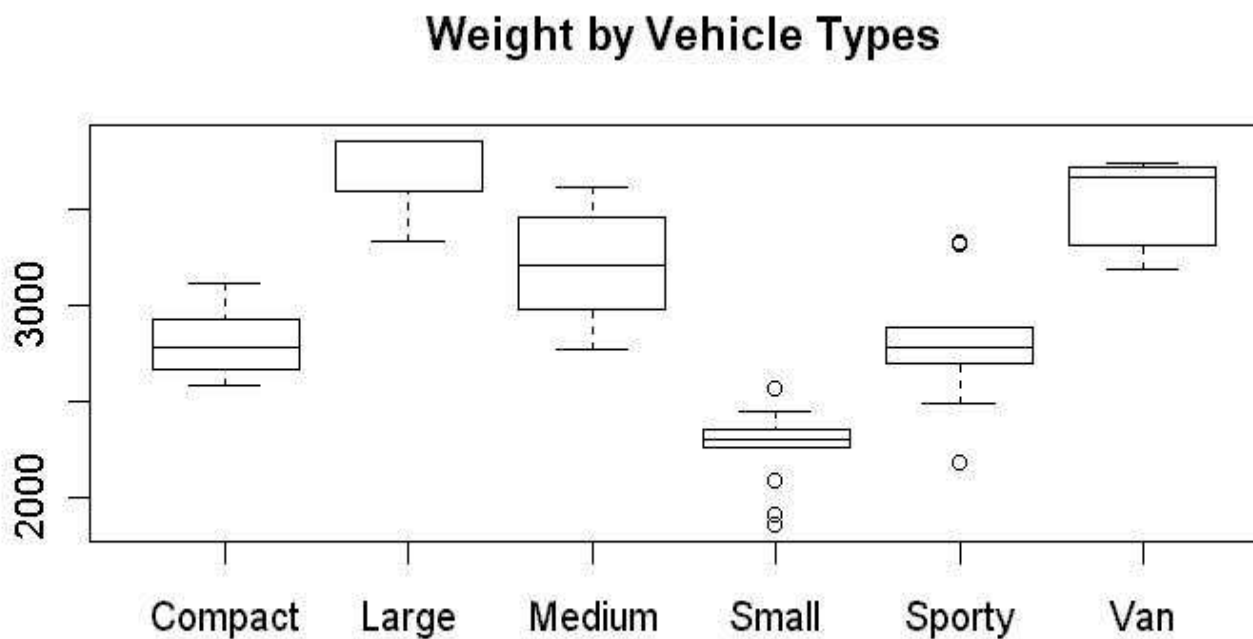
```

Boxplot is more useful when comparing grouped data. For example, side-by-side boxplots of weights grouped by vehicle types are shown below:

```

> boxplot(Weight ~Type)
> title("Weight by Vehicle Types")

```



On-line help is available for the commands:

```

> help(hist)
> help(boxplot)

```

3.3 plot()

plot() is a general graphic command with numerous options.

```

> plot(Weight)

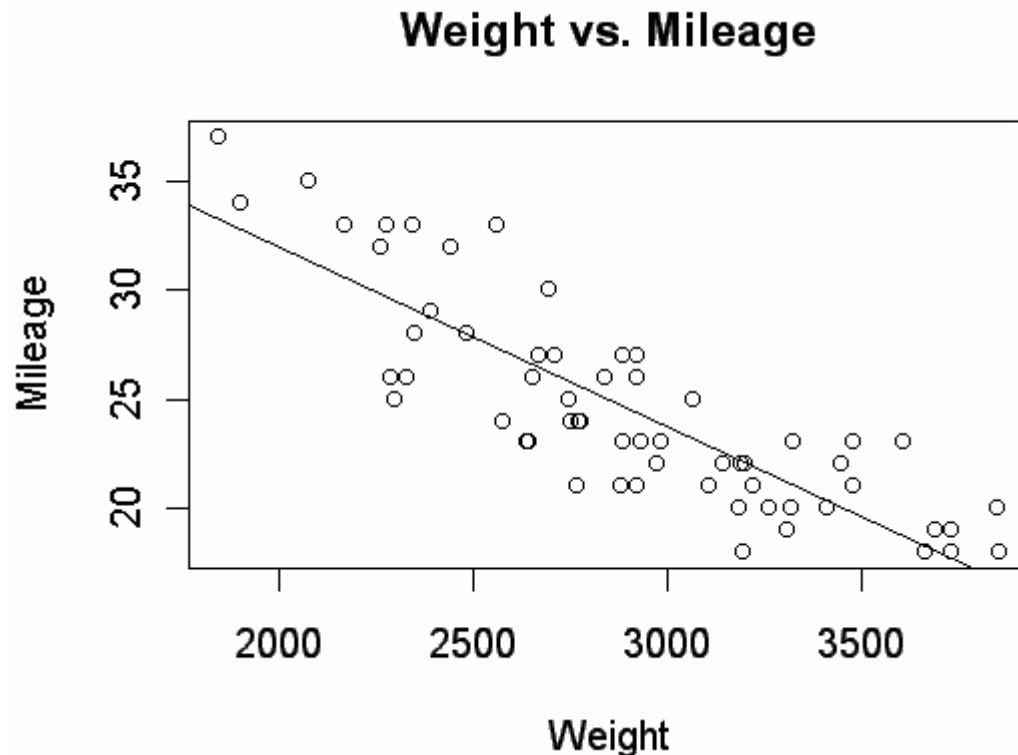
```

The following command produce a scatterplot with Weight on the x-axis and Mileage on the y-axis.

```
> plot(Weight, Mileage, main="Weight vs. Mileage")
```

A fitted straight line is shown in the plot by executing two more commands.

```
> fit<-lm(Mileage~Weight)
> abline(fit)
```



3.4 matplot()

`matplot()` is used to plot two or more vectors of equal length.

```
> y60<-c(316.27, 316.81, 317.42, 318.87, 319.87, 319.43, 318.01, 315.74, 314.00,
313.68, 314.84, 316.03)
> y70<-c(324.89, 325.82, 326.77, 327.97, 327.91, 327.50, 326.18, 324.53, 322.93,
322.90, 323.85, 324.96)
> y80<-c(337.84, 338.19, 339.91, 340.60, 341.29, 341.00, 339.39, 337.43, 335.72,
335.84, 336.93, 338.04)
> y90<-c(353.50, 354.55, 355.23, 356.04, 357.00, 356.07, 354.67, 352.76, 350.82,
351.04, 352.69, 354.07)
> y97<-c(363.23, 364.06, 364.61, 366.40, 366.84, 365.68, 364.52, 362.57, 360.24,
360.83, 362.49, 364.34)
> CO2<-data.frame(y60, y70, y80, y90, y97)
> row.names(CO2)<-c("Jan", "Feb",
"Mar", "Apr", "May", "Jun", "Jul", "Aug", "Sep", "Oct", "Nov", "Dec")
> CO2
```

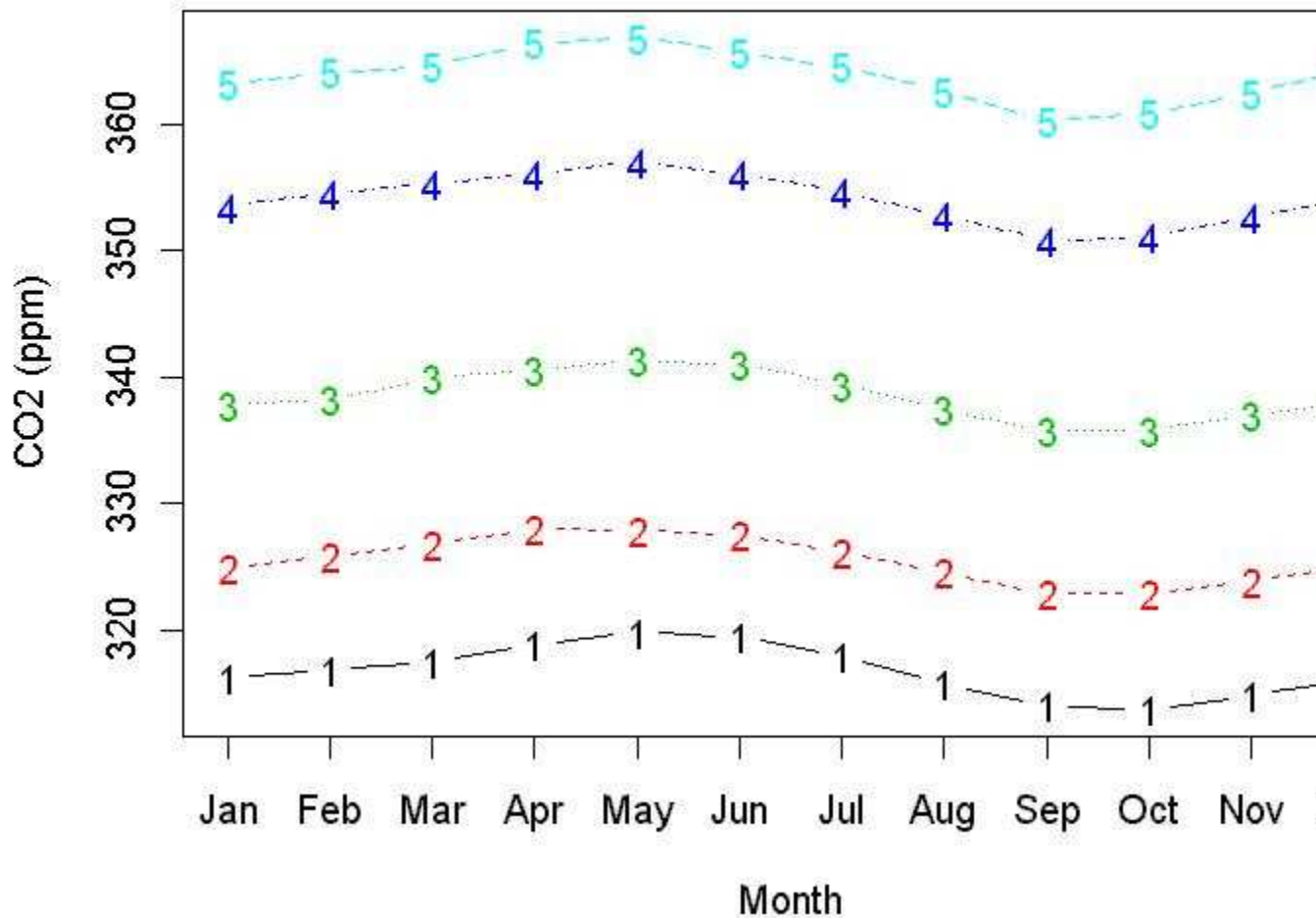
	y60	y70	y80	y90	y97
Jan	316.27	324.89	337.84	353.50	363.23
Feb	316.81	325.82	338.19	354.55	364.06
Mar	317.42	326.77	339.91	355.23	364.61
Apr	318.87	327.97	340.60	356.04	366.40
May	319.87	327.91	341.29	357.00	366.84
Jun	319.43	327.50	341.00	356.07	365.68

```
Jul 318.01 326.18 339.39 354.67 364.52
Aug 315.74 324.53 337.43 352.76 362.57
Sep 314.00 322.93 335.72 350.82 360.24
Oct 313.68 322.90 335.84 351.04 360.83
Nov 314.84 323.85 336.93 352.69 362.49
Dec 316.03 324.96 338.04 354.07 364.34
> matplot(CO2)
```

Note that the observations labeled 1 represents the monthly CO2 levels for 1960, 2 represents those for 1970, and so on. We can enhance the plot by changing the line types and adding axis labels and titles:

```
> matplot(CO2,axes=F,frame=T,type='b',ylab="")
> #axes=F: initially do not draw axis
> #frame=T: box around the plot is drawn;
> #type=b: both line and character represent a series;
> #ylab="": No label for y-axis is shown;
> #ylim=c(310,400): Specify the y-axis range
> axis(2) # put numerical annotations at the tickmarks in y-axis;
> axis(1, 1:12, row.names(CO2))
> # use the Monthly names for the tickmarks in x-axis; length is 12;
> title(xlab="Month") #label for x-axis;
> title(ylab="CO2 (ppm)")#label for y-axis;
> title("Monthly CO2 Concentration \n for 1960, 1970, 1980, 1990 and 1997")
> # two-line title for the matplot
```

Monthly CO2 Concentration for 1960, 1970, 1980, 1990 and 1997



4. Plot Options

4.1 Multiple plots in a single graphic window

You can have more than one plot in a graphic window. For example, `par(mfrow=c(1,2))` allows you to have two plots side by side. `par(mfrow=c(2,3))` allows 6 plots to appear on a page (2 rows of 3 plots each). Note that the arrangement remains in effect until you change it. If you want to go back to the one plot per page setting, type `par(mfrow=c(1,1))`.

4.2 Adjusting graphical parameters

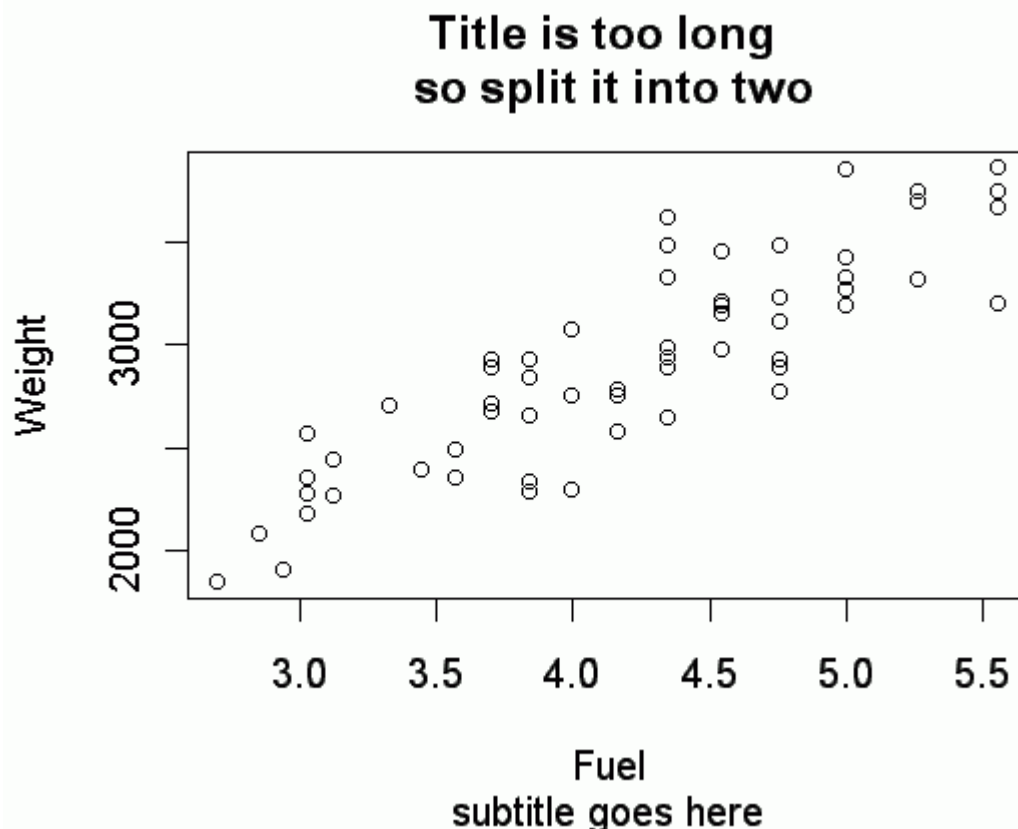
4.2.1 Labels and title; axis limits

Any plot benefits from clear and concise labels which greatly enhances the readability.

```
> plot(Fuel, Weight)
```

If the main title is too long, you can split it into two and adding a subtitle below the horizontal axis label is easy:

```
> title(main="Title is too long \n so split it into two",sub="subtitle goes here")
```



By default, when you issue a plot command R inserts variable name(s) if it is available and figures out the range of x axis and y axis by itself. Sometimes you may want to change these:

```
> plot(Fuel, Weight, ylab="Weight in pounds", ylim=c(1000,6000))
```

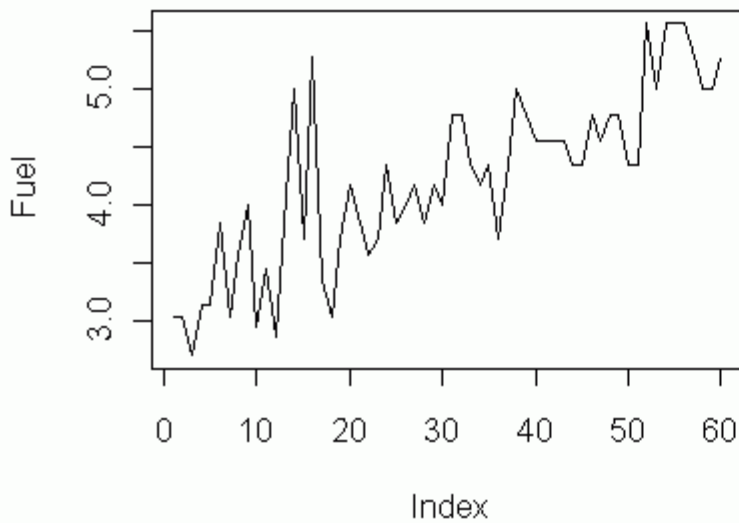
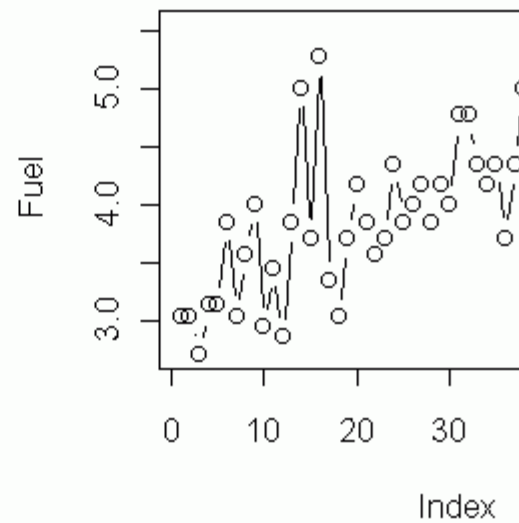
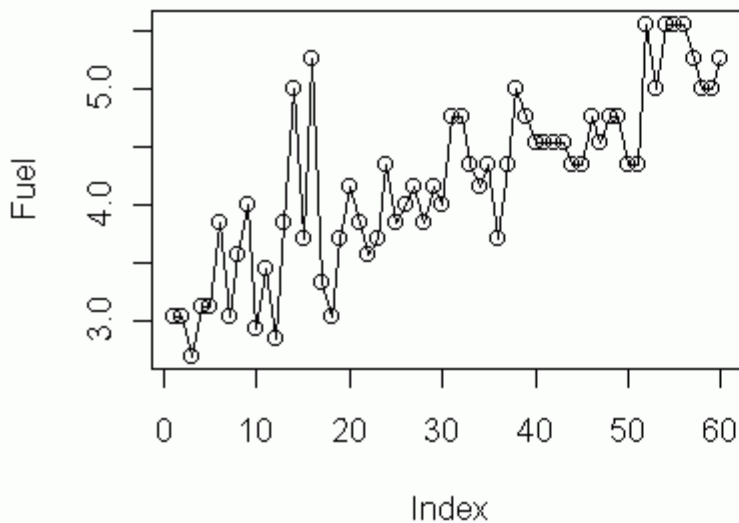
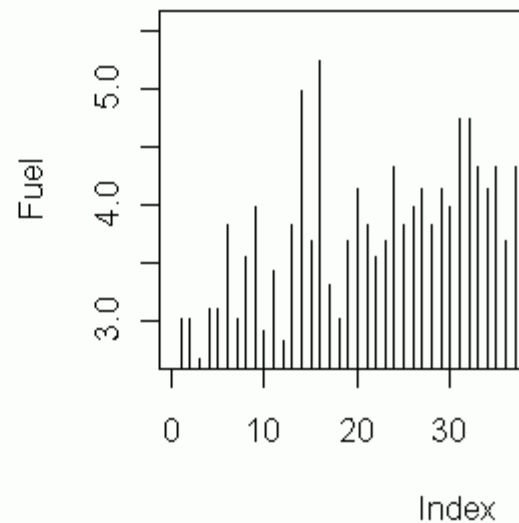
Similarly, you can specify xlab and xlim to change x-axis. If you do not want the default labels to appear, specify xlab=" ", ylab=" ". This give you a plot with no axis labels. Of course you can add the labels after using appropriate statements within title() statement.

```
> plot(Mileage, Weight, xlab="Miles per gallon", ylab="Weight in pounds", xlim=c(20,30),ylim=c(2000,4000))
> title(main="Weight versus Mileage \n data=fuel.frame;", sub="Figure 4.1")
```

4.2.2 Types for plots and lines

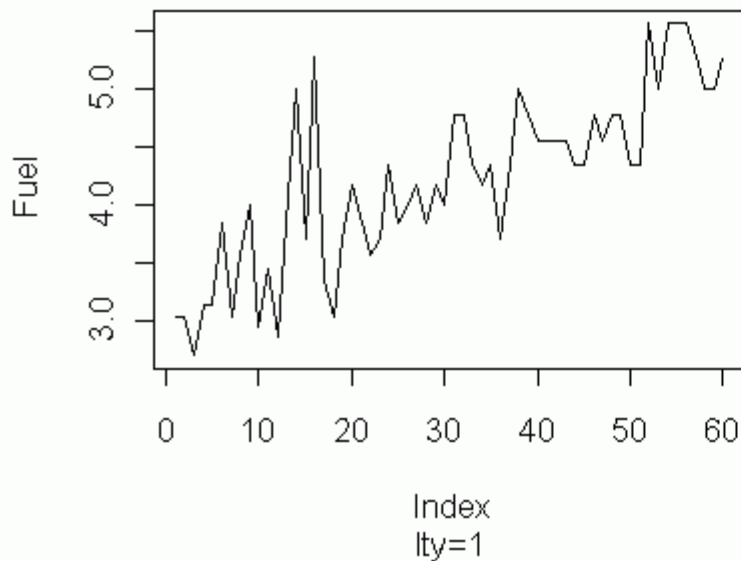
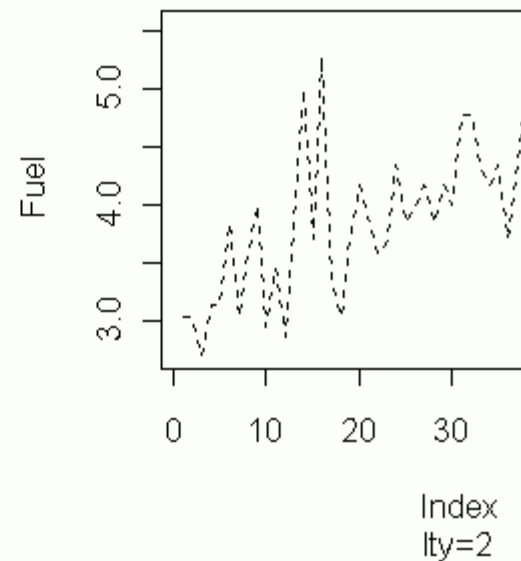
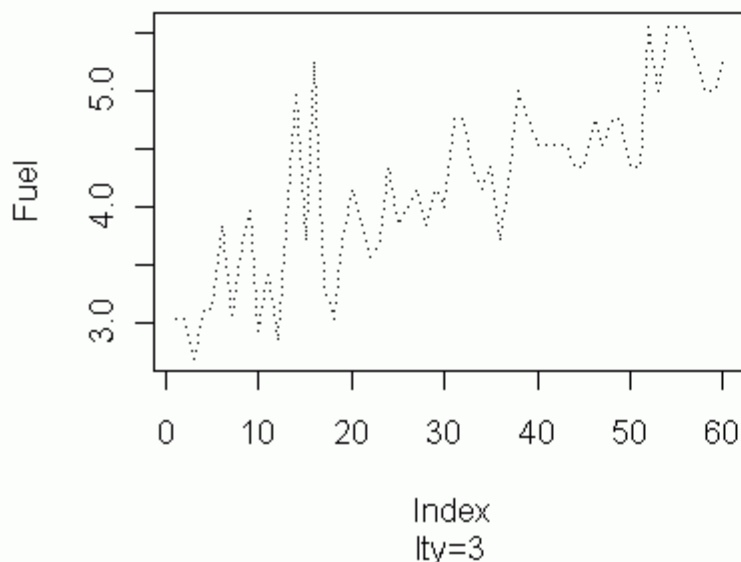
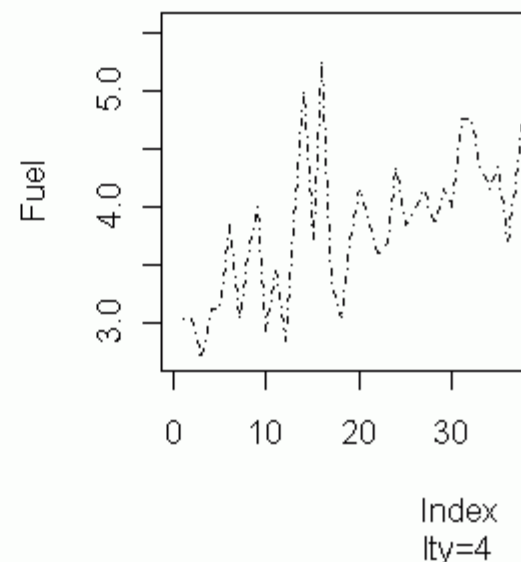
In a series plot (especially time series plot), type provides useful options:

```
> par(mfrow=c(2,2))
> plot(Fuel, type="l"); title("lines")
> plot(Fuel, type="b"); title("both")
> plot(Fuel, type="o"); title("overstruck")
> plot(Fuel, type="h"); title("high density")
```

lines**both****overstruck****high dens**

Also you can specify the line types using `lty` argument within `plot()` command:

```
> plot(Fuel, type="l", lty=1) #the usual series plot
> plot(Fuel, type="l", lty=2) #shows dotted line instead. lty can go up to 8.
> plot(Fuel, type="l", lty=1); title(main="Fuel data", sub="lty=1")
> plot(Fuel, type="l", lty=2); title(main="Fuel data", sub="lty=2")
> plot(Fuel, type="l", lty=3); title(main="Fuel data", sub="lty=3")
> plot(Fuel, type="l", lty=4); title(main="Fuel data", sub="lty=4")
```

Fuel data**Fuel dat****Fuel data****Fuel dat**

Note that we can control the thickness of the lines by `lwd=1` (default) through `lwd=5` (thickest).

4.3 Colors and characters

You can change the color by specifying

```
> plot(Fuel, col=2)
```

which shows a plot with different color. The default is `col=1`. The actual color assignment depends on the system you are using. You may want to experiment with different numbers. Of course you can specify the `col` option together with other options such as `type` or `lty`. `pch` option allows you to choose alternative plotting characters when making a points-type plot. For example, the command

```
> plot(Fuel, pch="*")      # plots with * characters
> plot(Fuel, pch="M")     # plots with M.
```

4.4 Controlling axis line

`bty="n"`; No box is drawn around the plot, although the x and y axes are still drawn.

`bty="o"`; The default box type; draws a four-sided box around the plot.

`bty="c"`; Draws a three-sided box around the plot in the shape of an uppercase "C."

`bty="l"`; Draws a two-sided box around the plot in the shape of an uppercase "L."

`bty="7"`; Draws a two-sided box around the plot in the shape of a square numeral "7."

```
> par(mfrow = c(2,2))
> plot(Fuel)
> plot(Fuel, bty="l")
> plot(Fuel, bty="7")
> plot(Fuel, bty="c")
```

4.5 Controlling tick marks

`tck` parameter is used to control the length of tick marks. `tck=1` draws grid lines. Any positive value between 0 and 1 draws inward tick marks for each axis. Also with some more work you can have tick marks of different lengths, as the following example shows.

```
> plot(Fuel, main="Default")
> plot(Fuel, tck=0.05, main="tck=0.05")
> plot(Fuel, tck=1, main="tck=1")
> plot(Fuel, axes=F, main="Different tick marks for each axis")
> #axes=F suppresses the drawing of axis
> axis(1)# draws x-axis.
> axis(2, tck=1, lty=2) # draws y-axis with horizontal grid of dotted line
> box()# draws box around the remaining sides.
```

4.6 Legend

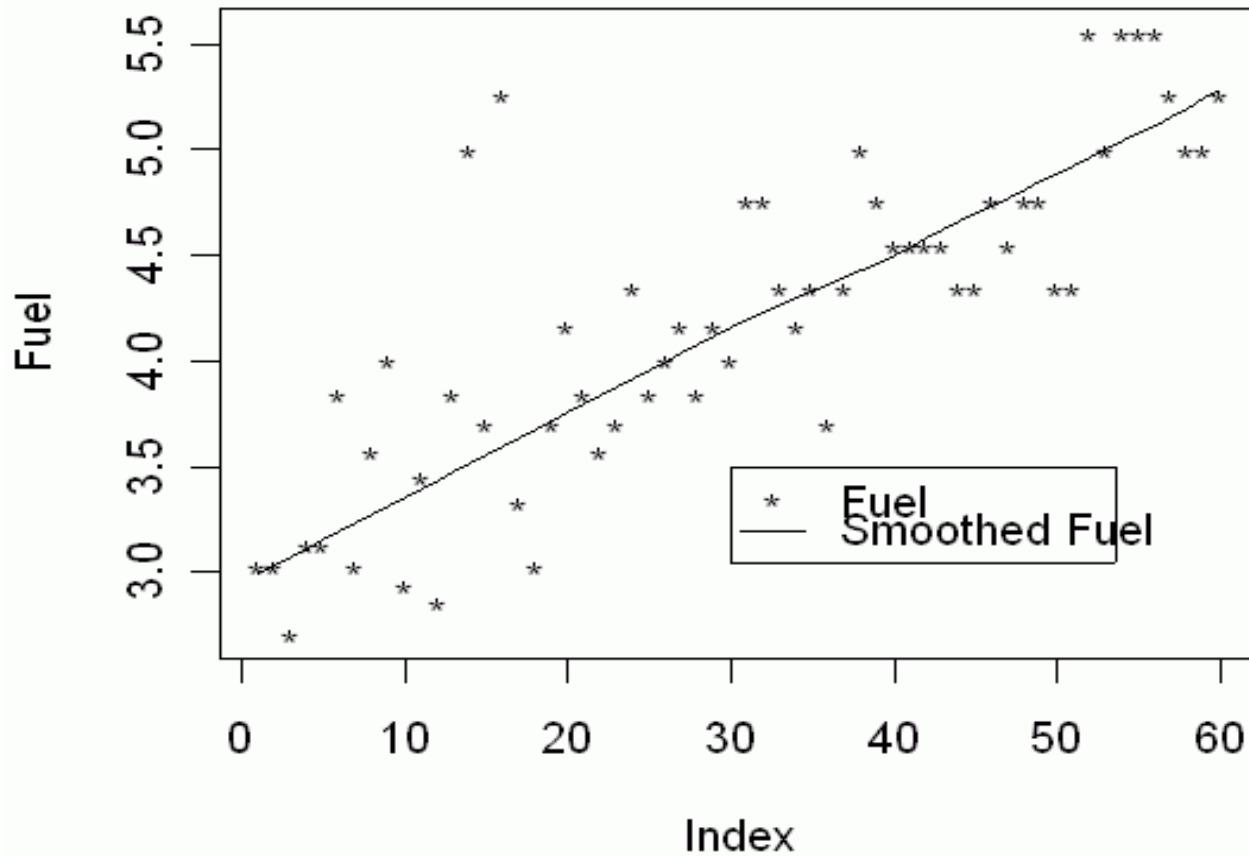
`legend()` is useful when adding more information to the existing plot.

In the following example, the `legend()` command says

(1) put a box whose upper left corner coordinates are `x=30` and `y=3.5`;

(2) write the two texts `Fuel` and `Smoothed Fuel` within the box together with corresponding symbols described in `pch` and `lty` arguments.

```
>par(mfrow = c(1,1))
>plot(Fuel)
>lines(lowess(Fuel))
>legend(30,3.5, c("Fuel","Smoothed Fuel"), pch="* ", lty=c(0,1))
```

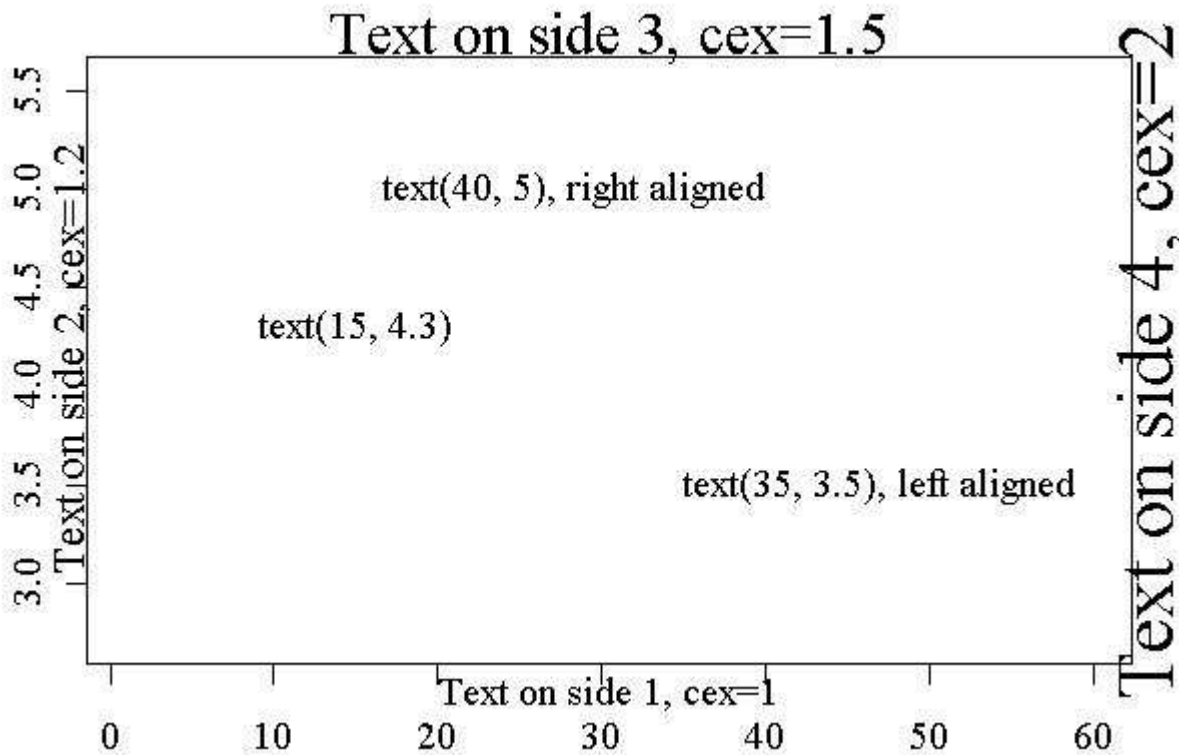


If you want to keep the legend box from appearing, add `bty="n"` to the legend command.

4.7 Putting text to the plot; controlling the text size

`mtext()` allows you to put texts to the four sides of the plot. Starting from the bottom (`side=1`), it goes clockwise to side 4. The plot command in the example suppresses axis labels and the plot itself. It just gives the frame. Also shown is the use of `cex` (character expansion) argument which controls the relative size of the text characters. By default, `cex` is set to 1, so graphics text and symbols appear in the default font size. With `cex=2`, text appears at twice the default font size. `text()` statement allows precise positioning of the text at any specified point. First text statement puts the text within the quotation marks centered at $x=15$, $y=4.3$. By using optional argument `adj`, you can align to the left (`adj=0`) such that the specified coordinates are the starting point of the text.

```
> plot(Fuel, xlab=" ", ylab=" ", type="n")
> mtext("Text on side 1, cex=1", side=1,cex=1)
> mtext("Text on side 2, cex=1.2", side=2,cex=1.2)
> mtext("Text on side 3, cex=1.5", side=3,cex=1.5)
> mtext("Text on side 4, cex=2", side=4,cex=2)
> text(15, 4.3, "text(15, 4.3)")
> text(35, 3.5, adj=0, "text(35, 3.5), left aligned")
> text(40, 5, adj=1, "text(40, 5), right aligned")
```



4.8 Adding symbols to plots

`abline()` can be used to draw a straight line to a plot.

`abline(a,b)` $a=y$ -intercept, b =slope.

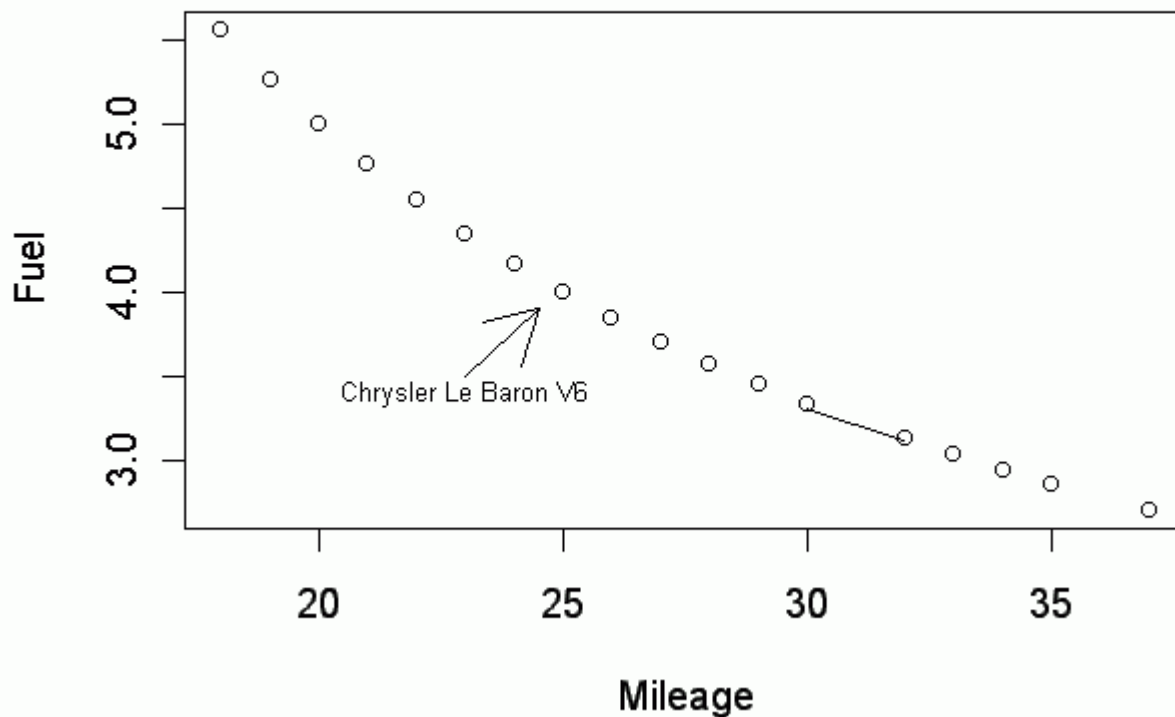
`abline(h=30)` draws a horizontal line at $y=30$.

`abline(v=12)` draws a vertical line at $x=12$.

4.9 Adding arrow and line segment

```
> plot(Mileage, Fuel)
> arrows(23,3.5,24.5,3.9)
> segments(31.96713,3.115541, 29.97309,3.309592)
> title("arrow and segment")
> text(23,3.4,"Chrysler Le Baron V6", cex=0.7)
```

arrow and segment



4.10 Identifying plotted points

While examining a plot, identifying a data point such as possible outliers can be achieved using `identify()` function.

```
> plot(Fuel)
> identify(Fuel, n=3)
```

After pressing return, R waits for you to identify ($n=3$) points with the mouse. Moving the mouse cursor over the graphics window and click on a data point. Then the observation number appears next to the point, thus making the point identifiable.

4.11 Managing graphics windows

Normally high level graphic commands (`hist()`, `plot()`, `boxplot()`, ...) produce a plot which replaces the previous one. To avoid this, use `win.graph()` to open a separate graphic window. Even if more than one graphics windows are open, only one window is active, i.e., as long as you don't change the active window, subsequent plotting commands will show the plot in that particular window. `dev.cur()` gives the current active window, `dev.list()` lists all available graphics windows, `dev.set()` changes the active window, `dev.off()` closes the current graphic window, and `graphics.off()` closes all the open graphics windows at once. The following examples assume that currently no graphic window is open.

```
> for (i in 1:3) win.graph()      #open three graphic windows
> dev.list()
windows windows windows
      2       3       4
> dev.cur()
windows
```

```

4
> dev.set(3)      #change the current window to window 3
windows
3
> dev.cur()      #check it
windows
3
> dev.off()      #close the current window and window 4 is active
windows
4
> dev.list()
windows windows
2 4
> graphics.off() # now close all three
> dev.list()
NULL

```

5. Statistical Analysis

5.1 Descriptive statistics

`summary()` returns five number summary plus mean for numeric vector and returns the frequencies for categorical vector.

```

> fuel.frame<-read.table("c:/fuel_frame.txt", header=T, sep=",")
> names(fuel.frame)
[1] "row.names" "Weight"      "Disp."      "Mileage"    "Fuel"      "Type"
> attach(fuel.frame)
> summary(Mileage)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
 18.00  21.00  23.00  24.58  27.00  37.00
> summary(fuel.frame)
      row.names      Weight      Disp.
Acura.Legend.V6   : 1  Min.    :1845      Min.    : 0.73
Audi.80.4         : 1  1st Qu.:2571      1st Qu.:113.75
Buick.Century.4   : 1  Median :2885      Median :144.50
Buick.Le.Sabre.V6 : 1  Mean    :2901      Mean    :136.92
Buick.Skylark.4   : 1  3rd Qu.:3231      3rd Qu.:180.00
Chevrolet.Beretta.4: 1  Max.    :3855      Max.    :305.00
(Other)          :54

      Mileage      Fuel      Type
Min.    :18.00      Min.    :2.703      Compact:15
1st Qu.:21.00      1st Qu.:3.704      Large  : 3
Median :23.00      Median :4.348      Medium :13
Mean    :24.58      Mean    :4.210      Small  :13
3rd Qu.:27.00      3rd Qu.:4.762      Sporty : 9
Max.    :37.00      Max.    :5.556      Van    : 7

```

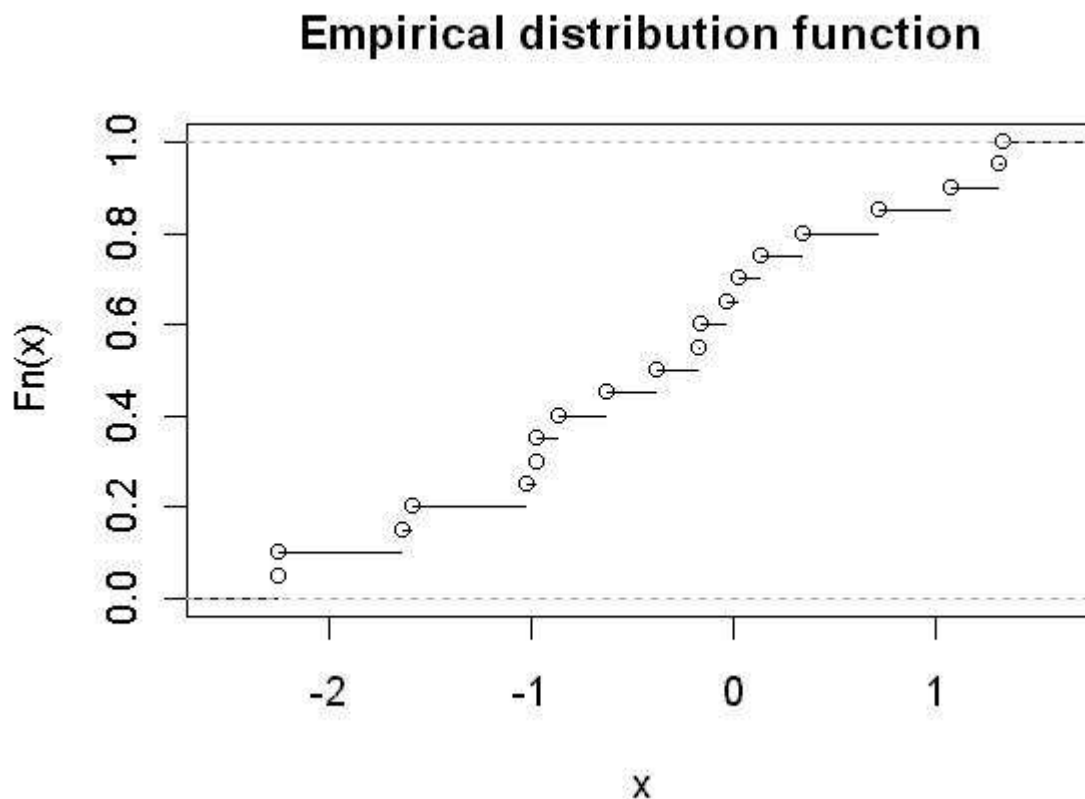
`var()` returns the sample variance, `sd()` the sample standard deviation, and `cor()` the sample correlation coefficient between two vectors:

```
> var(Mileage)
```

```
[1] 22.95904
> sd(Mileage)
[1] 4.791559
> cor(Mileage,Weight)
[1] -0.8478541
```

5.2 Empirical distribution function

```
> library(stepfun) # call a library of functions
> F20<-rnorm(20) # generate a normal random sample of size 20
> plot.ecdf(F20,main="Empirical distribution function")
```



5.3 One sample and two sample t tests

Recall the CO2 data (CO2 concentration in the atmosphere). The following command performs a one-sample t-test whether the average CO2 level for the year 1960 is 320 ppm. By default, it does a two-sided test and extremely small p-value indicates that the null hypothesis is rejected for any reasonable choice of alpha.

```
> t.test(CO2$y60, mu=320)
```

```
One Sample t-test
```

```
data: CO2$y60
```

```
t = -5.5183, df = 11, p-value = 0.0001812
```

```
alternative hypothesis: true mean is not equal to 320
```

```
95 percent confidence interval:
```

```

315.4502 318.0448
sample estimates:
mean of x
316.7475

```

Now we perform a two-sample independent t-test of equal mean for the CO2 level of 1960 and 1970. We assume that the variances for the two populations are equal. The average concentrations are significantly different, just as the test shows.

```
> t.test(CO2$y60, CO2$y70, var.equal=T)
```

```
Two Sample t-test
```

```

data: CO2$y60 and CO2$y70
t = -11.1522, df = 22, p-value = 1.602e-10
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -10.40088 -7.13912
sample estimates:
mean of x mean of y
316.7475 325.5175

```

Paired t-test is also available. All you have to do is to include `paired=T` within `t.test()` argument.

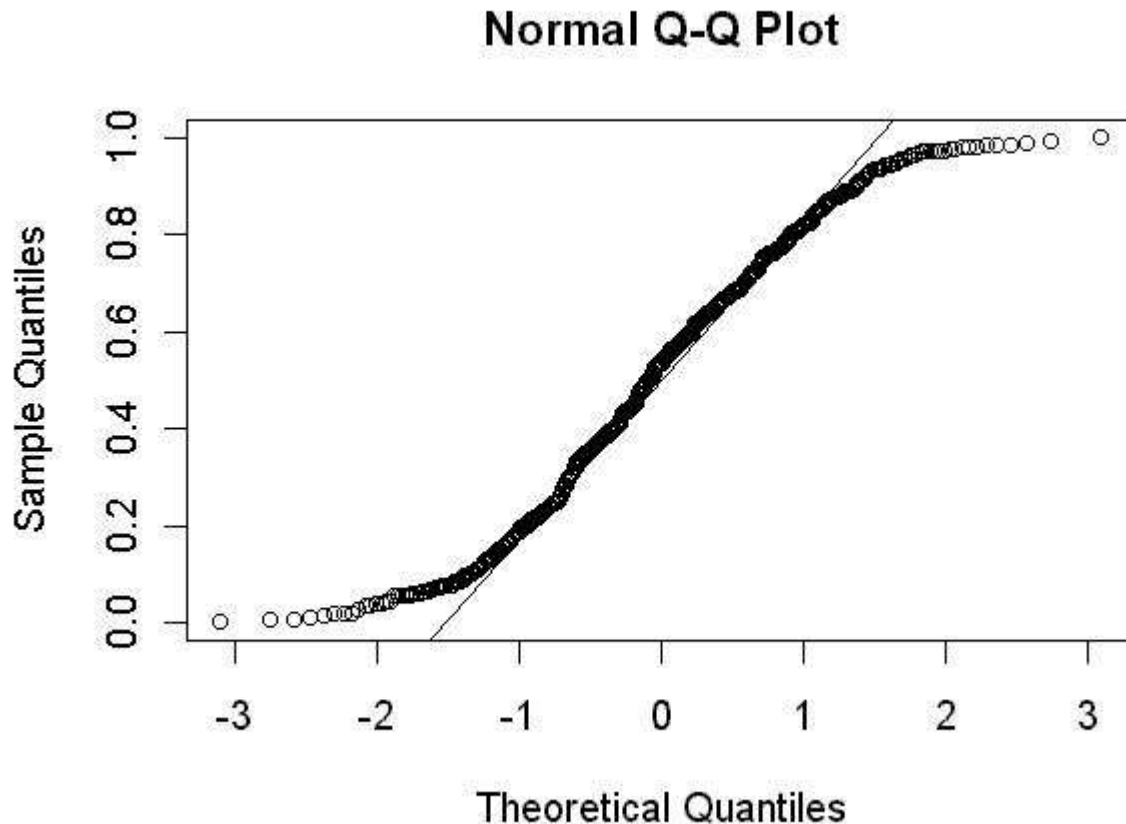
5.4 Checking normality

Quite a few statistical tests are based on the normality of the underlying population. Here we illustrate normal plot and Kolmogorov-Smirnov test to check the normality assumption.

```

> #generate 500 observations from uniform (0,1) distribution
> F500<-runif(500);a<-c(mean(F500),sd(F500))
> qqnorm(F500)      #normal probability plot
> qqline(F500)     #ideal sample will fall near the straight line

```



Obviously the curve is far from the straight line so we strongly suspect the normality (if we didn't know that the generated data came from uniform). We formally test the normality by performing Kolmogorov-Smirnov test, comparing the empirical distribution of F500 to a comparable normal distribution with the mean and standard deviation same as that of F500.

```
> ks.test(F500, "pnorm", mean=a[1], sd=a[2])
```

One-sample Kolmogorov-Smirnov test

```
data: F500
D = 0.0655, p-value = 0.02742
alternative hypothesis: two.sided
```

5.5 Analysis of variance

ANOVA is an extension of a two-sample t test, testing the equality of means of more than two groups. In the example below, we use `aov()` function to test the equality of average weight per vehicle type.

```
> a<-aov(Weight~Type)
> summary(a)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Type	5	11161777	2232355	36.035	4.855e-16 ***
Residuals	54	3345331	61951		

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> boxplot(Weight~Type) #side-by-side boxplot
```

5.6 Linear regression model

`lm()` fits a linear regression model. `summary()` returns rich information for the fit.

```
> attach(fuel.frame)
> names(fuel.frame)
[1] "row.names" "Weight"      "Disp."      "Mileage"    "Fuel"      "Type"

> fit1<-lm(Mileage~Weight+Disp.)
> fit1      #gives model specification and regression coefficients
```

Call:

```
lm(formula = Mileage ~ Weight + Disp.)
```

Coefficients:

```
(Intercept)      Weight      Disp.
  44.379702    -0.006044    -0.016540
```

```
> summary(fit1)      #more information is available
```

Call:

```
lm(formula = Mileage ~ Weight + Disp.)
```

Residuals:

```
      Min       1Q   Median       3Q      Max
-4.5726 -1.5814 -0.2569  1.8499  4.6783
```

Coefficients:

```
              Estimate Std. Error t value Pr(>|t|)
(Intercept)  44.379702   2.654526  16.719 < 2e-16 ***
Weight       -0.006044   0.001188  -5.088 4.23e-06 ***
Disp.        -0.016540   0.007641  -2.165  0.0346 *
---

```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Residual standard error: 2.485 on 57 degrees of freedom

Multiple R-Squared: 0.7402, Adjusted R-squared: 0.7311

F-statistic: 81.21 on 2 and 57 DF, p-value: < 2.2e-16

```
> names(fit1)
[1] "coefficients" "residuals"      "effects"        "rank"
[5] "fitted.values" "assign"         "qr"            "df.residual"
[9] "xlevels"      "call"          "terms"         "model"
```

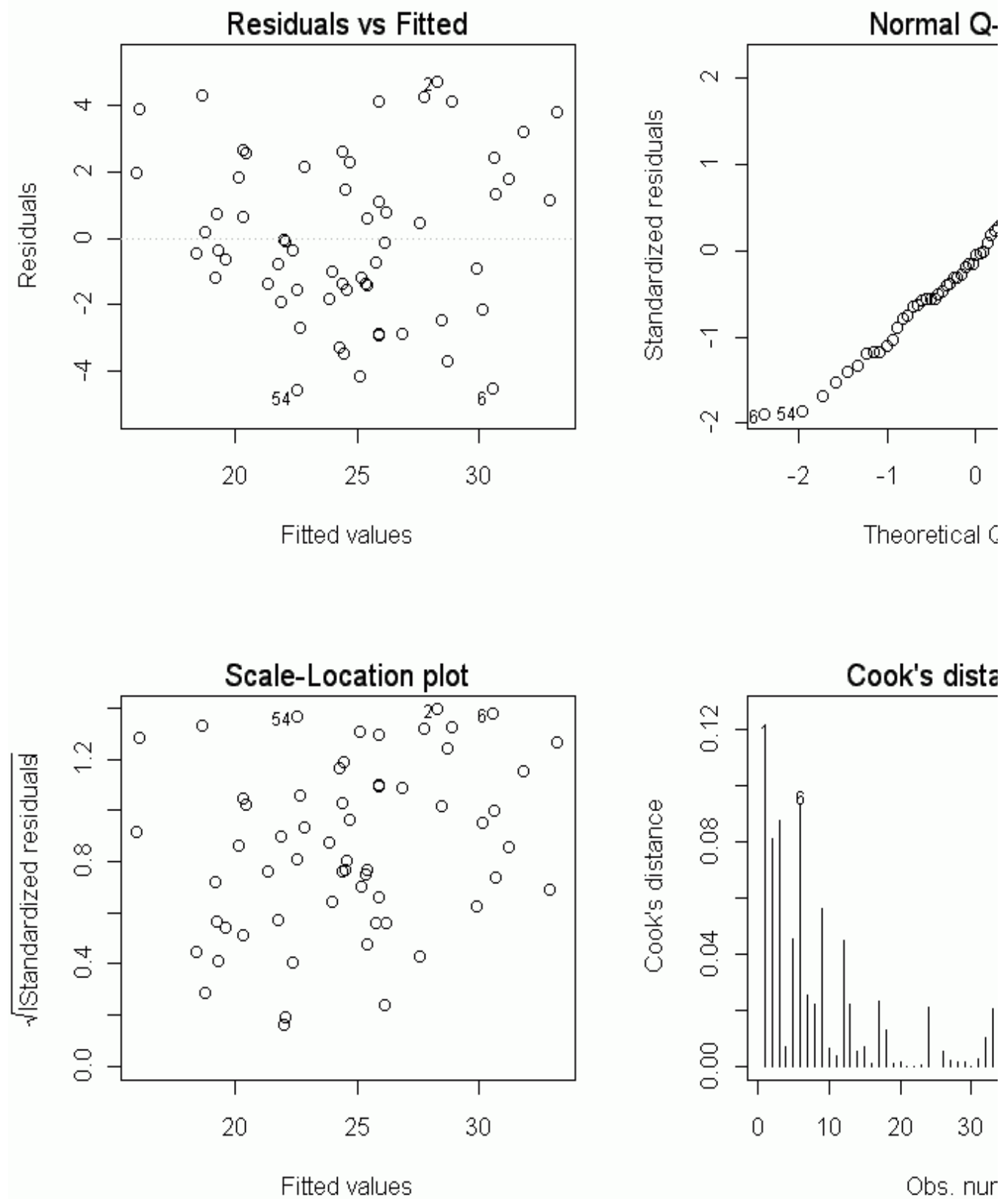
```
> plot(fit1$fitted.values,fit1$residuals) #residual plot
```

If no intercept term is desired, use the following command:

```
> fit2<-lm(Mileage~ -1+Weight+Disp.)
```

Usual diagnostic plots can be obtained by requesting `plot(fit1)`:

```
> par(mfrow=c(2,2))
> plot(fit1)
```



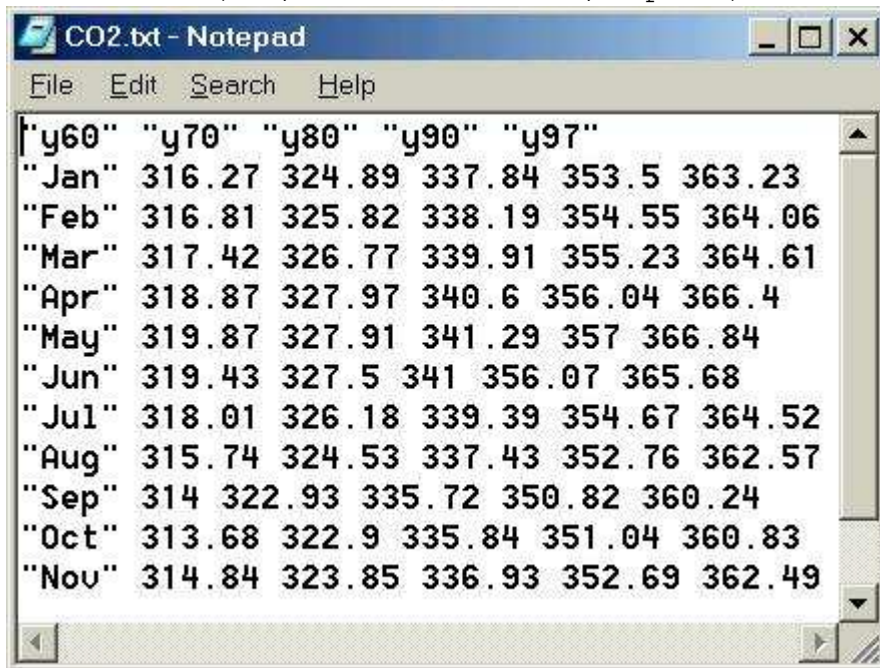
6. Miscellanies

6.1 Data import/export

Small to moderate size data sets can be easily handled using tools presented so far. However, quite often we have a garden variety of data sources from data handling programs. By far the easiest way to import and export the data in R is using text files. Save the data in plain text format which may be imported to a different software. That way, you can easily view the data using any of the capable text editor even when the original software that produced the data is no longer available.

`write.table()` outputs the specified data frame to a file. A blank space is used to separate columns when `sep=" "` is specified within its argument. Other popular choices include comma (`sep=","`), and tab (`sep="\t"`).

```
> CO2      # data frame
> write.table(CO2, file="c:/CO2.txt", sep=" ")
```



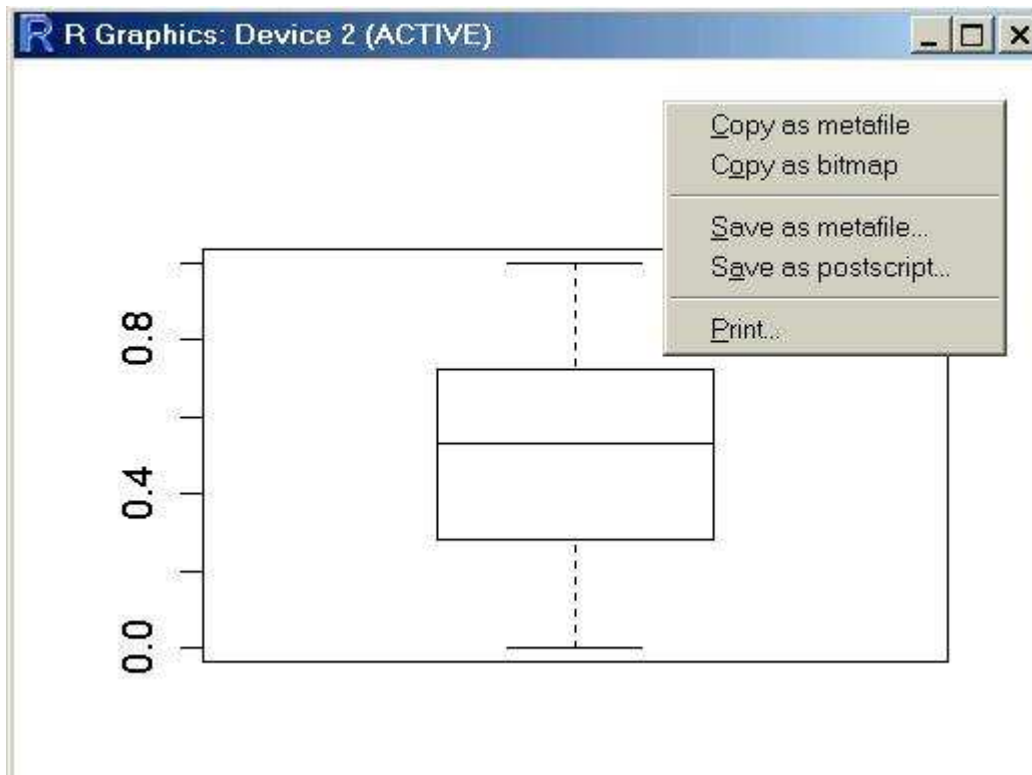
On the other hand, `read.table()` reads in an external text file and creates a data frame. For example, if the first line of the text data file `file.dat` consists of variable names, the following command will do the job:

```
> data1<-read.table("c:/file.dat", header=TRUE)
```

`getwd()` returns the current working directory and `setwd()` changes it.

```
> getwd()
[1] "C:\\Program Files\\R\\rw1070"
> setwd("c:/")      # set the root directory as the working directory
> getwd()
[1] "c:\\\"
> read.table(file="CO2.txt")
> # now pathname is not required to read data files in the root directory
```

6.2 Saving graphical output



Right clicking anywhere inside the active graphics window shows a context sensitive menu, allowing either saving the plot as metafile (EMF) or postscript format (PS). On the other hand, Copy as metafile or Copy as bitmap (BMP) puts the information in the clipboard, a temporary memory area used by Windows. In the latter, you need to immediately paste it in some applications which understand the graphics format, e.g., MS Word. More graphical formats are available from the main menu. While the graphic window is active, click File| Save As from the menu and it lists six file formats (metafile, postscript, PDF, PNG, BMP, and JPG at three quality levels) in total so you have plenty of choices.

Some comments on the choice of graphic formats are in order. In general metafile format retains graphic quality even when it is resized in the application. On the other hand, JPG is a very popular choice on the Internet and file size is usually much smaller than metafile. Except for rare circumstances, I would not recommend BMP file format because it is usually very large and shows very poor picture quality when resized. Postscript file format is useful when including the graphic file in another postscript file or when postscript printer is available. Picture quality does not deteriorate when resized, and it is the default file format to be included in TeX documents.

6.3 Missing values

NA (Not Applicable) is used to denote missing values. Since many functions returns NA if missing values are present, we illustrate how to handle them.

```
> x                                #contains a missing value
[1] 1 2 3 4 5 NA
> mean(x)                           #doesn't work
[1] NA
> is.na(x)                          #returns a logical vector
[1] FALSE FALSE FALSE FALSE FALSE TRUE
> sum(is.na(x))                      #number of NA's in the vector
```

```
[1] 1
> x1<-x[!is.na(x)];x1      #retain only non-missing cases
[1] 1 2 3 4 5
> a<-mean(x[!is.na(x)]);a  #compute the average value of the non-missing cases
[1] 3
> x2<-x
> x2[is.na(x)]<-a;x2      #impute the missing by the average value
[1] 1 2 3 4 5 3
```

The following example shows how to select those listwise nonmissing cases.

```
> data2
  var1 var2 var3
1    1  2.3  aa
2    4  3.2 <NA>
3    3  5.4  bc
4   NA  2.7  ed
5    3  4.1  dd
> a1<-!is.na(data2$var1);a1  #TRUE if nonmissing for var1
[1] TRUE TRUE TRUE FALSE TRUE
> a2<-!is.na(data2$var2);a2
[1] TRUE TRUE TRUE TRUE TRUE
> a3<-!is.na(data2$var3);a3
[1] TRUE FALSE TRUE TRUE TRUE
> data3<-data2[a1*a2*a3==1,]
> #select those rows if all of the elements are nonmissing.
> data3
  var1 var2 var3
1    1  2.3  aa
3    3  5.4  bc
5    3  4.1  dd
```

6.4 Getting help

By default, R has a couple of excellent manuals in PDF format. "An Introduction to R" is almost a required reading to begin using R. To access the manual, click [Help | Manuals](#) and the list of available documents will be shown. Also use `help()` to get command-specific information.

```
> help(read.table)
```

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Page location: <http://www.math.ilstu.edu/dhkim/Rstuff/Rtutor.html>

