

101 Default R Graphs

Cevi Herdian

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Chapter 1

Preface

Note:

- An R Notebook is an R Markdown document with chunks that can be executed independently and interactively, with output visible immediately beneath the input.
- Notebook output are available as HTML, PDF, Word, or Latex.
- This Notebook as HTML is preferably open with Google Chrome.
- R-Code can be extracted as Rmd file under the button “Code” in the notebook.

Why R?:

- Free and open source
- Built for statistical computing
- Visualization tools
- Tools for building Models

Learning Objectives:

- How to create beautiful, useful, and insightful graphics and charts
- How to customize the look of them

All languages have their inconsistencies include R Programming. This documentation helps us to create visualizing in default R graphics before to far away using the packages.

Change log update:

- 18.12.2018
- 19.12.2018
- 23.12.2018
- 24.12.2018
- 13.05.2019
- 14.05.2019
- 15.05.2019

Preferences:

- R Programming/Graphics
- Tutorials for learning R
- RDocumentation
- Statistical tools for high-throughput data analysis



Figure 1.1:

- [statmethods](#)
- [cognitiveclass.ai](#)-Data Visualization with R
- [Comprehensive Guide to Data Visualization in R](#)
- [datascienceplus](#)
- [Hands-On Programming with R](#)
- [R for Data Science](#)
- [R Markdown: The Definitive Guide](#)
- [R bookdown: Authoring Books and Technical Documents with R Markdown](#)
- [blogdown: Creating Websites with R Markdown](#)

License:

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Chapter 2

Creating and Saving Graphs in R

Creating graphs:

Used `mtcars` (Motor Trend Car Road Tests from default dataset) dataset for this section:

```
?mtcars
```

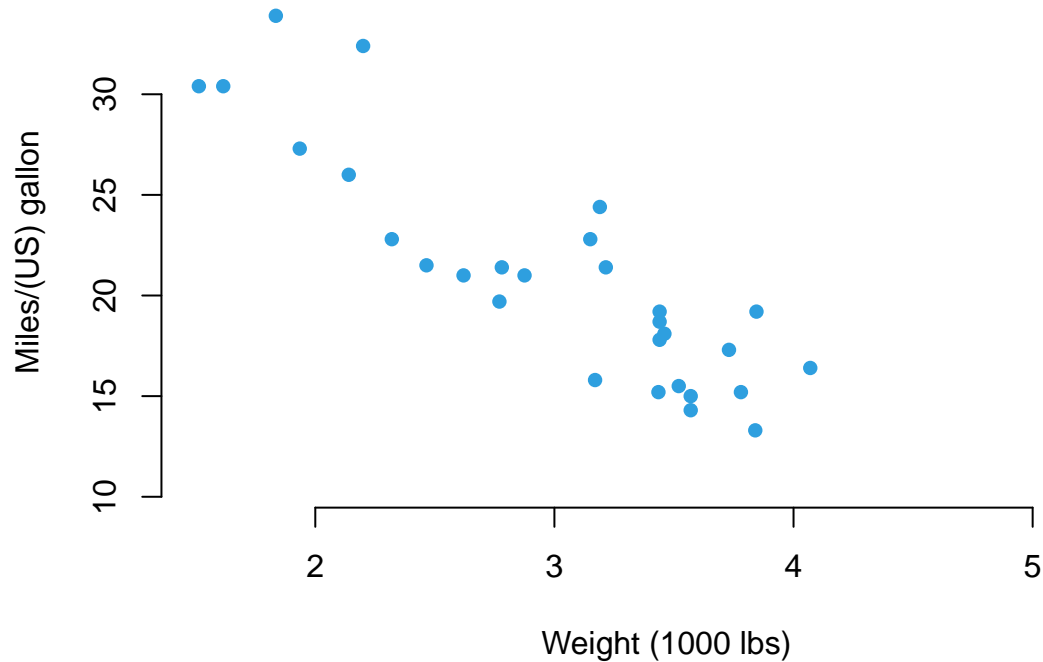
```
## starting httpd help server ... done
```

```
head(mtcars,5)
```

```
##           mpg cyl disp  hp drat   wt  qsec vs am gear carb
## Mazda RX4      21.0   6  160 110 3.90 2.620 16.46  0  1    4    4
## Mazda RX4 Wag  21.0   6  160 110 3.90 2.875 17.02  0  1    4    4
## Datsun 710      22.8   4  108  93 3.85 2.320 18.61  1  1    4    1
## Hornet 4 Drive  21.4   6  258 110 3.08 3.215 19.44  1  0    3    1
## Hornet Sportabout 18.7   8  360 175 3.15 3.440 17.02  0  0    3    2
```

Created a graphs with `plot()` function:

```
plot(x = mtcars$wt, y = mtcars$mpg,
     pch = 16, frame = FALSE,
     xlab = "Weight (1000 lbs)", ylab = "Miles/(US) gallon", col = "#2E9FDF")
```



Saving graphs:

There are two ways of saving graphs in R

1. In RStudio IDE:

Plots panel -> Export -> Save as Image or Save as PDF

#preference: <http://www.sthda.com/english/wiki/r-base-graphs>

2. Using R codes:

- 1. Choose the format
- 2. Create the graphs
- 3. Enter the `dev.off()` command

Example->

Note: The file you save are in the current working directory.

The command to get directory is `getwd()`

```
# 1. Choose the format
png("rplot.jpg") #width = 25, height = "25"

# 2. Create the graphs
plot(x = mtcars$wt, y = mtcars$mpg,
     pch = 16, frame = FALSE,
     xlab = "wt", ylab = "mpg", col = "#2E9FDF")
```

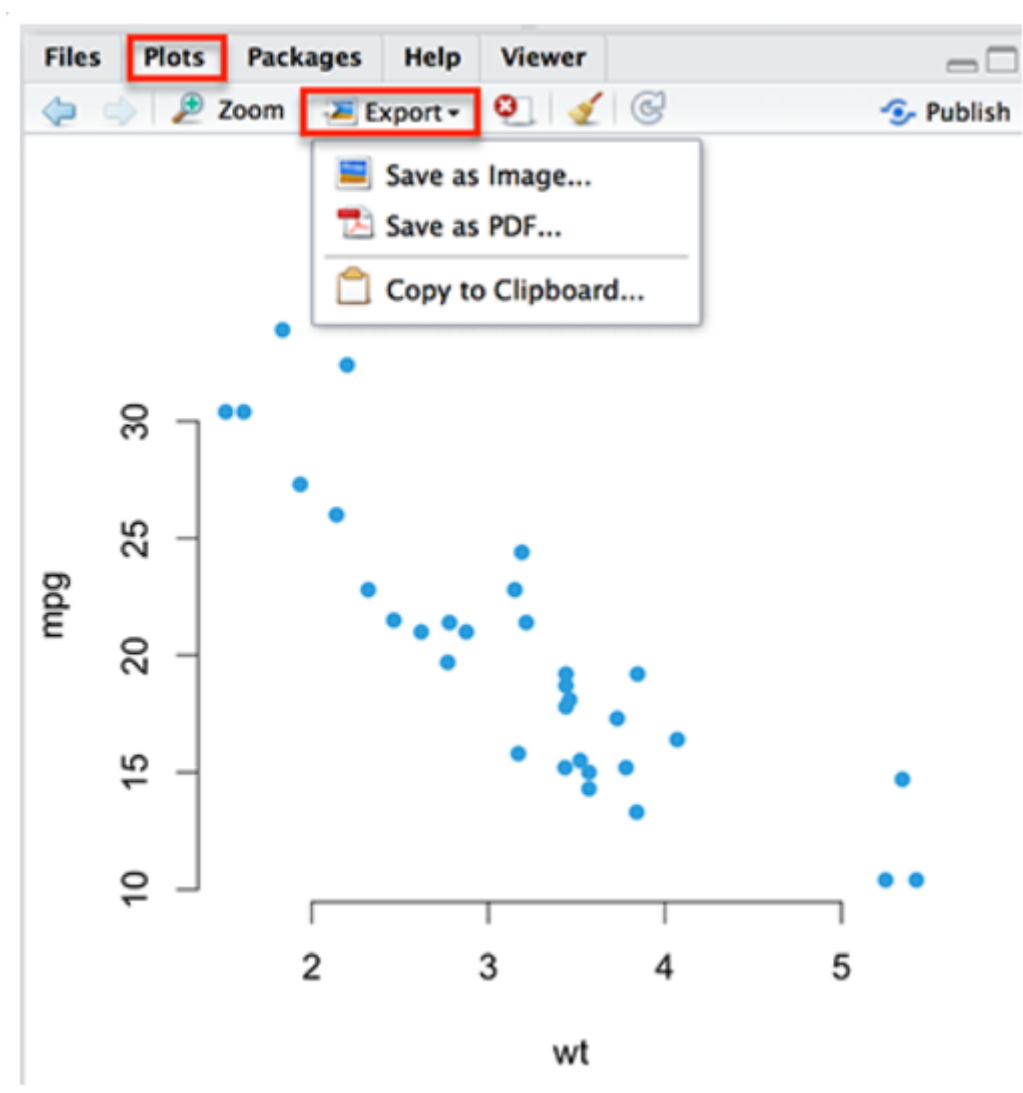


Figure 2.1:

```
# 3. Save the file with dev.off() command in directory  
dev.off()
```

```
## pdf  
## 2
```

File formats for exporting plots are:

- `pdf("rplot.pdf")` : pdf file
- `png("rplot.png")` : png file
- `jpeg("rplot.jpg")` : jpeg file
- `postscript("rplot.ps")`: postscript file
- `bmp("rplot.bmp")` : bmp file
- `win.metafile("rplot.wmf")` : windows metafile

Chapter 3

Standard graphical formatting in R

Introduction `plot()` function:

`plot()` function is generic function for plotting of R objects in basic graphs.

- `par()` : the default settings (rows x columns) for plots.
- `plot()` : the main function.
- There are many other plot functions which are specific to some tasks such as `hist()`, `boxplot()`, etc. Most of them take the same arguments as the `plot()` function.

Formula:

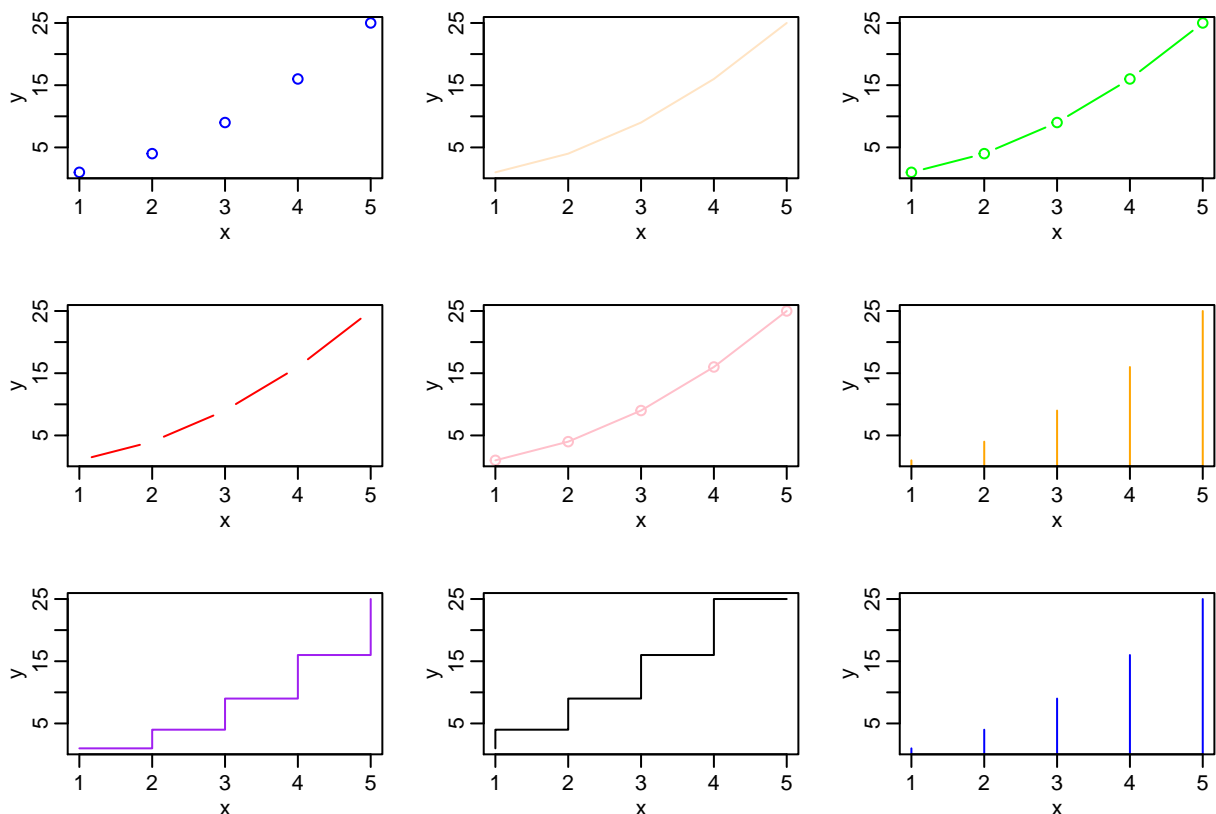
```
plot(x, y, type="p")
```

- **x** and **y**: the coordinates of points to plot
- **type**:
 - “p” for points,
 - “l” for lines,
 - “b” for both,
 - “c” for the lines part alone of “b”,
 - “o” for both ‘overplotted’,
 - “h” for ‘histogram’ like (or ‘high-density’) vertical lines,
 - “s” for stair steps,
 - “S” for other steps, see ‘Details’ below,
 - “n” for no plotting.

Example:

```
par(mfrow=c(3,3), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
x<-1:5; y=x*x
plot(x, y, type="p", col="blue")
plot(x, y, type="l", col="bisque")
plot(x,y, type="b", col="green")
plot(x, y, type="c", col="red")
plot(x, y, type="o", col="pink")
plot(x,y, type="h", col="orange")
```

```
plot(x, y, type="s",col="purple")
plot(x, y, type="S",col="black")
plot(x,y, type="h",col="blue")
```



For more color type used the sintaks `color()`

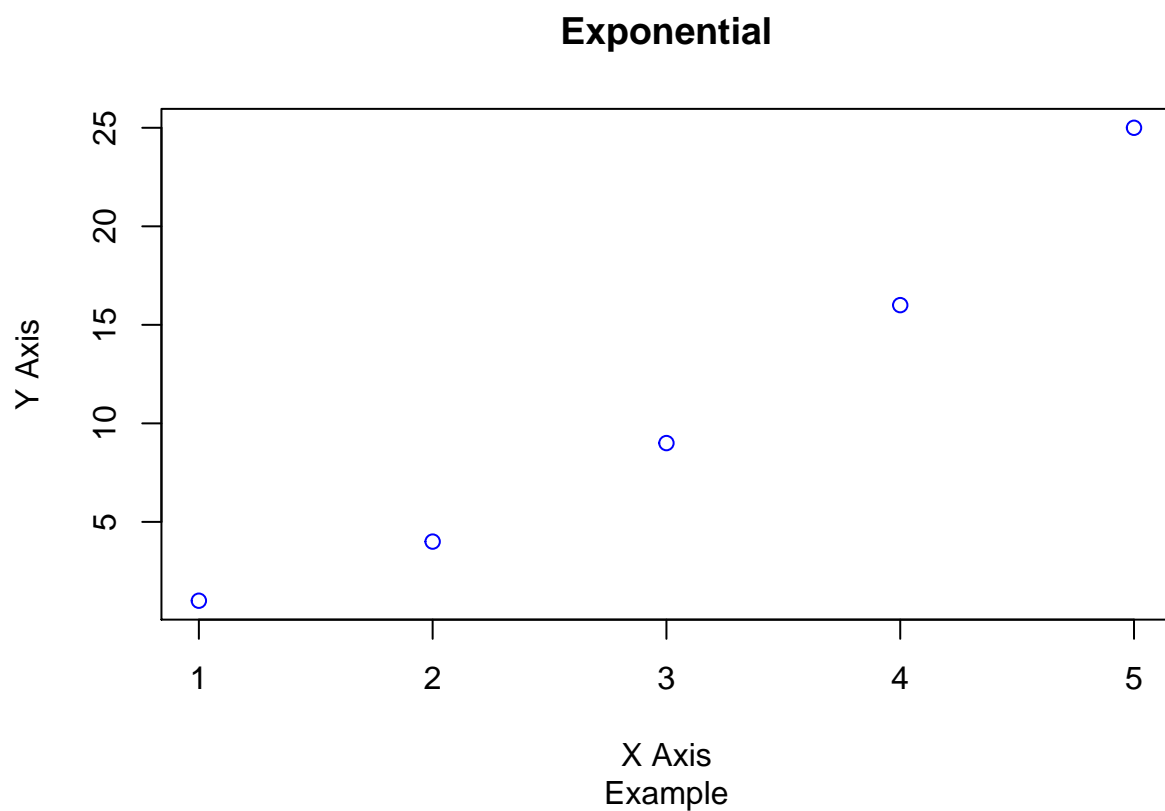
More info about `plot()` go to [RDocumentation](#) here

Titles:

- `main="test"`: main titles
- `sub="text"`: subtitle
- `xlab="test"`: the name of the x axis
- `ylab="test"`: the name of the y axis.

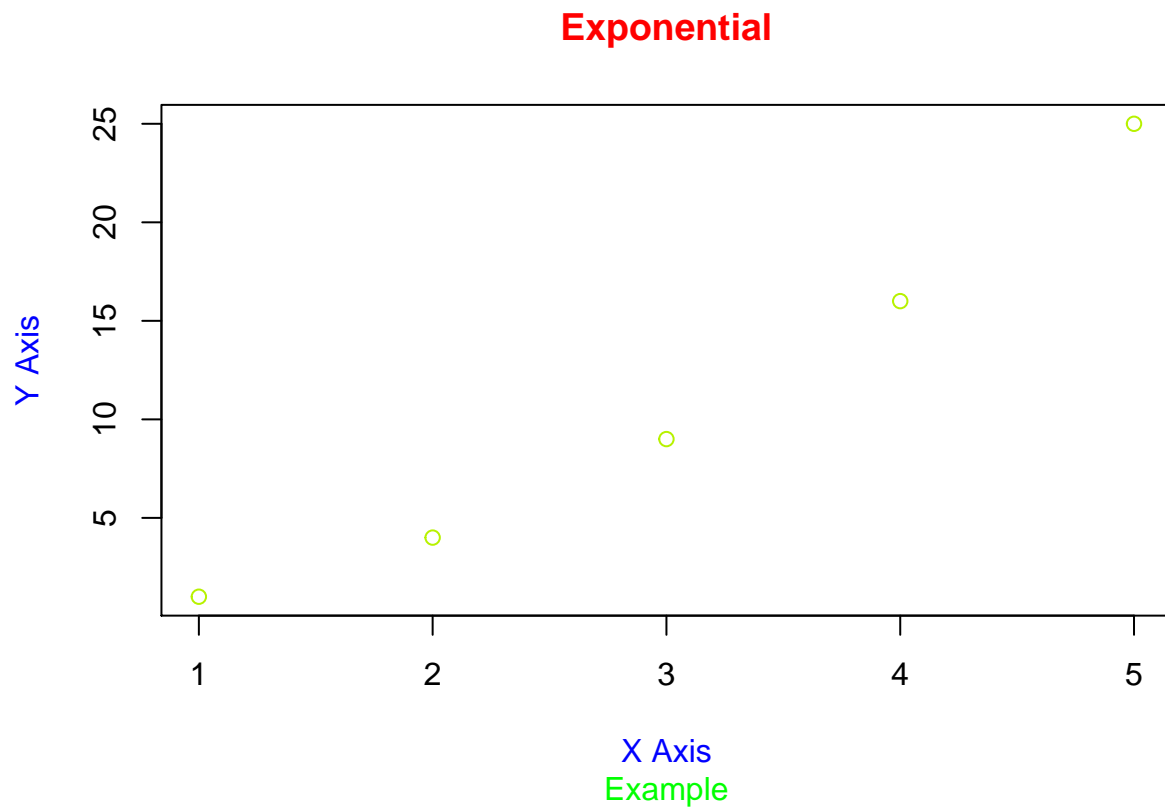
====Main title and axis labels=====

```
x<-1:5; y=x*x
plot(x, y, type="p",col="blue",main="Exponential",sub="Example",ylab="Y Axis",xlab="X Axis")
```



====Title colors====

```
x<-1:5; y=x*x
plot(x, y, type="p",col="#B7F200",main="Exponential",sub="Example",ylab="Y Axis",xlab="X Axis",
     col.main="red", col.lab="blue", col.sub="green")
```

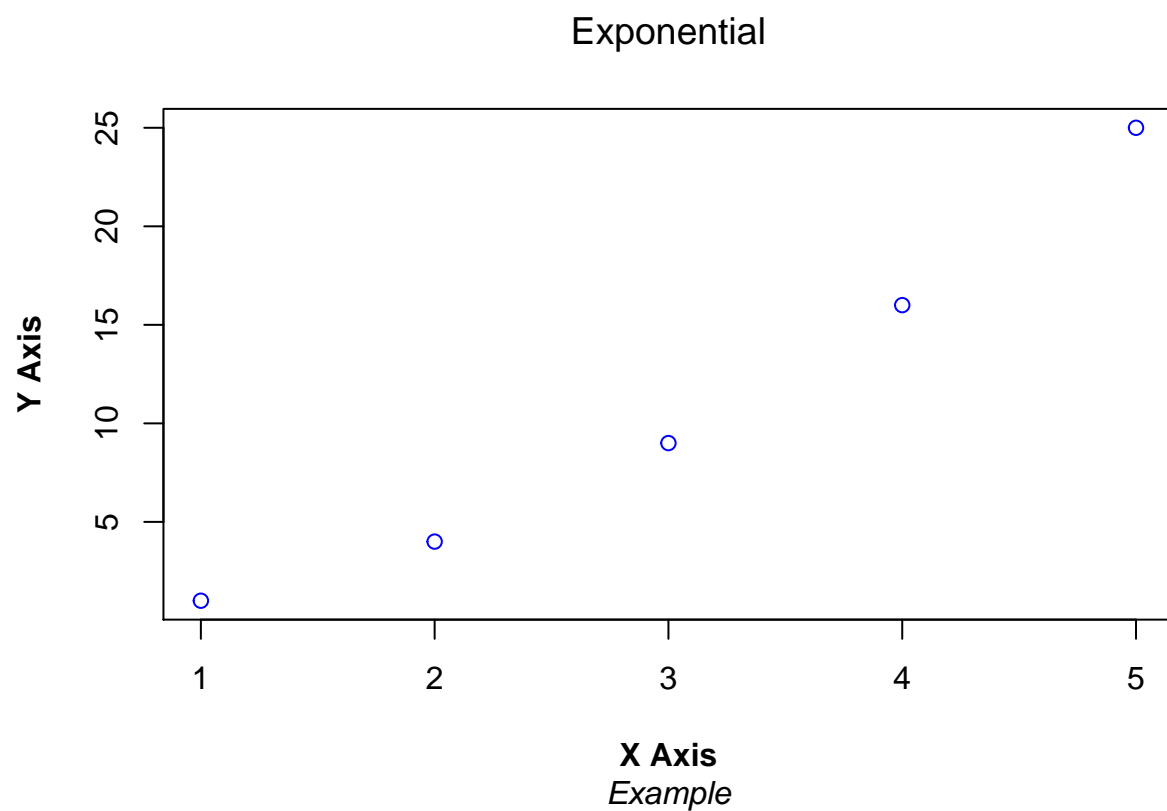


====The font style=====

The possible values for the font style are :

1: normal text 2: bold 3: italic 4: bold and italic 5 : Symbol font

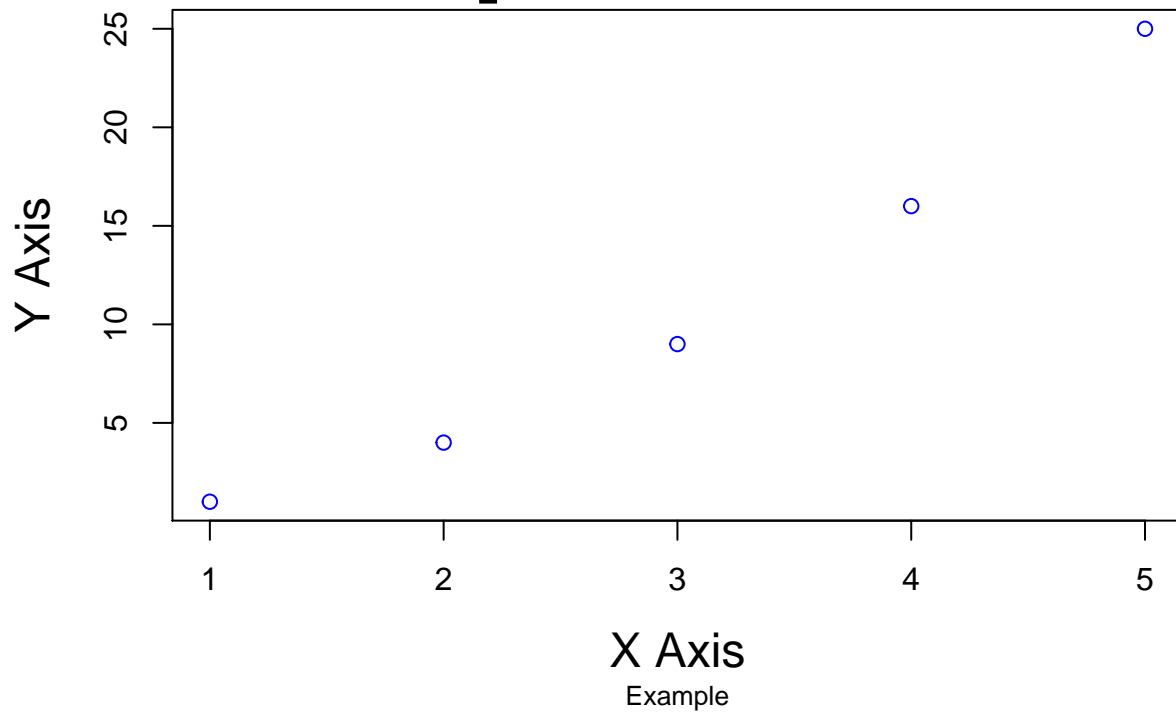
```
x<-1:5; y=x*x
plot(x, y, type="p", col="blue", main="Exponential", sub="Example", ylab="Y Axis", xlab="X Axis",
     font.main=1, font.lab=2, font.sub=3
)
```

====The font size====

```
x<-1:5; y=x*x
plot(x, y, type="p", col="blue", main="Exponential", sub="Example", ylab="Y Axis", xlab="X Axis",
      cex.main=4, cex.lab=1.5, cex.sub=0.8
)
```

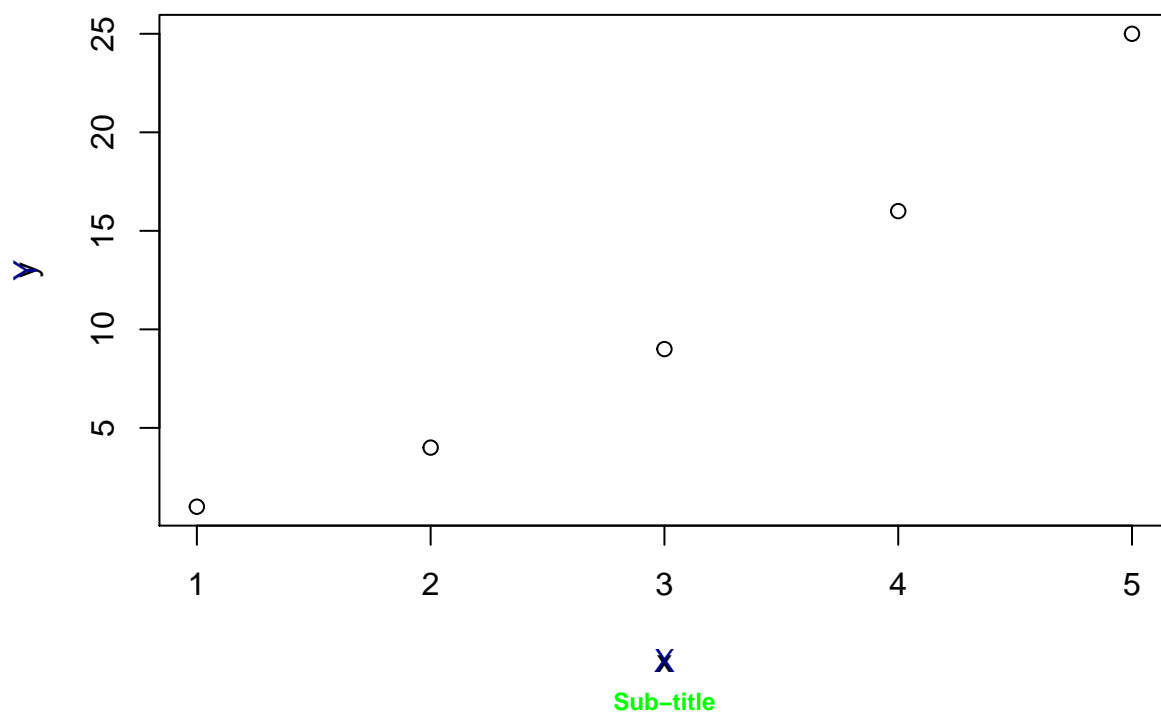
Exponential



====Use the title() function====

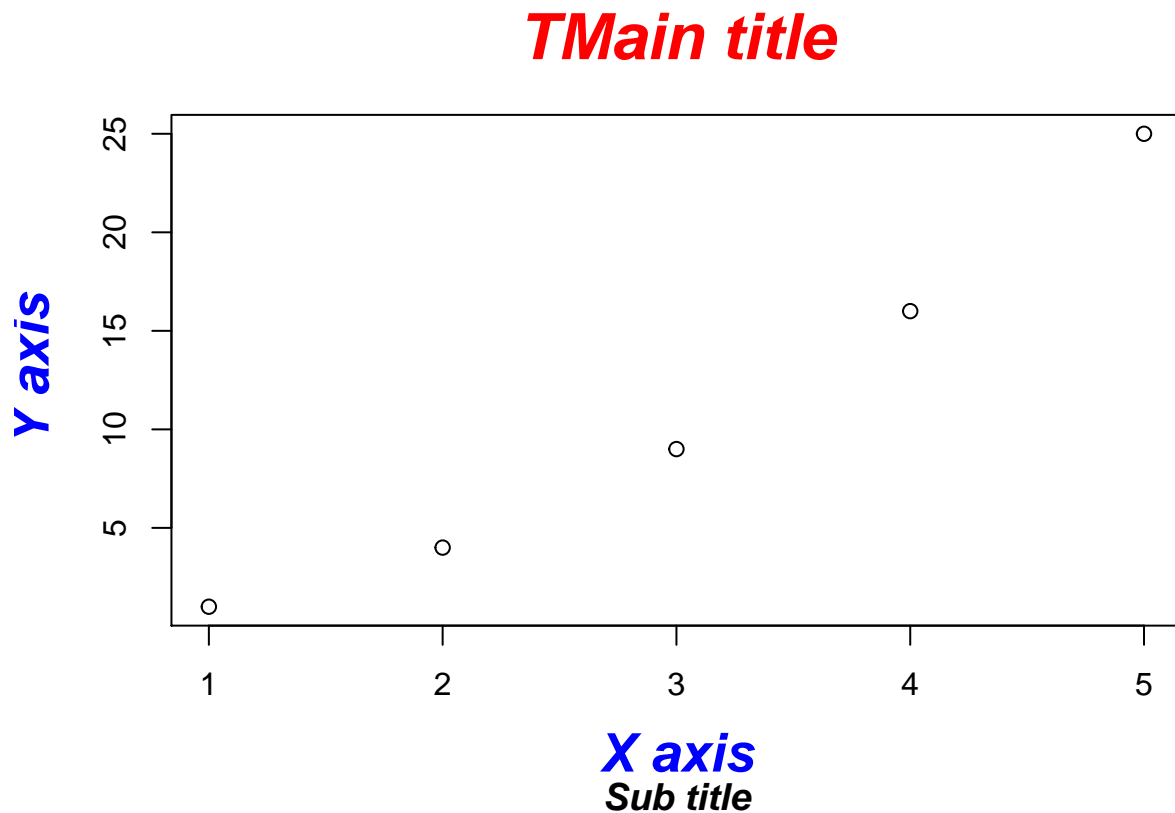
```
x<-1:5; y=x*x
plot(x, y, type="p")
title(main = "Main title", sub = "Sub-title",
      xlab = "X", ylab = "Y",
      cex.main = 2, font.main= 4, col.main= "red",
      cex.sub = 0.75, font.sub = 2, col.sub = "green",
      col.lab ="darkblue"
)
```

Main title



====Customize the titles using `par()` function=====

```
par(
  # Change the colors
  col.main="red", col.lab="blue", col.sub="black",
  # Titles in italic and bold
  font.main=4, font.lab=4, font.sub=4,
  # Change font size
  cex.main=2, cex.lab=1.7, cex.sub=1.2
)
plot(x, y, type="p",
     main="Main title",
     xlab="X axis",
     ylab="Y axis",
     sub="Sub title"
)
```

**Legend:**

`legend()`: the position can be “bottomleft”, “bottomright”, “topleft”, “topright” or exact coordinates.

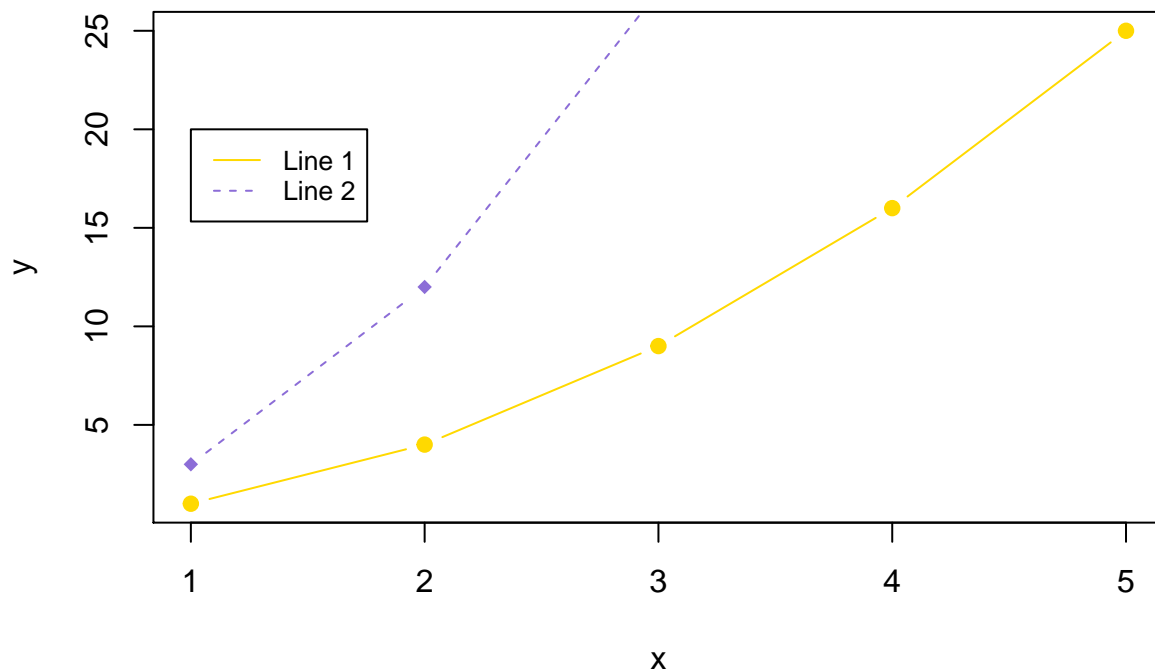
====R legend function=====

Formula:

```
legend(x, y=NULL, legend, fill, col, bg)
```

Example:

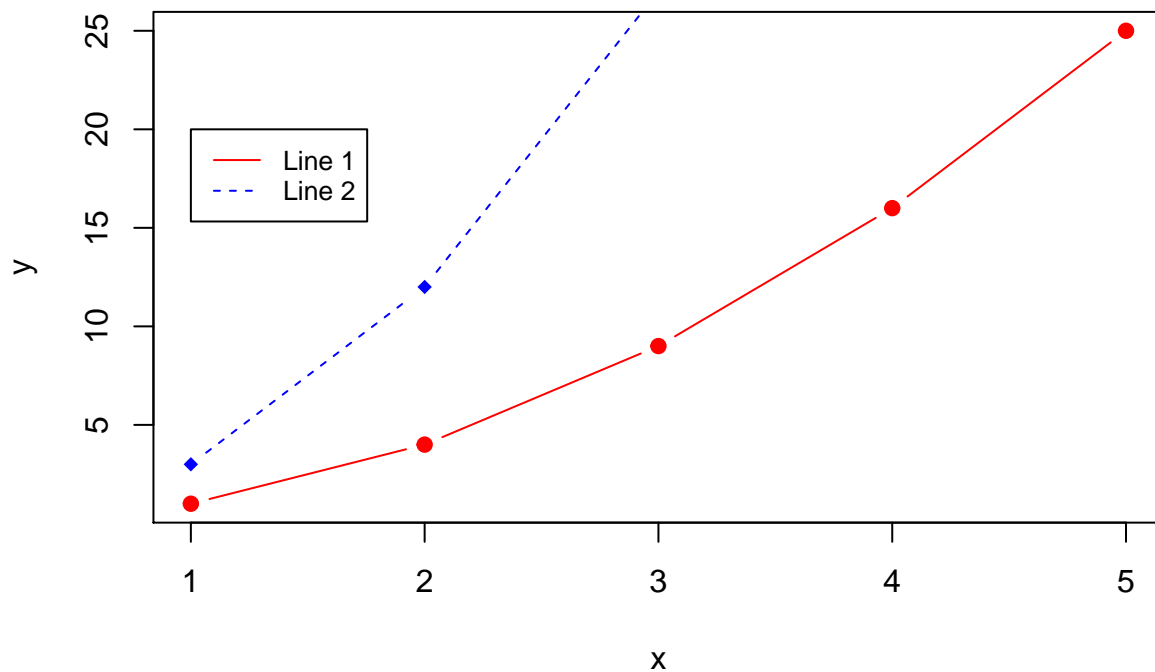
```
# Generate example data
x<-1:5; y1=x*x; y2=3*y1
plot(x, y1, type="b", pch=19, col="#FFD900", xlab="x", ylab="y")
# Add a line
lines(x, y2, pch=18, col="#8C6DD7", type="b", lty=2)
# Add a legend
legend(1, 20, legend=c("Line 1", "Line 2"),
      col=c("#FFD900", "#8C6DD7"), lty=1:2, cex=0.8)
```



Create R function to avoid repeating graphs code:

```
# Generate example data
make_plot<-function(){
x<-1:5; y1=x*x; y2=3*y1
plot(x, y1, type="b", pch=19, col="red", xlab="x", ylab="y")
# Add a line
lines(x, y2, pch=18, col="blue", type="b", lty=2)
# Add a legend
legend(1, 20, legend=c("Line 1", "Line 2"),
      col=c("red", "blue"), lty=1:2, cex=0.8)
}
```

```
make_plot()
```

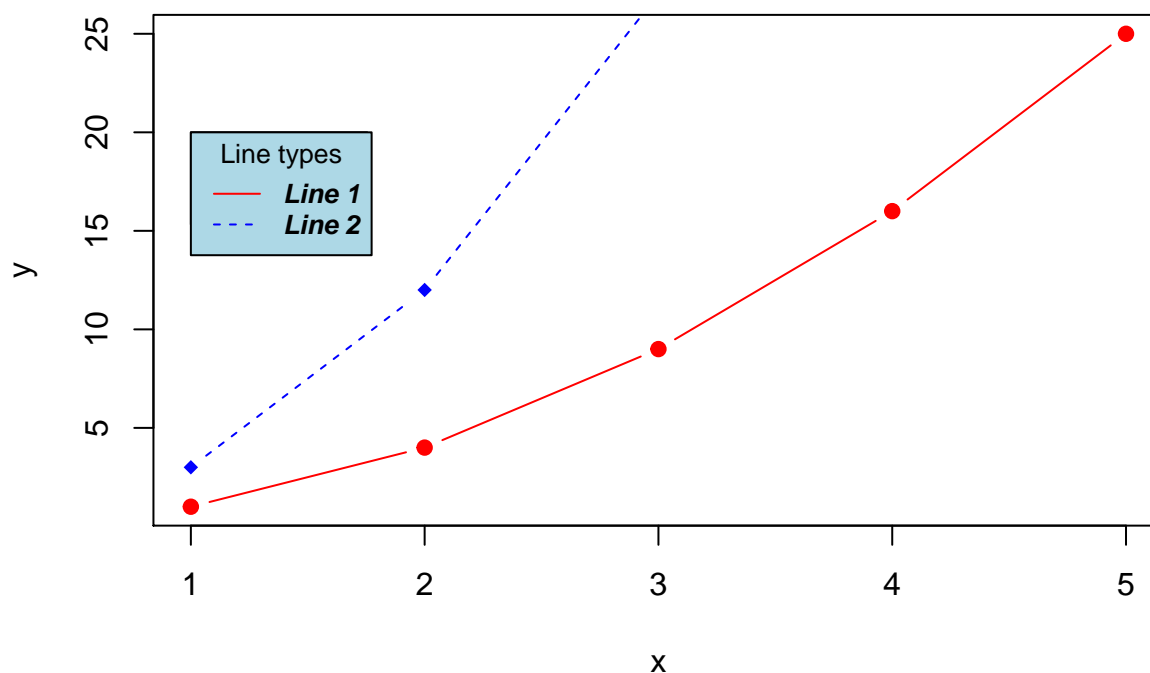


====Title, text font and background color of the legend box=====

- title: The title of the legend
- text.font: an integer specifying the font style of the legend text; possible values are :
 - 1: normal
 - 2: bold
 - 3: italic
 - 4: bold and italic
 - bg: background color of the legend box

```
make_plot()

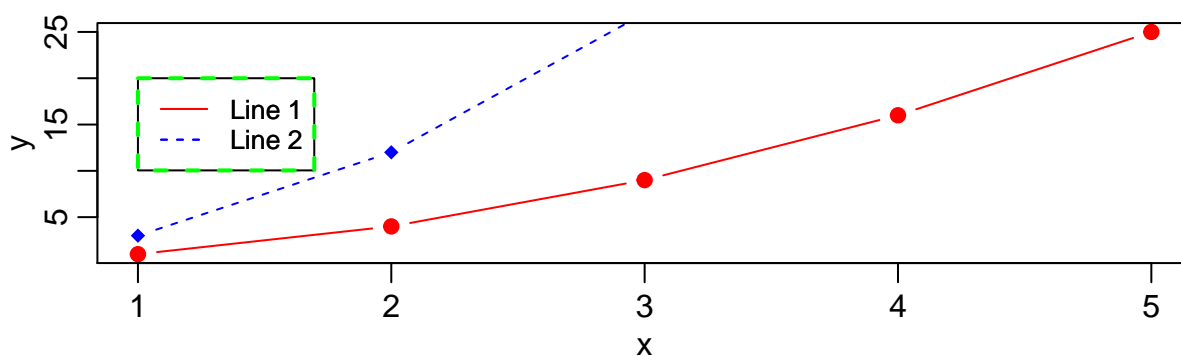
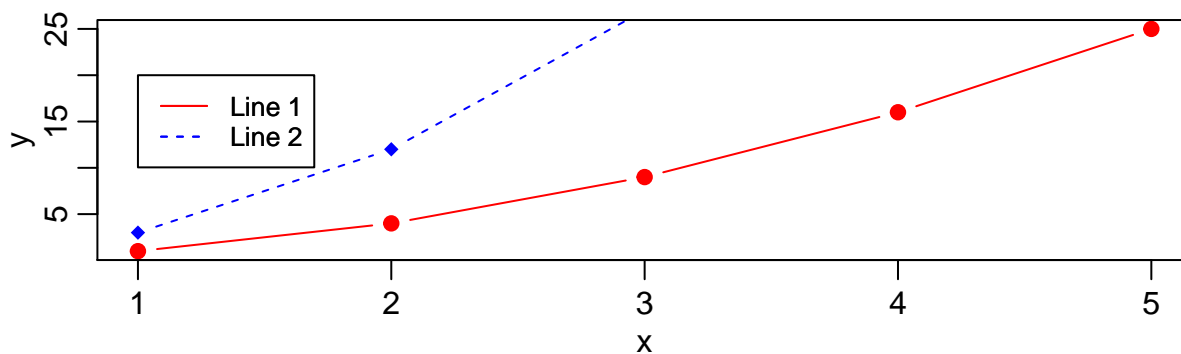
# Add a legend to the plot
legend(1,20, legend=c("Line 1", "Line 2"),
      col=c("red", "blue"), lty=1:2, cex=0.8,
      title="Line types", text.font=4, bg="lightblue")
```



====Border of the legend box=====

- box.lty: modify the line type
- box.lwd: modify the width
- box.col: modify the color

```
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
# Remove legend border using box.lty = 0
make_plot()
legend(1,20, legend=c("Line 1", "Line 2"),
      col=c("red", "blue"), lty=1:2, cex=0.8,
      box.lty=0)
# Change the border
make_plot()
legend(1,20, legend=c("Line 1", "Line 2"),
      col=c("red", "blue"), lty=1:2, cex=0.8,
      box.lty=2, box.lwd=2, box.col="green")
```



====The legend position by keywords====

Using the following keywords : “bottomright”, “bottom”, “bottomleft”, “left”, “topleft”, “top”, “topright”, “right” and “center”.

```
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
# Remove legend border using box.lty = 0
make_plot()
legend("center", legend=c("Line 1", "Line 2"),
      col=c("red", "blue"), box.lty=0)
make_plot()
legend("topright", legend=c("Line 1", "Line 2"),
      col=c("red", "blue"),
      box.lty=0)
```

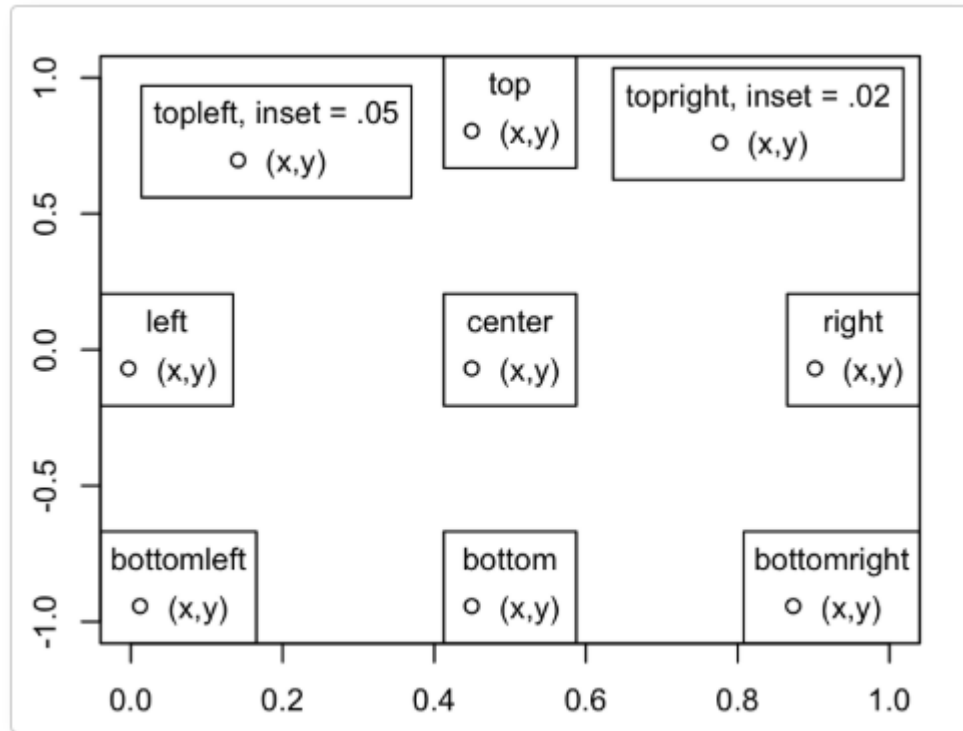
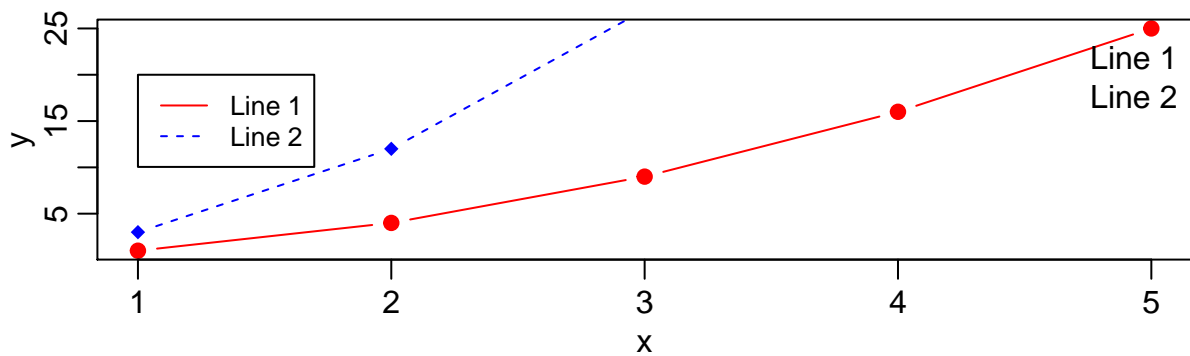
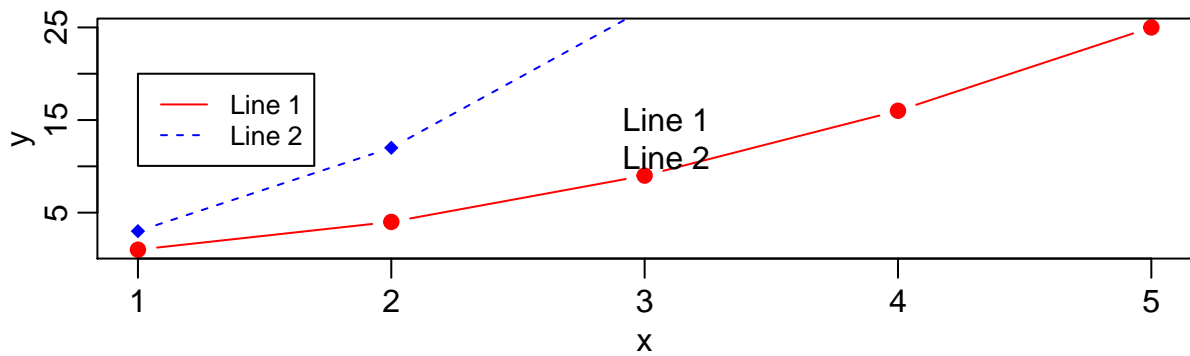



Figure 3.1:



Text:

`text()` and `mtext()` R functions can be used To add a text to a plot.

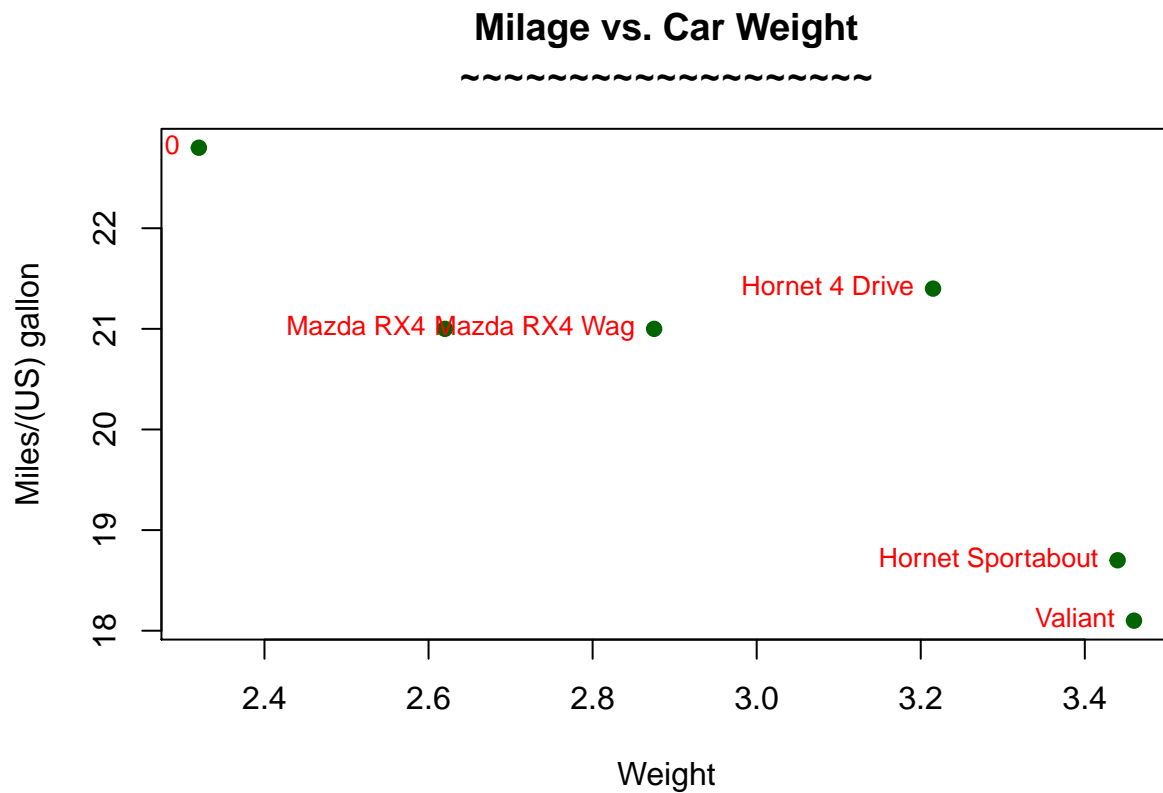
====Texts within the graph=====

Formula:

```
text(x, y, labels)
```

Example:

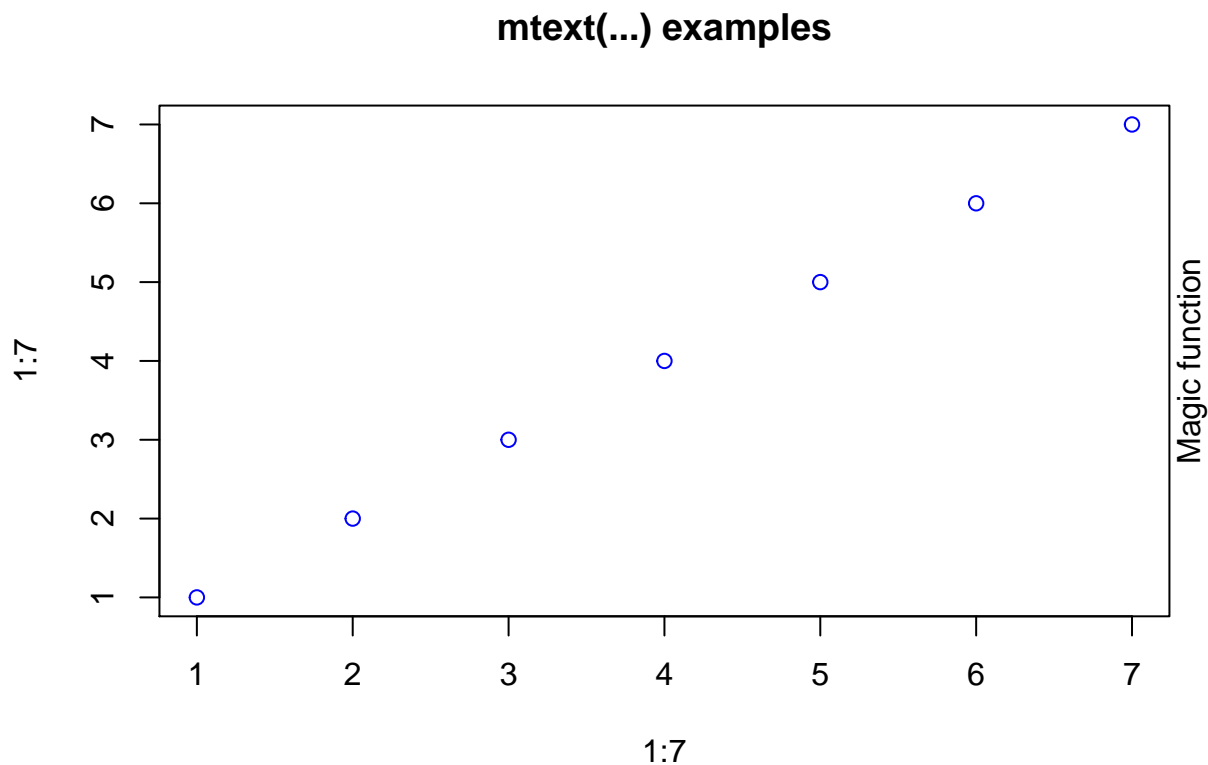
```
d<-head(mtcars)
plot(d[, 'wt'], d[, 'mpg'],
     main="Milage vs. Car Weight\n~~~~~",
     xlab="Weight", ylab="Miles/(US) gallon",
     pch=19, col="darkgreen")
text(d[, 'wt'], d[, 'mpg'], row.names(d),
     cex=0.85, pos=2, col="red")
```



====Text in the margins of the graph=====

- text : the text to be written
- side : an integer specifying the side of the plot; Possible values are :
 - 1: bottom
 - 2: left
 - 3: top
 - 4: right

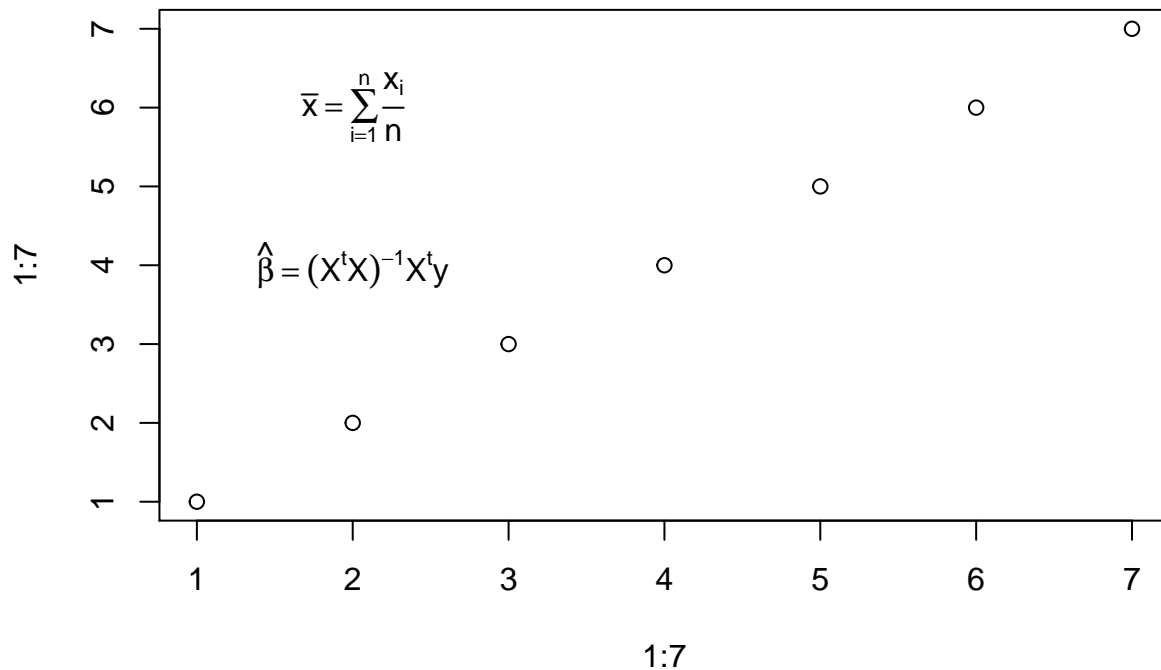
```
plot(1:7, 1:7,
     main="mtext(...) examples",col="blue")
mtext("Magic function", side=4)
```



====Mathematical annotation within the graph=====

Example:

```
plot(1:7, 1:7,
     main="text(...) examples",col="black")
text(2, 4, expression(hat(beta) == (X^t * X)^{-1} * X^t * y))
text(2, 6, expression(bar(x) == sum(frac(x[i], n), i==1, n)))
```

text(...) examples**Types of Plot:**

An option for plot types can be:

type: * "p" for points,

* "l" for lines,

* "b" for both,

* "c" for the lines part alone of "b",

* "o" for both 'overplotted',

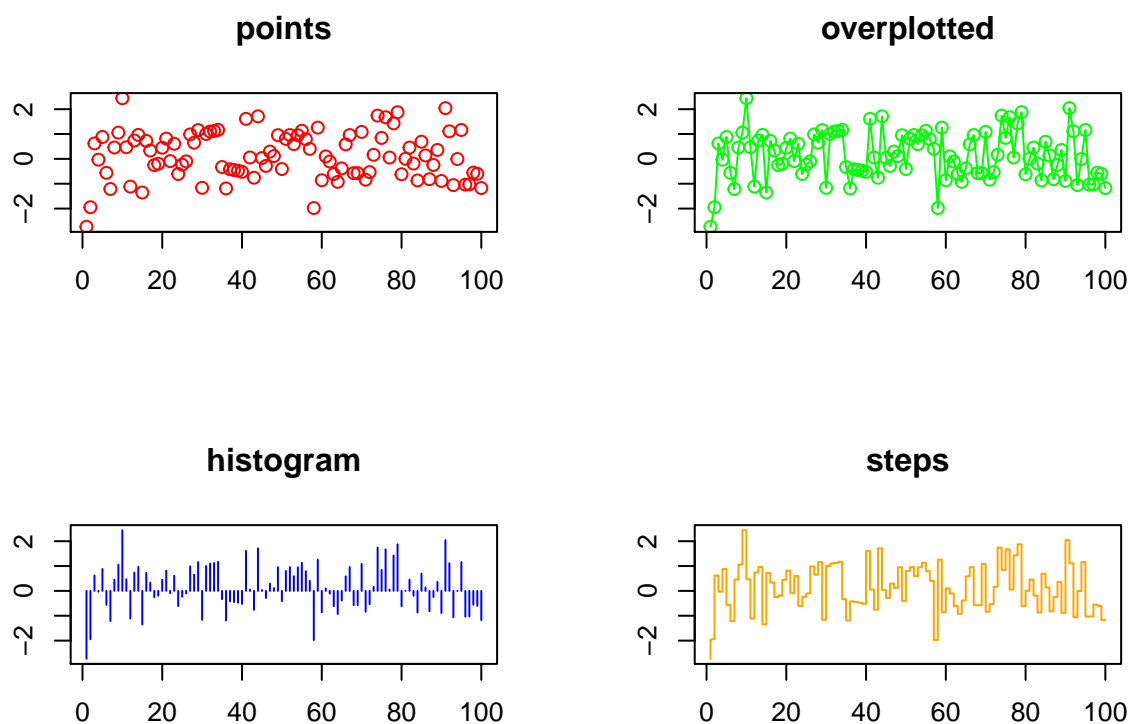
* "h" for 'histogram' like (or 'high-density') vertical lines,

* "s" for stair steps,

* "S" for other steps, see 'Details' below,

* "n" for no plotting.

```
x <- rnorm(100)
par(mfrow = c(2,2))
plot(x, type = "p", main = "points", ylab = "", xlab = "", col="red")
plot(x, type = "o", main = "overplotted", ylab = "", xlab = "", col="green")
plot(x, type = "h", main = "histogram", ylab = "", xlab = "", col="blue")
plot(x, type = "S", main = "steps", ylab = "", xlab = "", col="orange")
```



Axis:

====Add an axis to a plot=====

Formula:

```
axis(side, at=NULL, labels=TRUE)
```

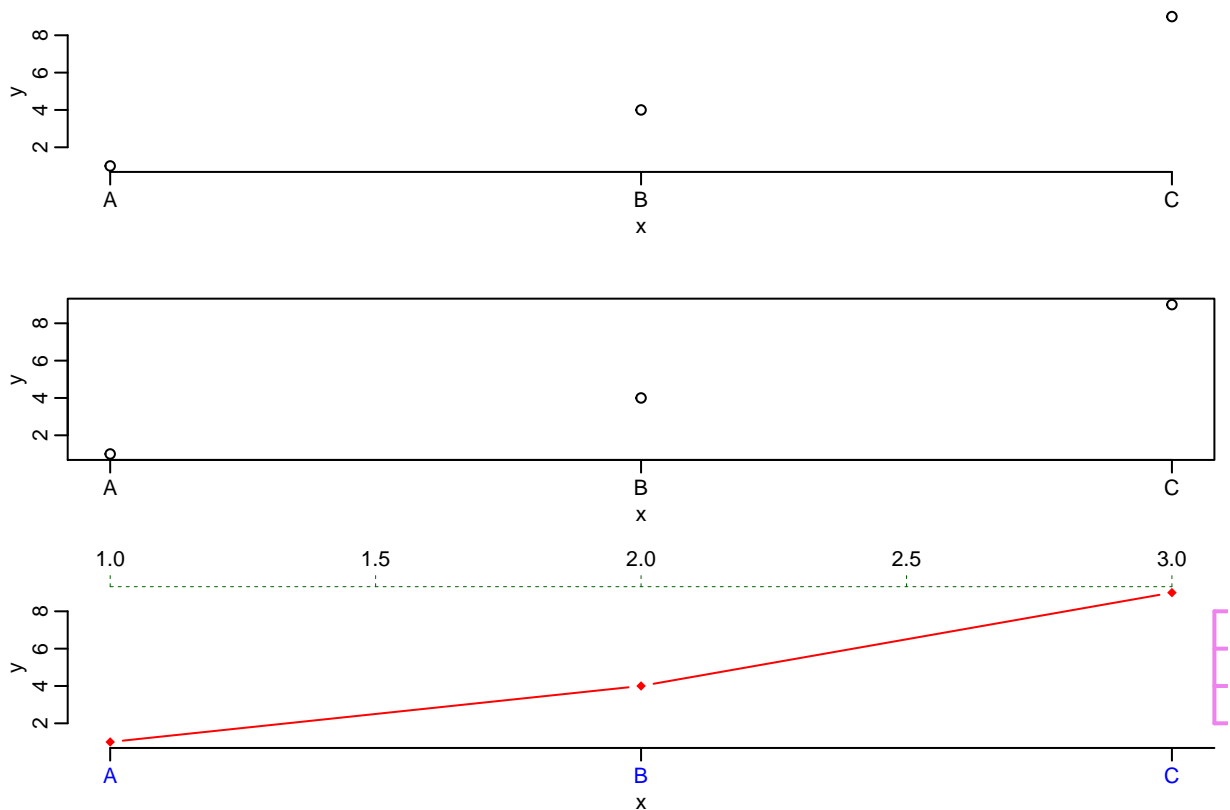
Side :

- 1: below
- 2: left
- 3: above
- 4: right
- at: The position
- labels: Texts for tick-mark labels.

Example :

```
par(mfrow=c(3,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
x<-1:3; y=x*x
# Example 1
plot(x, y, axes = FALSE)
axis(side=1, at = 1:3, labels=LETTERS[1:3])
axis(2)
# Example 2
plot(x, y, axes = FALSE)
```

```
axis(side=1, at=1:3, labels=LETTERS[1:3])
axis(2)
box() #- To make it look like "usual" plot
#Example3:
plot(x, y, pch=18, col="red", type="b",
     frame=FALSE, xaxt="n") # Remove x axis
axis(1, 1:4, LETTERS[1:4], col.axis="blue")
axis(3, col = "darkgreen", lty = 2, lwd = 0.5)
axis(4, col = "violet", col.axis = "dark violet", lwd = 2)
```



====Axis scale=====

- xlim: the limit of x axis; format : xlim = c(min, max)
- ylim: the limit of y axis; format: ylim = c(min, max)

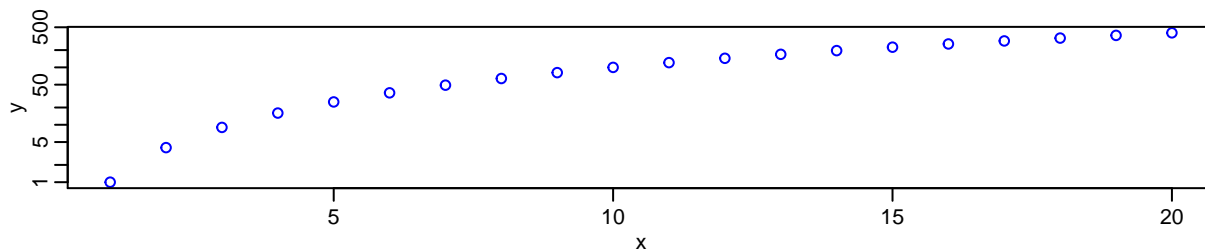
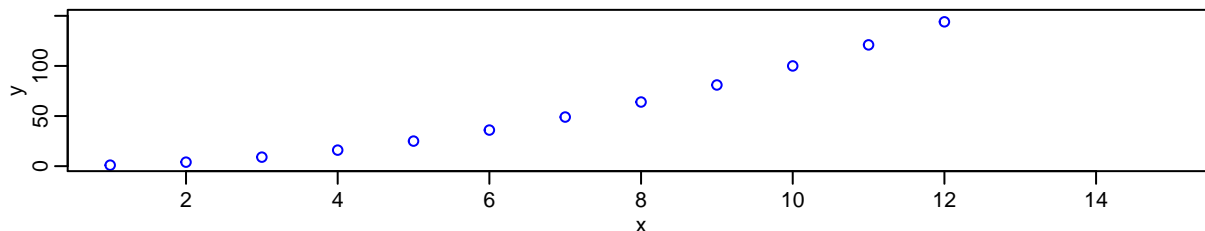
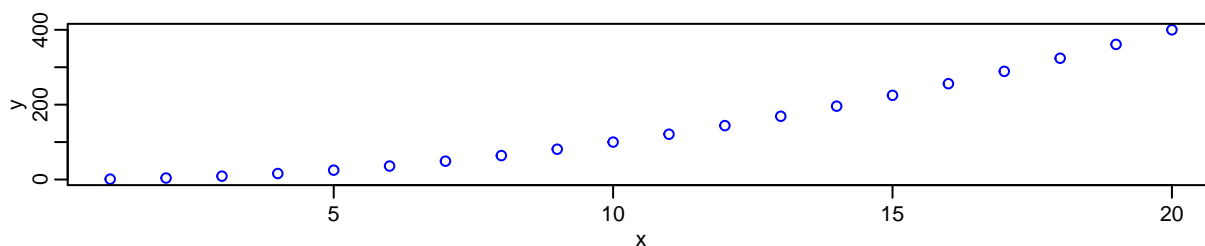
Transformation to log scale:

- log = "x"
- log = "y"
- log = "xy"*

Example:

```
par(mfrow=c(3,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
x<-1:20; y=x*x
# Simple graph
plot(x, y,col="blue")
```

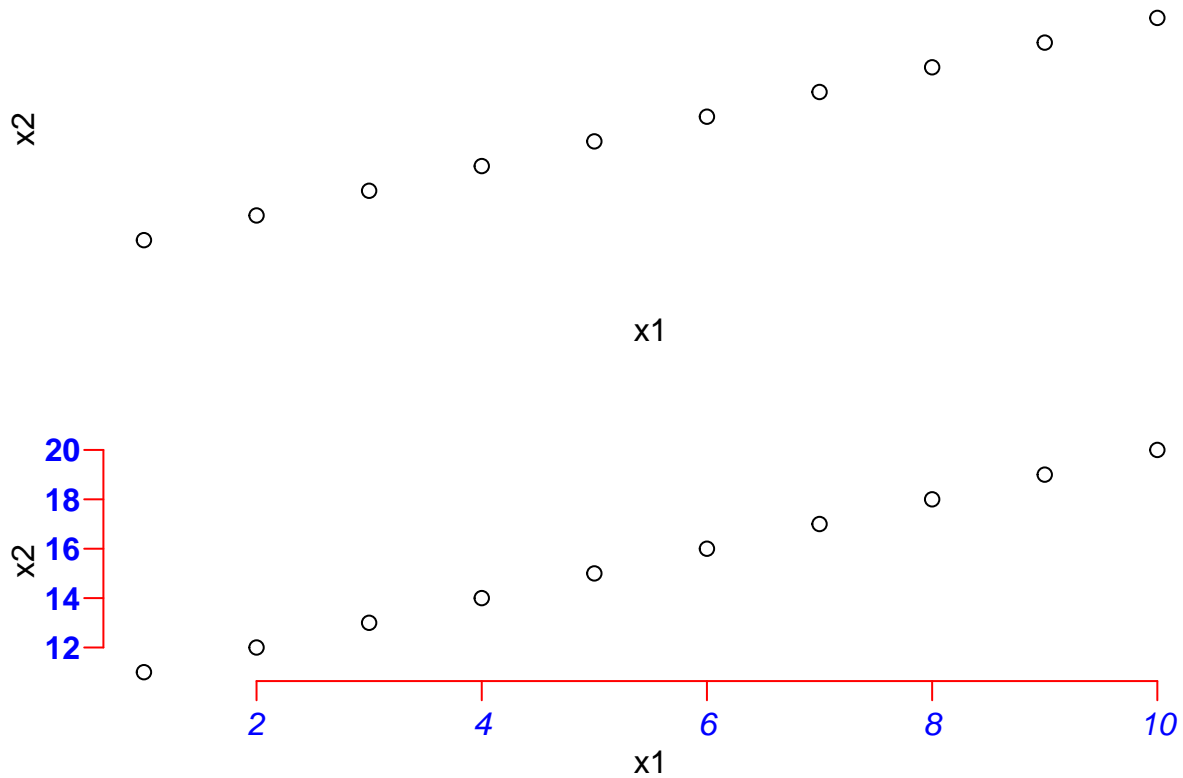
```
# Enlarge the scale
plot(x, y, xlim=c(1,15), ylim=c(1,150),col="blue")
# Log scale
plot(x, y, log="y",col="blue")
```



====Axes fonts=====

Remove them with `axes=FALSE`:

```
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
x1<-1:10
x2<-11:20
plot(x1,x2,axes=FALSE)
plot(x1,x2,axes=FALSE)
axis(1,col="red",col.axis="blue",font.axis=3)
axis(2,col="red",col.axis="blue",font.axis=2,las=2)
```

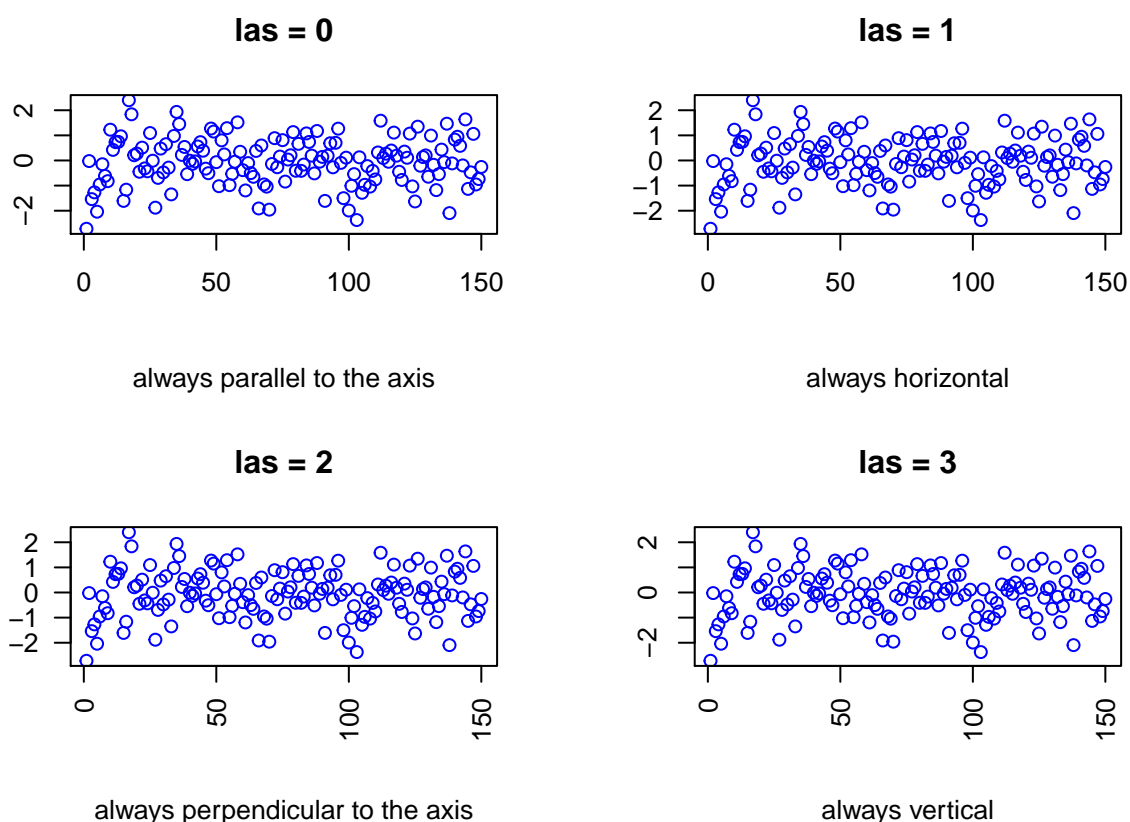


====Axes positions=====

Use `las` function to specified the style of axis labels:

- 0 : always parallel to the axis [default]
- 1 : always horizontal
- 2 : always perpendicular to the axis
- 3 : always vertical

```
x1 <- rnorm(150)
par(mfrow = c(2,2))
plot(x1, las = 0, main = "las = 0", sub = "always parallel to the axis", xlab = "", ylab = "",col="blue")
plot(x1, las = 1, main = "las = 1", sub = "always horizontal", xlab = "", ylab = "",col="blue")
plot(x1, las = 2, main = "las = 2", sub = "always perpendicular to the axis", xlab = "", ylab = "",col="blue")
plot(x1, las = 3, main = "las = 3", sub = "always vertical", xlab = "", ylab = "",col="blue")
```

Margins `par()`:

Formula:

```
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
```

See `?par` to learn more about the topic.

Colors:

Use by name (e.g `col = "red"`) or as a hexadecimal RGB triplet (such as `col = "#FFCC00"`).

====Default color in R====

```
colors() # list the r colors
```

## [1] "white"	"aliceblue"	"antiquewhite"
## [4] "antiquewhite1"	"antiquewhite2"	"antiquewhite3"
## [7] "antiquewhite4"	"aquamarine"	"aquamarine1"
## [10] "aquamarine2"	"aquamarine3"	"aquamarine4"
## [13] "azure"	"azure1"	"azure2"
## [16] "azure3"	"azure4"	"beige"
## [19] "bisque"	"bisque1"	"bisque2"
## [22] "bisque3"	"bisque4"	"black"
## [25] "blanchedalmond"	"blue"	"blue1"
## [28] "blue2"	"blue3"	"blue4"
## [31] "blueviolet"	"brown"	"brown1"
## [34] "brown2"	"brown3"	"brown4"
## [37] "burlywood"	"burlywood1"	"burlywood2"
## [40] "burlywood3"	"burlywood4"	"cadetblue"

## [43]	"cadetblue1"	"cadetblue2"	"cadetblue3"
## [46]	"cadetblue4"	"chartreuse"	"chartreuse1"
## [49]	"chartreuse2"	"chartreuse3"	"chartreuse4"
## [52]	"chocolate"	"chocolate1"	"chocolate2"
## [55]	"chocolate3"	"chocolate4"	"coral"
## [58]	"coral1"	"coral2"	"coral3"
## [61]	"coral4"	"cornflowerblue"	"cornsilk"
## [64]	"cornsilk1"	"cornsilk2"	"cornsilk3"
## [67]	"cornsilk4"	"cyan"	"cyan1"
## [70]	"cyan2"	"cyan3"	"cyan4"
## [73]	"darkblue"	"darkcyan"	"darkgoldenrod"
## [76]	"darkgoldenrod1"	"darkgoldenrod2"	"darkgoldenrod3"
## [79]	"darkgoldenrod4"	"darkgray"	"darkgreen"
## [82]	"darkgrey"	"darkkhaki"	"darkmagenta"
## [85]	"darkolivegreen"	"darkolivegreen1"	"darkolivegreen2"
## [88]	"darkolivegreen3"	"darkolivegreen4"	"darkorange"
## [91]	"darkorange1"	"darkorange2"	"darkorange3"
## [94]	"darkorange4"	"darkorchid"	"darkorchid1"
## [97]	"darkorchid2"	"darkorchid3"	"darkorchid4"
## [100]	"darkred"	"darksalmon"	"darkseagreen"
## [103]	"darkseagreen1"	"darkseagreen2"	"darkseagreen3"
## [106]	"darkseagreen4"	"darkslateblue"	"darkslategray"
## [109]	"darkslategray1"	"darkslategray2"	"darkslategray3"
## [112]	"darkslategray4"	"darkslategrey"	"darkturquoise"
## [115]	"darkviolet"	"deeppink"	"deeppink1"
## [118]	"deeppink2"	"deeppink3"	"deeppink4"
## [121]	"deepskyblue"	"deepskyblue1"	"deepskyblue2"
## [124]	"deepskyblue3"	"deepskyblue4"	"dimgray"
## [127]	"dimgrey"	"dodgerblue"	"dodgerblue1"
## [130]	"dodgerblue2"	"dodgerblue3"	"dodgerblue4"
## [133]	"firebrick"	"firebrick1"	"firebrick2"
## [136]	"firebrick3"	"firebrick4"	"floralwhite"
## [139]	"forestgreen"	"gainsboro"	"ghostwhite"
## [142]	"gold"	"gold1"	"gold2"
## [145]	"gold3"	"gold4"	"goldenrod"
## [148]	"goldenrod1"	"goldenrod2"	"goldenrod3"
## [151]	"goldenrod4"	"gray"	"gray0"
## [154]	"gray1"	"gray2"	"gray3"
## [157]	"gray4"	"gray5"	"gray6"
## [160]	"gray7"	"gray8"	"gray9"
## [163]	"gray10"	"gray11"	"gray12"
## [166]	"gray13"	"gray14"	"gray15"
## [169]	"gray16"	"gray17"	"gray18"
## [172]	"gray19"	"gray20"	"gray21"
## [175]	"gray22"	"gray23"	"gray24"
## [178]	"gray25"	"gray26"	"gray27"
## [181]	"gray28"	"gray29"	"gray30"
## [184]	"gray31"	"gray32"	"gray33"
## [187]	"gray34"	"gray35"	"gray36"
## [190]	"gray37"	"gray38"	"gray39"
## [193]	"gray40"	"gray41"	"gray42"
## [196]	"gray43"	"gray44"	"gray45"
## [199]	"gray46"	"gray47"	"gray48"
## [202]	"gray49"	"gray50"	"gray51"

## [205]	"gray52"	"gray53"	"gray54"
## [208]	"gray55"	"gray56"	"gray57"
## [211]	"gray58"	"gray59"	"gray60"
## [214]	"gray61"	"gray62"	"gray63"
## [217]	"gray64"	"gray65"	"gray66"
## [220]	"gray67"	"gray68"	"gray69"
## [223]	"gray70"	"gray71"	"gray72"
## [226]	"gray73"	"gray74"	"gray75"
## [229]	"gray76"	"gray77"	"gray78"
## [232]	"gray79"	"gray80"	"gray81"
## [235]	"gray82"	"gray83"	"gray84"
## [238]	"gray85"	"gray86"	"gray87"
## [241]	"gray88"	"gray89"	"gray90"
## [244]	"gray91"	"gray92"	"gray93"
## [247]	"gray94"	"gray95"	"gray96"
## [250]	"gray97"	"gray98"	"gray99"
## [253]	"gray100"	"green"	"green1"
## [256]	"green2"	"green3"	"green4"
## [259]	"greenyellow"	"grey"	"grey0"
## [262]	"grey1"	"grey2"	"grey3"
## [265]	"grey4"	"grey5"	"grey6"
## [268]	"grey7"	"grey8"	"grey9"
## [271]	"grey10"	"grey11"	"grey12"
## [274]	"grey13"	"grey14"	"grey15"
## [277]	"grey16"	"grey17"	"grey18"
## [280]	"grey19"	"grey20"	"grey21"
## [283]	"grey22"	"grey23"	"grey24"
## [286]	"grey25"	"grey26"	"grey27"
## [289]	"grey28"	"grey29"	"grey30"
## [292]	"grey31"	"grey32"	"grey33"
## [295]	"grey34"	"grey35"	"grey36"
## [298]	"grey37"	"grey38"	"grey39"
## [301]	"grey40"	"grey41"	"grey42"
## [304]	"grey43"	"grey44"	"grey45"
## [307]	"grey46"	"grey47"	"grey48"
## [310]	"grey49"	"grey50"	"grey51"
## [313]	"grey52"	"grey53"	"grey54"
## [316]	"grey55"	"grey56"	"grey57"
## [319]	"grey58"	"grey59"	"grey60"
## [322]	"grey61"	"grey62"	"grey63"
## [325]	"grey64"	"grey65"	"grey66"
## [328]	"grey67"	"grey68"	"grey69"
## [331]	"grey70"	"grey71"	"grey72"
## [334]	"grey73"	"grey74"	"grey75"
## [337]	"grey76"	"grey77"	"grey78"
## [340]	"grey79"	"grey80"	"grey81"
## [343]	"grey82"	"grey83"	"grey84"
## [346]	"grey85"	"grey86"	"grey87"
## [349]	"grey88"	"grey89"	"grey90"
## [352]	"grey91"	"grey92"	"grey93"
## [355]	"grey94"	"grey95"	"grey96"
## [358]	"grey97"	"grey98"	"grey99"
## [361]	"grey100"	"honeydew"	"honeydew1"
## [364]	"honeydew2"	"honeydew3"	"honeydew4"

## [367] "hotpink"	"hotpink1"	"hotpink2"
## [370] "hotpink3"	"hotpink4"	"indianred"
## [373] "indianred1"	"indianred2"	"indianred3"
## [376] "indianred4"	"ivory"	"ivory1"
## [379] "ivory2"	"ivory3"	"ivory4"
## [382] "khaki"	"khaki1"	"khaki2"
## [385] "khaki3"	"khaki4"	"lavender"
## [388] "lavenderblush"	"lavenderblush1"	"lavenderblush2"
## [391] "lavenderblush3"	"lavenderblush4"	"lawngreen"
## [394] "lemonchiffon"	"lemonchiffon1"	"lemonchiffon2"
## [397] "lemonchiffon3"	"lemonchiffon4"	"lightblue"
## [400] "lightblue1"	"lightblue2"	"lightblue3"
## [403] "lightblue4"	"lightcoral"	"lightcyan"
## [406] "lightcyan1"	"lightcyan2"	"lightcyan3"
## [409] "lightcyan4"	"lightgoldenrod"	"lightgoldenrod1"
## [412] "lightgoldenrod2"	"lightgoldenrod3"	"lightgoldenrod4"
## [415] "lightgoldenrodyellow"	"lightgray"	"lightgreen"
## [418] "lightgrey"	"lightpink"	"lightpink1"
## [421] "lightpink2"	"lightpink3"	"lightpink4"
## [424] "lightsalmon"	"lightsalmon1"	"lightsalmon2"
## [427] "lightsalmon3"	"lightsalmon4"	"lightseagreen"
## [430] "lightskyblue"	"lightskyblue1"	"lightskyblue2"
## [433] "lightskyblue3"	"lightskyblue4"	"lightslateblue"
## [436] "lightslategray"	"lightslategrey"	"lightsteelblue"
## [439] "lightsteelblue1"	"lightsteelblue2"	"lightsteelblue3"
## [442] "lightsteelblue4"	"lightyellow"	"lightyellow1"
## [445] "lightyellow2"	"lightyellow3"	"lightyellow4"
## [448] "limegreen"	"linen"	"magenta"
## [451] "magenta1"	"magenta2"	"magenta3"
## [454] "magenta4"	"maroon"	"maroon1"
## [457] "maroon2"	"maroon3"	"maroon4"
## [460] "mediumaquamarine"	"mediumblue"	"mediumorchid"
## [463] "mediumorchid1"	"mediumorchid2"	"mediumorchid3"
## [466] "mediumorchid4"	"mediumpurple"	"mediumpurple1"
## [469] "mediumpurple2"	"mediumpurple3"	"mediumpurple4"
## [472] "mediumseagreen"	"mediumslateblue"	"mediumspringgreen"
## [475] "mediumturquoise"	"mediumvioletred"	"midnightblue"
## [478] "mintcream"	"mistyrose"	"mistyrose1"
## [481] "mistyrose2"	"mistyrose3"	"mistyrose4"
## [484] "moccasin"	"navajowhite"	"navajowhite1"
## [487] "navajowhite2"	"navajowhite3"	"navajowhite4"
## [490] "navy"	"navyblue"	"oldlace"
## [493] "olivedrab"	"olivedrab1"	"olivedrab2"
## [496] "olivedrab3"	"olivedrab4"	"orange"
## [499] "orange1"	"orange2"	"orange3"
## [502] "orange4"	"orangered"	"orangered1"
## [505] "orangered2"	"orangered3"	"orangered4"
## [508] "orchid"	"orchid1"	"orchid2"
## [511] "orchid3"	"orchid4"	"palegoldenrod"
## [514] "palegreen"	"palegreen1"	"palegreen2"
## [517] "palegreen3"	"palegreen4"	"paleturquoise"
## [520] "paleturquoise1"	"paleturquoise2"	"paleturquoise3"
## [523] "paleturquoise4"	"palevioletred"	"palevioletred1"
## [526] "palevioletred2"	"palevioletred3"	"palevioletred4"

## [529] "papayawhip"	"peachpuff"	"peachpuff1"
## [532] "peachpuff2"	"peachpuff3"	"peachpuff4"
## [535] "peru"	"pink"	"pink1"
## [538] "pink2"	"pink3"	"pink4"
## [541] "plum"	"plum1"	"plum2"
## [544] "plum3"	"plum4"	"powderblue"
## [547] "purple"	"purple1"	"purple2"
## [550] "purple3"	"purple4"	"red"
## [553] "red1"	"red2"	"red3"
## [556] "red4"	"rosybrown"	"rosybrown1"
## [559] "rosybrown2"	"rosybrown3"	"rosybrown4"
## [562] "royalblue"	"royalblue1"	"royalblue2"
## [565] "royalblue3"	"royalblue4"	"saddlebrown"
## [568] "salmon"	"salmon1"	"salmon2"
## [571] "salmon3"	"salmon4"	"sandybrown"
## [574] "seagreen"	"seagreen1"	"seagreen2"
## [577] "seagreen3"	"seagreen4"	"seashell"
## [580] "seashell1"	"seashell2"	"seashell3"
## [583] "seashell4"	"sienna"	"sienna1"
## [586] "sienna2"	"sienna3"	"sienna4"
## [589] "skyblue"	"skyblue1"	"skyblue2"
## [592] "skyblue3"	"skyblue4"	"slateblue"
## [595] "slateblue1"	"slateblue2"	"slateblue3"
## [598] "slateblue4"	"slategray"	"slategray1"
## [601] "slategray2"	"slategray3"	"slategray4"
## [604] "slategrey"	"snow"	"snow1"
## [607] "snow2"	"snow3"	"snow4"
## [610] "springgreen"	"springgreen1"	"springgreen2"
## [613] "springgreen3"	"springgreen4"	"steelblue"
## [616] "steelblue1"	"steelblue2"	"steelblue3"
## [619] "steelblue4"	"tan"	"tan1"
## [622] "tan2"	"tan3"	"tan4"
## [625] "thistle"	"thistle1"	"thistle2"
## [628] "thistle3"	"thistle4"	"tomato"
## [631] "tomato1"	"tomato2"	"tomato3"
## [634] "tomato4"	"turquoise"	"turquoise1"
## [637] "turquoise2"	"turquoise3"	"turquoise4"
## [640] "violet"	"violetred"	"violetred1"
## [643] "violetred2"	"violetred3"	"violetred4"
## [646] "wheat"	"wheat1"	"wheat2"
## [649] "wheat3"	"wheat4"	"whitesmoke"
## [652] "yellow"	"yellow1"	"yellow2"
## [655] "yellow3"	"yellow4"	"yellowgreen"

More info about color scheme extractor code in R:

- Adobe Color (Adobe Kuler)
- Color Schema Designer
- Color Calculator
- Another products

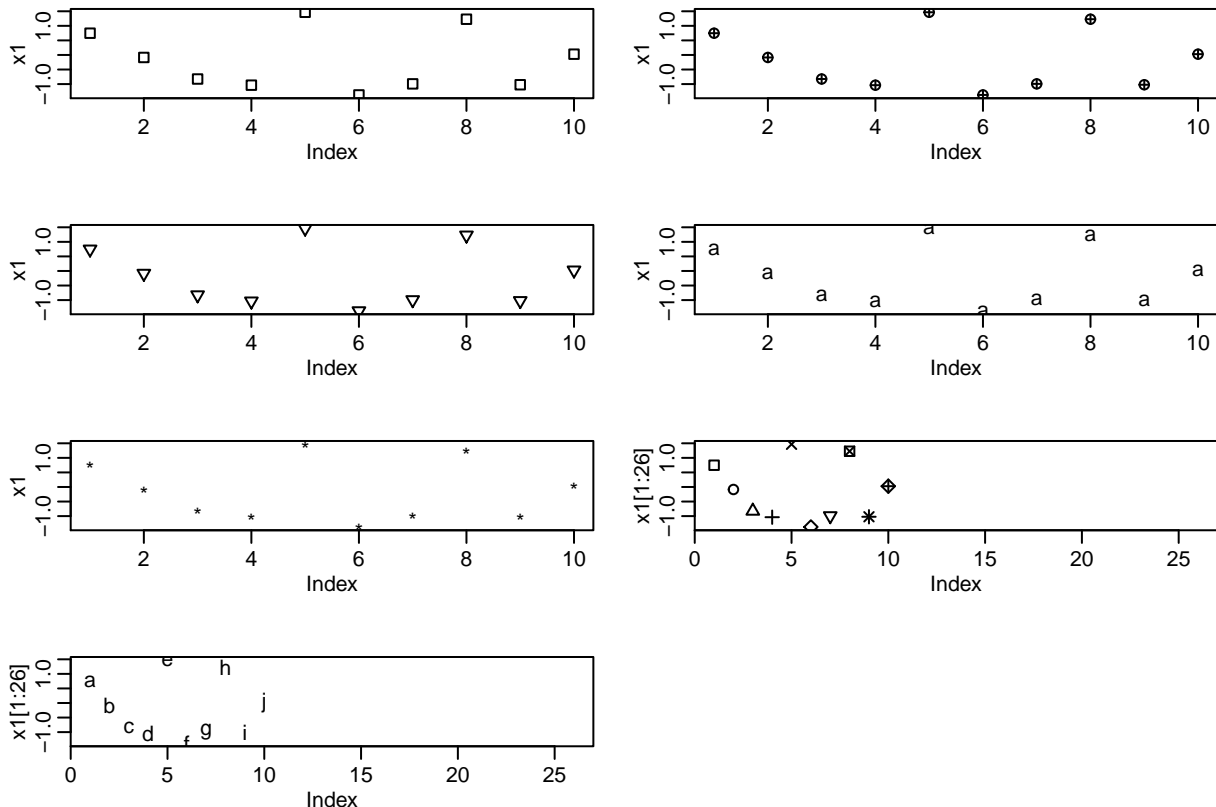
Points:

Using `pch=""`. It has values between 1 till 25 or a single character.

```

par(mfrow=c(4,2), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
x1<-rnorm(10)
plot(x1, type = "p", pch = 0)
plot(x1, type = "p", pch = 10)
plot(x1, type = "p", pch = 25)
plot(x1, type = "p", pch = "a")
plot(x1, type = "p", pch = "*")
plot(x1[1:26], type = "p", pch = 0:25)
plot(x1[1:26], type = "p", pch = letters)

```



Lines:

====Line types in R : lty====

`lty` for changing the lines type and `lwd` to change line width. The argument is a string ("blank", "solid", "dashed", "dotted", "dotdash", "longdash", or "twodash") or an integer (0=blank, 1=solid (default), 2=dashed, 3=dotted, 4=dotdash, 5=longdash, 6=twodash)

```

#Line types
#####
generateRLineTypes<-function(){
  oldPar<-par()
  par(font=2, mar=c(0,0,0,0))
  plot(1, pch="", ylim=c(0,6), xlim=c(0,0.7), axes=FALSE,xlab="", ylab="")
  for(i in 0:6) lines(c(0.3,0.7), c(i,i), lty=i, lwd=3)
  text(rep(0.1,6), 0:6, labels=c("0.'blank'", "1.'solid'", "2.'dashed'", "3.'dotted'",
    "4.'dotdash'", "5.'longdash'", "6.'twodash'"))
}

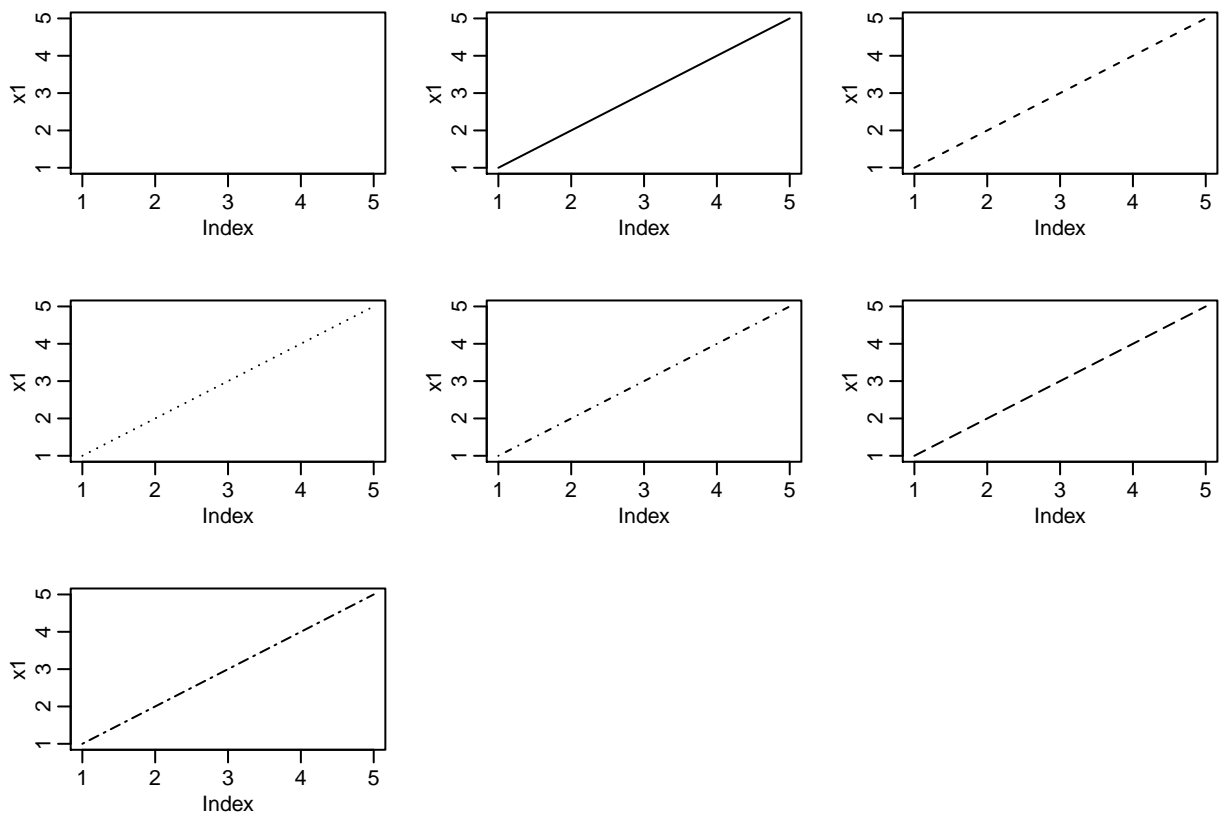
```

```
par(mar=oldPar$mar,font=oldPar$font )
}
generateRLineTypes()
```

6. 'twodash'	- - - - -
5. 'longdash'	- - - - - .
4. 'dotdash'	. - . - . - . - . - . - . - . -
3. 'dotted'
2. 'dashed'	- - - - -
1. 'solid'	_____
0. 'blank'	

Example:

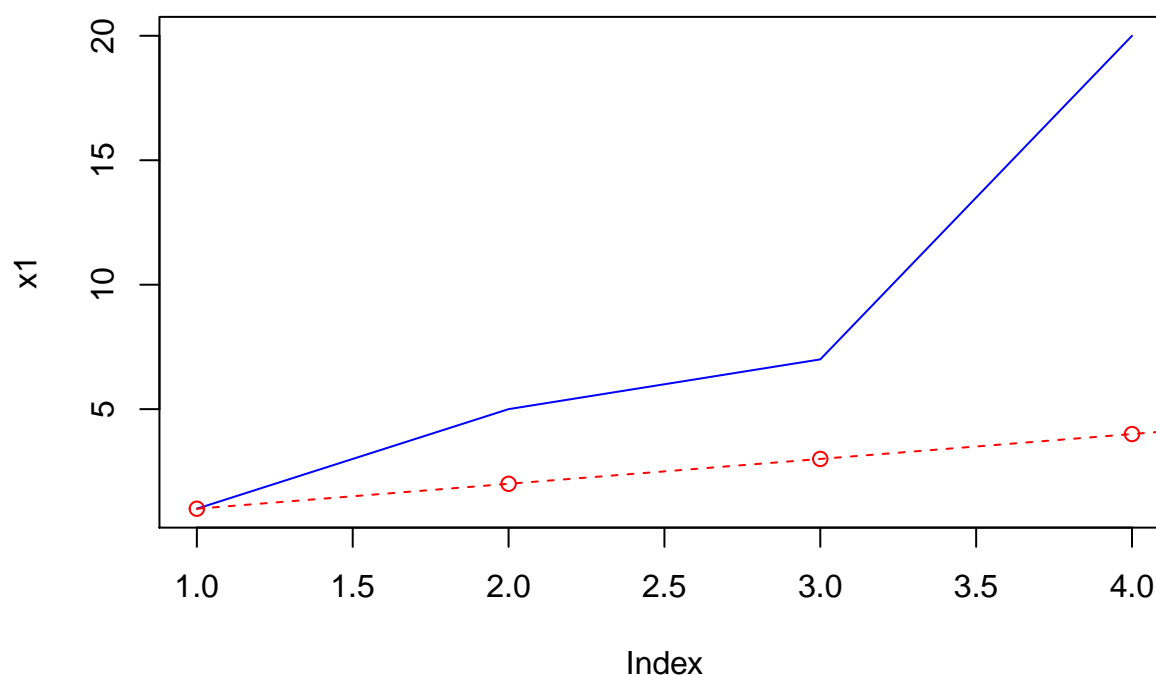
```
par(mfrow=c(3,3), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
x1<-1:5
plot(x1, type = "l", lty = "blank")
plot(x1, type = "l", lty = "solid")
plot(x1, type = "l", lty = "dashed")
plot(x1, type = "l", lty = "dotted")
plot(x1, type = "l", lty = "dotdash")
plot(x1, type = "l", lty = "longdash")
plot(x1, type = "l", lty = "twodash")
```



====An additional `lines()` on a graph=====

Example:

```
x1<-c(1,5,7,20)
x2<-1:10
plot(x1, type = "l", lty = "solid",col="blue")
lines(x2, type = "o", lty = "dashed", col = "red")
```

====Straight `abline()`====

`abline()` can be used to add vertical, horizontal or regression lines.

Formula:

`abline(a=NULL, b=NULL, h=NULL, v=NULL, ...)`

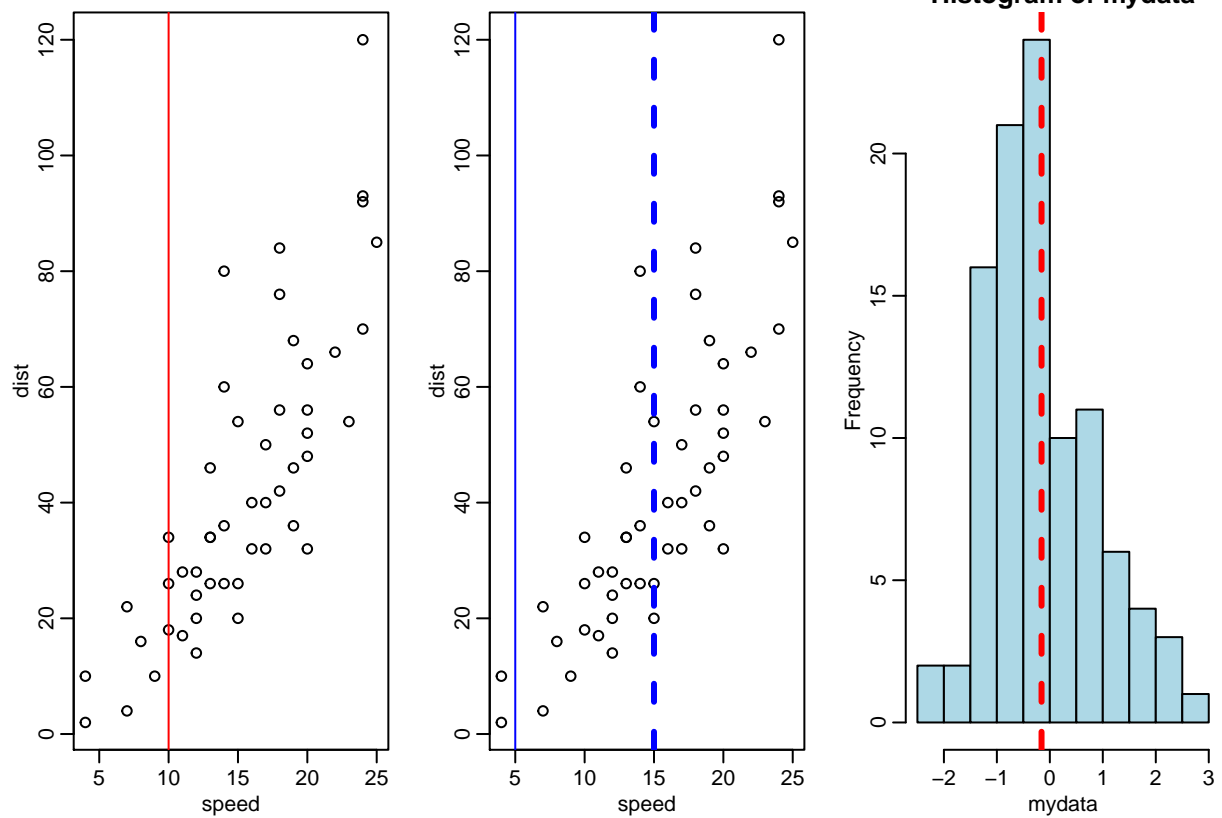
Vertical line:

```
par(mfrow=c(1,3), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics

#Example-1: Add one line
plot(cars)
abline(v=10, col="red")

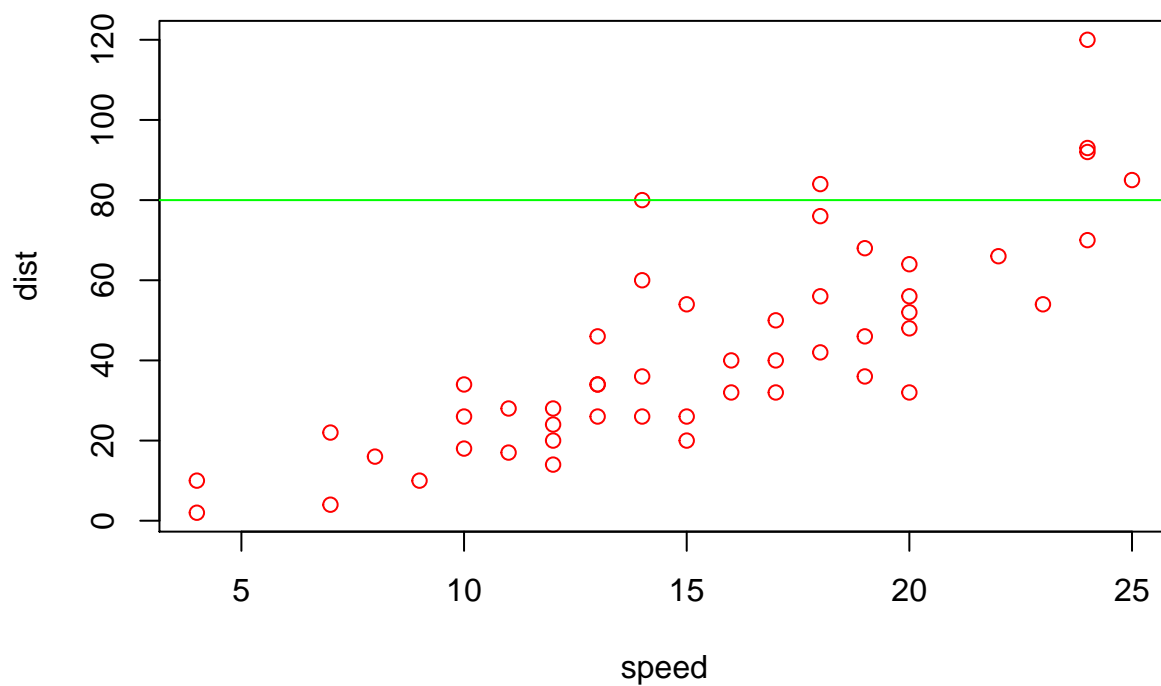
#Example-2: add 2 lines
# change line colors, sizes and types
plot(cars)
abline(v=c(5,15), col=c("blue", "blue"), lty=c(1,2), lwd=c(1, 3))

#Example-3:
set.seed(1234); mydata<-rnorm(100)
hist(mydata, col="lightblue")
abline(v = mean(mydata), col="red", lwd=3, lty=2)
```



Horizontal line:

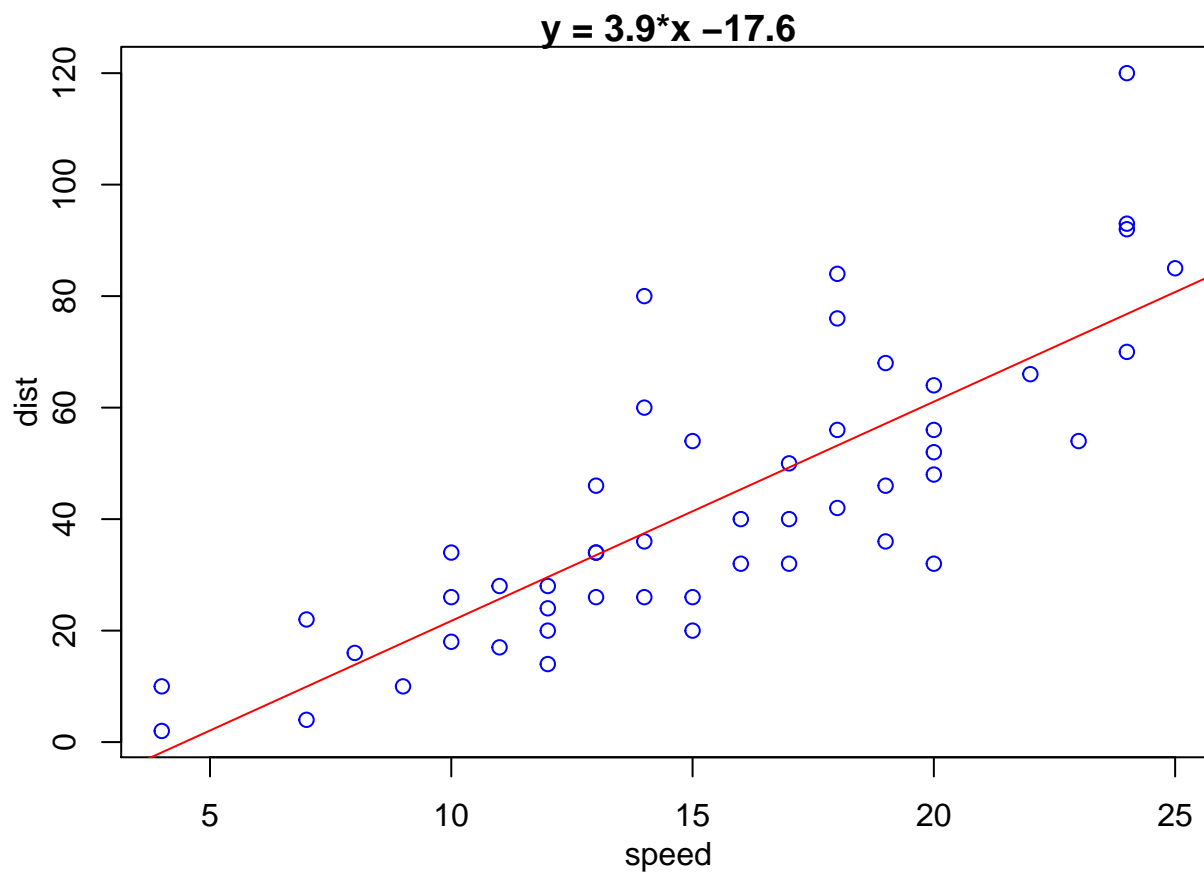
```
plot(cars,col="red")
abline(h=80, col="green")
```



Regression line:

lm() function to fit linear model.

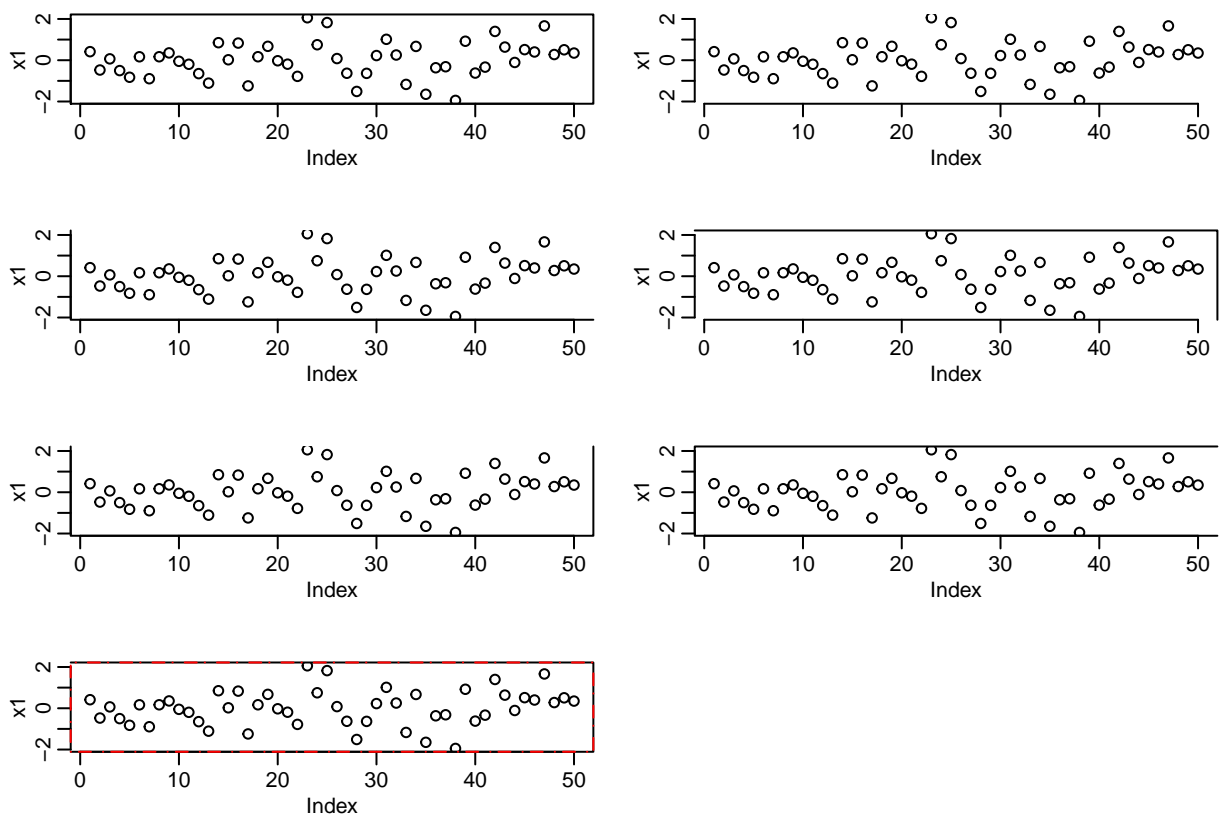
```
par(mgp=c(2,1,0), mar=c(3,3,1,1))
# Fit regression line
require(stats)
reg<-lm(dist ~ speed, data = cars)
coeff=coefficients(reg)
# equation of the line :
eq = paste0("y = ", round(coeff[2],1), "*x ", round(coeff[1],1))
# plot
plot(cars, main=eq,col="blue")
abline(reg, col="red")
```



Boxes:

bty specifies the box type.

```
par(mfrow=c(4,2), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0)) # set up the graphics
x1<-rnorm(50)
plot(x1, bty = "o") # the default
plot(x1, bty = "n") # no box
plot(x1, bty = "l")
plot(x1, bty = "7")
plot(x1, bty = "u")
plot(x1, bty = "c")
plot(x1, bty = "]")
box(lty = 'l375', col = 'red')
```



Grid:

`grid()` adds a grid to the current graph.

Example:

```
x1<-rnorm(100)
plot(x1, col="#FF8100")
grid()
```

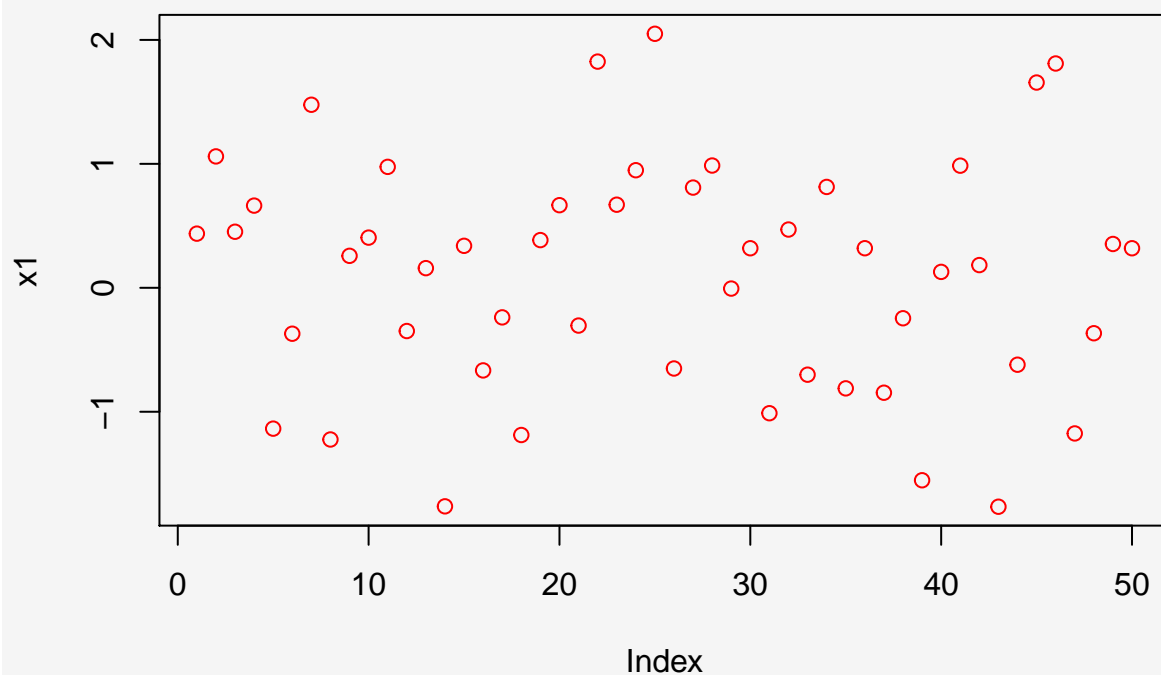


Background:

Change the background color with `par(bg=)`.

Example:

```
x1<-rnorm(50)
par(bg="whitesmoke")
plot(x1, bty="o",col="red") # the default
```

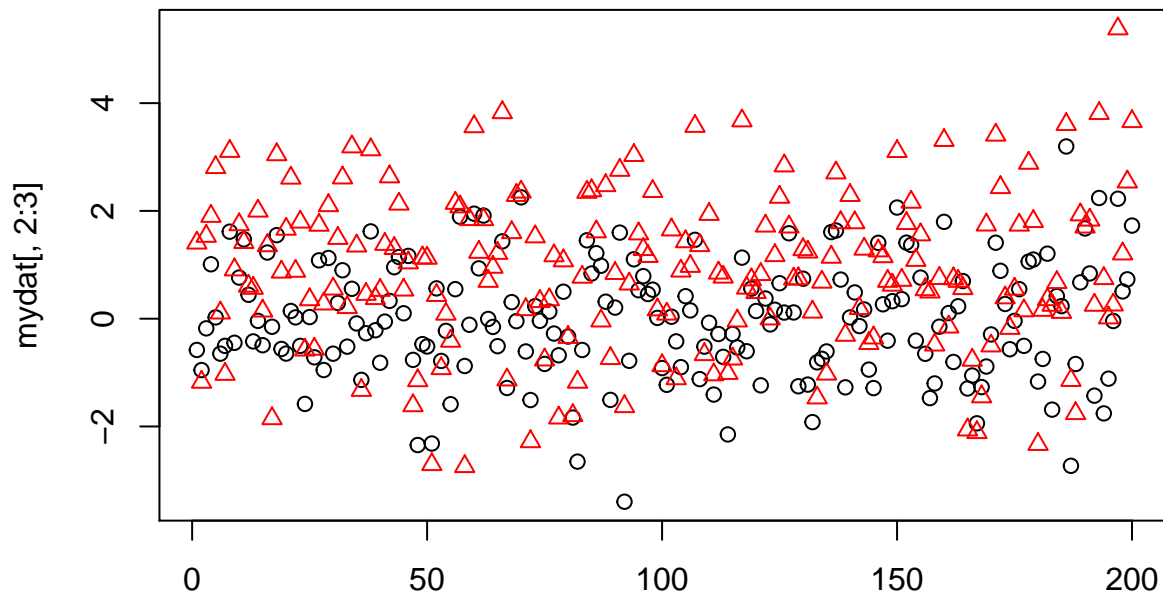


Overlaying plots:

`matplot()`

Example:

```
N <- 200
x1 <- rnorm(N)
x2 <- rnorm(N) + x1 + 1
y <- 1 + x1 + x2 + rnorm(N)
mydat <- data.frame(y,x1,x2)
matplot(mydat[,2:3], pch = 1:2)
```

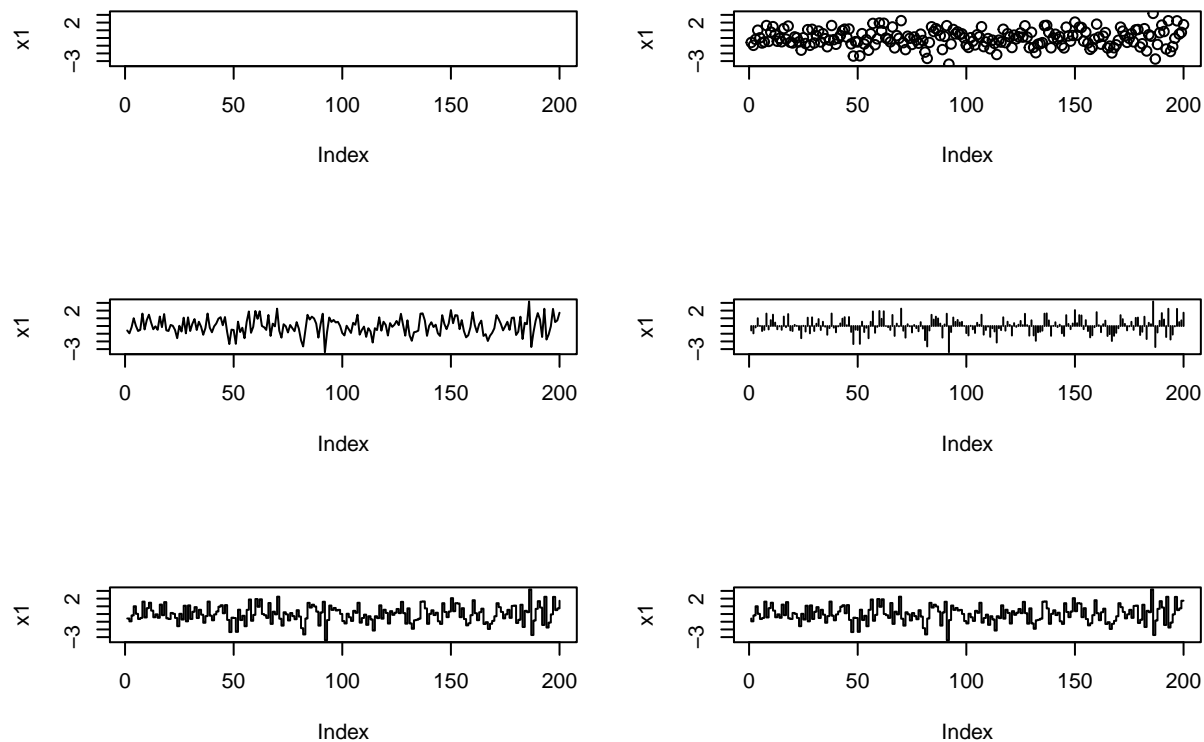


Multiple plots:

- `par()` for display multiple figures
- `mfrow = c(3,2)` : 3 row x 2 column
- `mfcol = c(3,2)` : same with `mfrow` but another positions

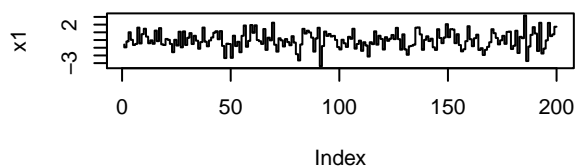
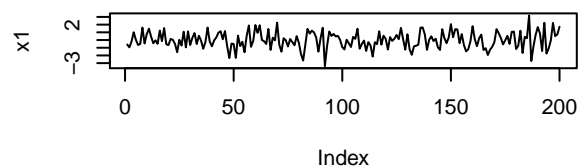
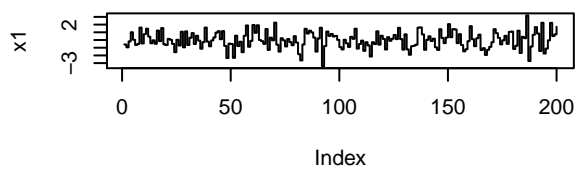
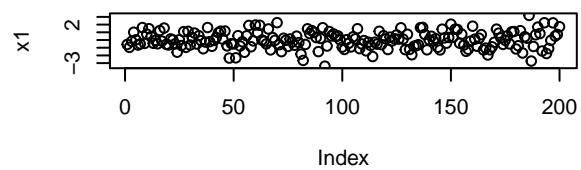
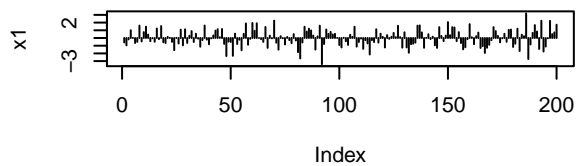
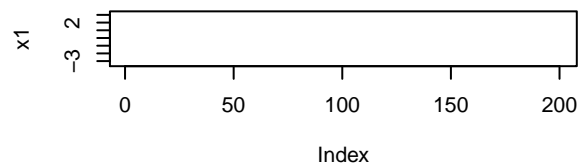
Example-1:

```
par(mfrow = c(3,2))
plot(x1, type = "n")
plot(x1, type = "p")
plot(x1, type = "l")
plot(x1, type = "h")
plot(x1, type = "s")
plot(x1, type = "S")
```

Example-2:

```
par(mfcol = c(3,2))
plot(x1, type = "n")
plot(x1, type = "p")
plot(x1, type = "l")
plot(x1, type = "h")
plot(x1, type = "s")
plot(x1, type = "S")
```

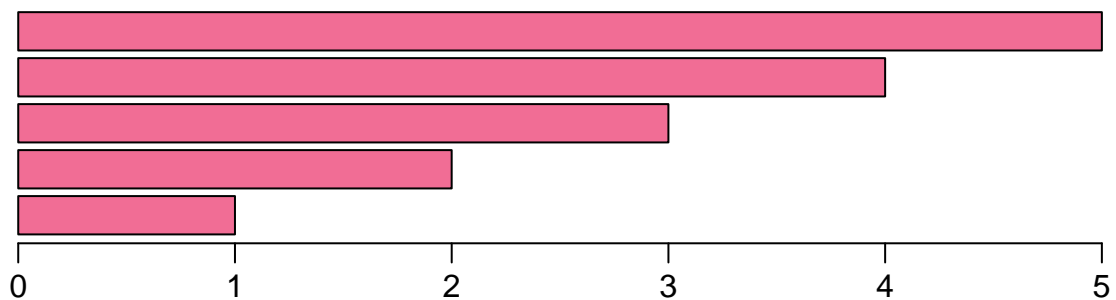
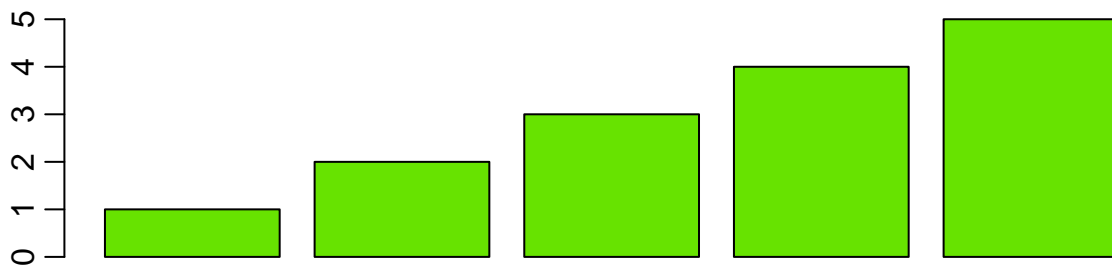


Chapter 4

Default Bar plots in R

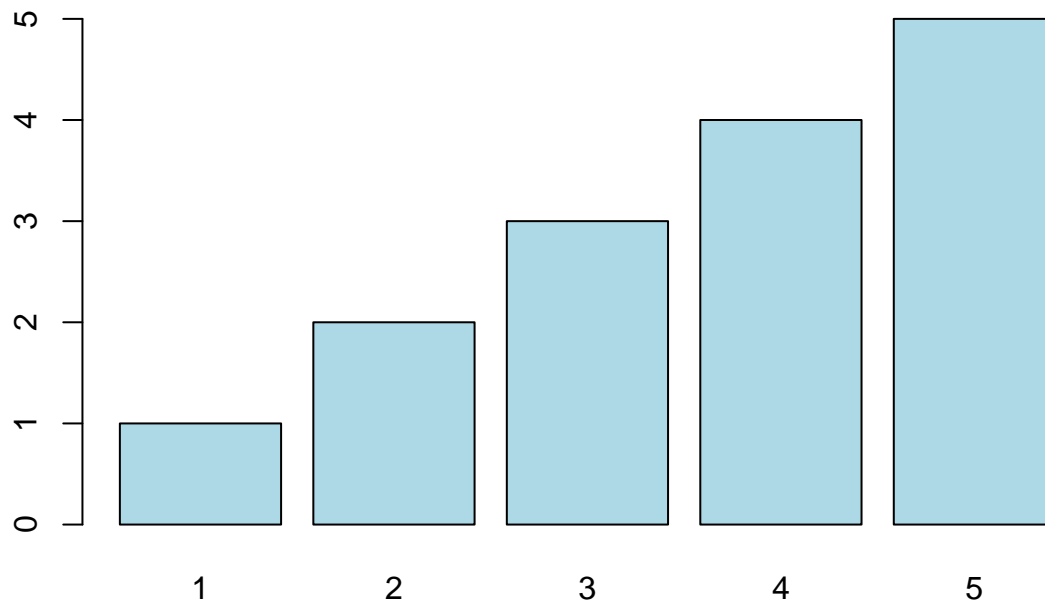
Basic bar plots:

```
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0))
x<-c(1:5)
barplot(x,col="#67E300")
barplot(x,hORIZ=TRUE,col="#F16D95")
```



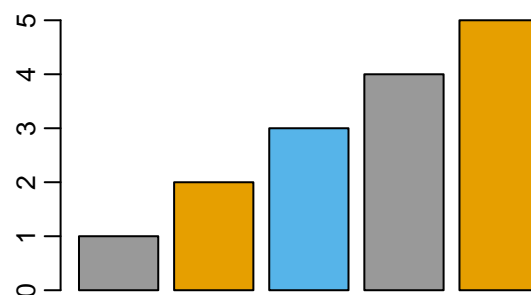
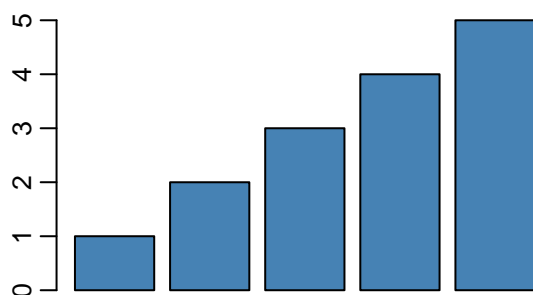
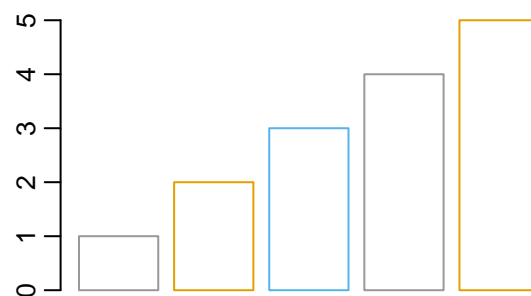
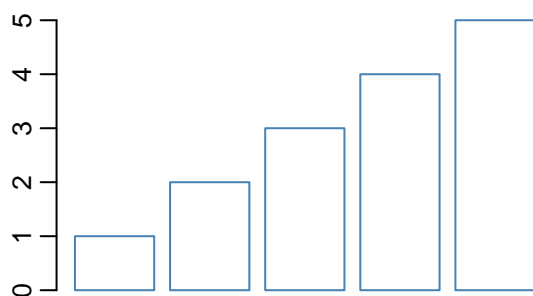
Change group names:

```
barplot(x, names.arg = c("1", "2", "3","4","5"), col="lightblue")
```



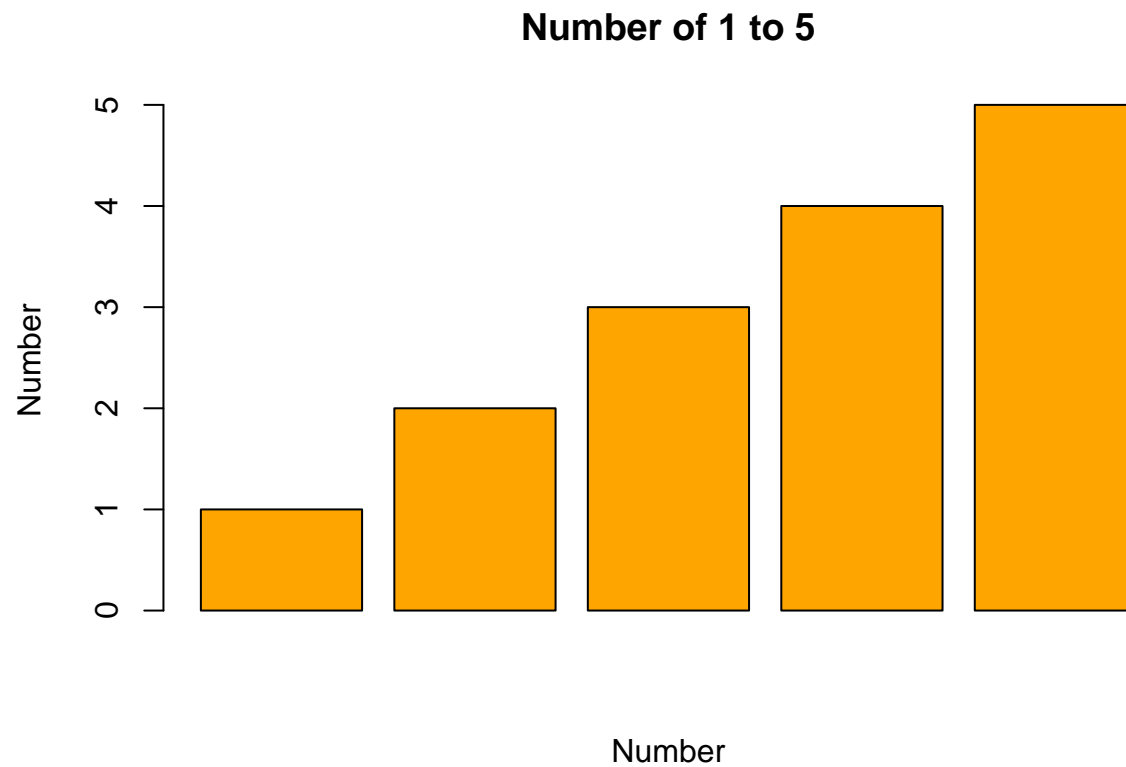
Change color:

```
par(mfrow=c(2,2), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0))
x<-c(1:5)
# Change border and fill color using one single color
barplot(x, col = "white", border = "steelblue")
# Change the color of border.
# Use different colors for each group
barplot(x, col = "white",
        border = c("#999999", "#E69F00", "#56B4E9"))
# Change fill color : single color
barplot(x, col = "steelblue")
# Change fill color: multiple colors
barplot(x, col = c("#999999", "#E69F00", "#56B4E9"))
```



Change main title and axis labels:

```
# Change axis titles
# Change color (col = "orange") and remove frame
x<-c(1:5)
barplot(x, main = "Number of 1 to 5",
        xlab = "Number", ylab = "Number", col="orange")
```



Stacked bar plots:

Example-1:

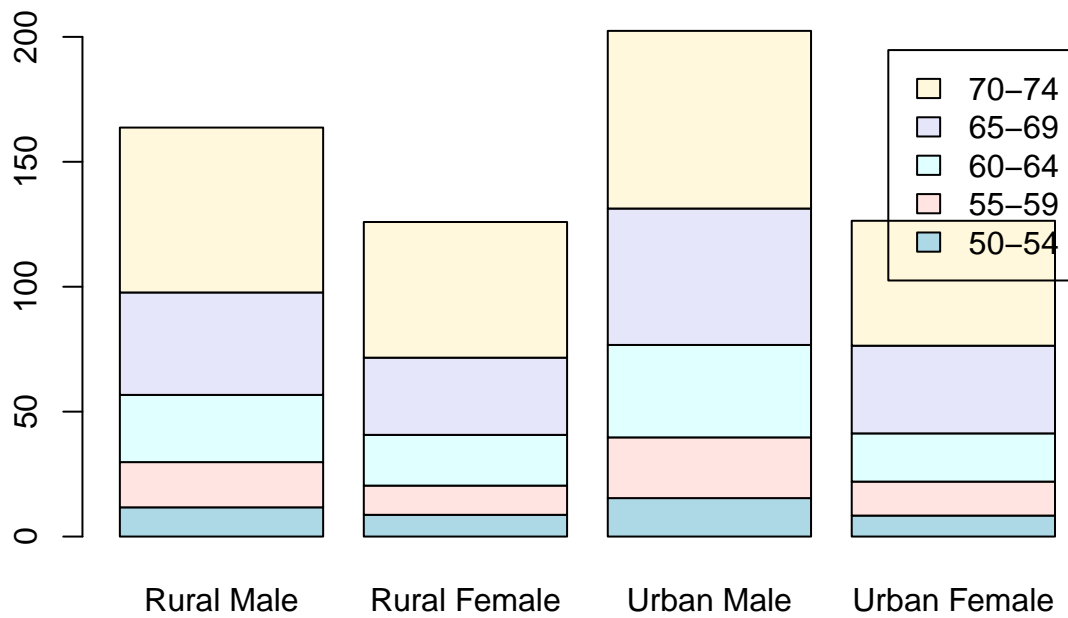
```
barplot(VADeaths,  
        col = c("lightblue", "mistyrose", "lightcyan",  
                "lavender", "cornsilk"),  
        legend = rownames(VADeaths))
```



Stacked bar plots:

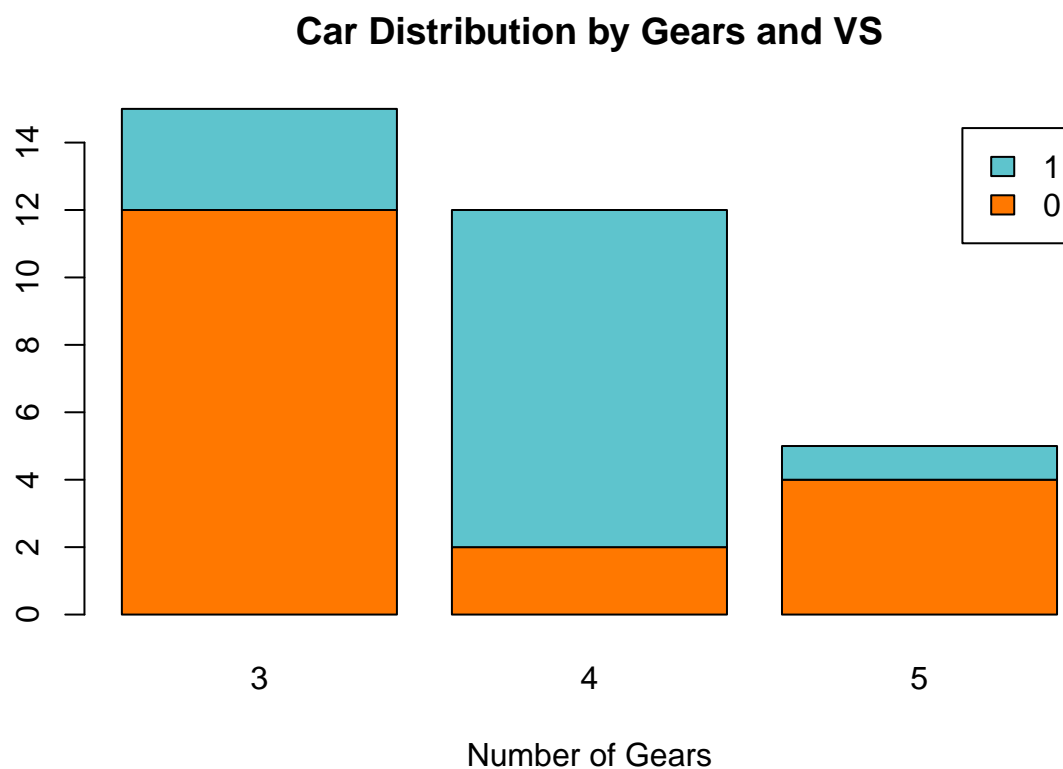
Example-1:

```
barplot(VADeaths,
  col = c("lightblue", "mistyrose", "lightcyan",
    "lavender", "cornsilk"),
  legend = rownames(VADeaths))
```



Example-2:

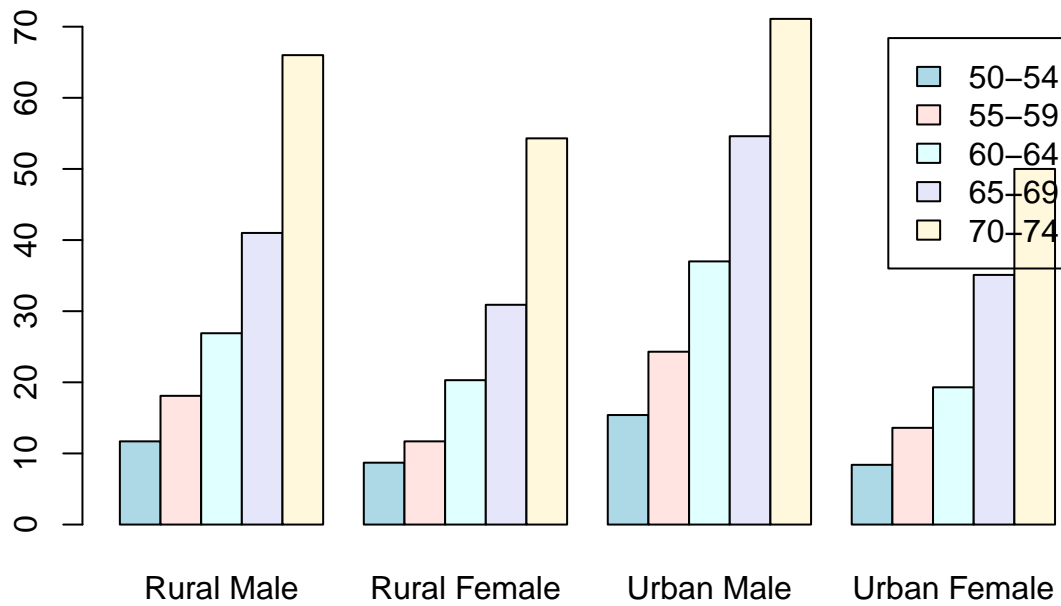
```
# Stacked Bar Plot with Colors and Legend
counts <- table(mtcars$vs, mtcars$gear)
barplot(counts, main="Car Distribution by Gears and VS",
        xlab="Number of Gears", col=c("#FF7800", "#5EC4CD"),
        legend = rownames(counts))
```

Grouped bar plots:

Example-1:

```
barplot(VADeaths,  
        col = c("lightblue", "mistyrose", "lightcyan",  
                "lavender", "cornsilk"),  
        legend = rownames(VADeaths), beside = TRUE)
```



Example-2:

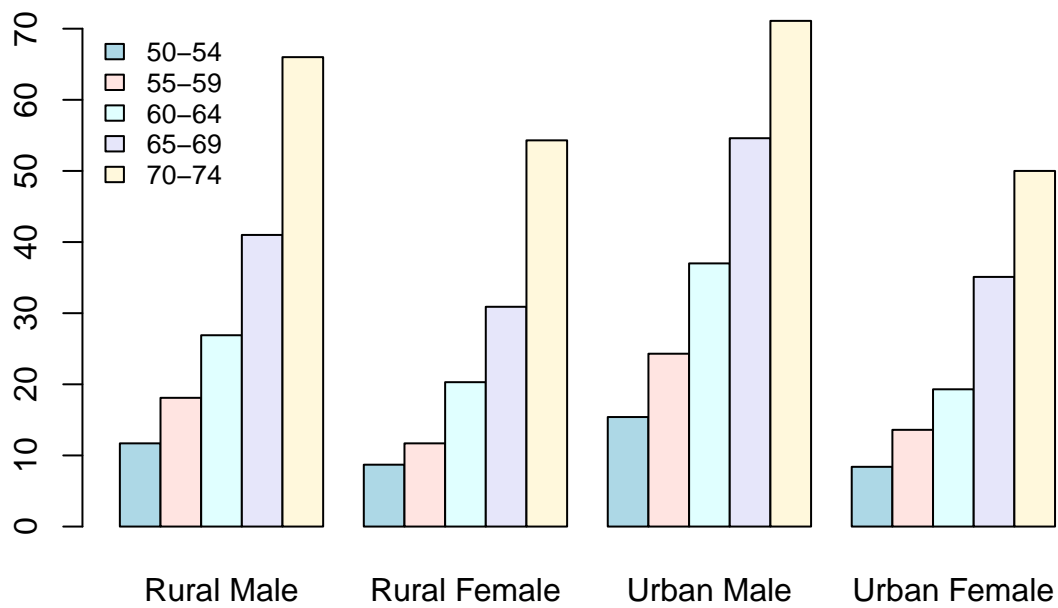
- `box.lty = 0`: Remove the box around the legend

- `cex = 0.8`: legend text size

```
# Define a set of colors
my_colors <- c("lightblue", "mistyrose", "lightcyan",
               "lavender", "cornsilk")

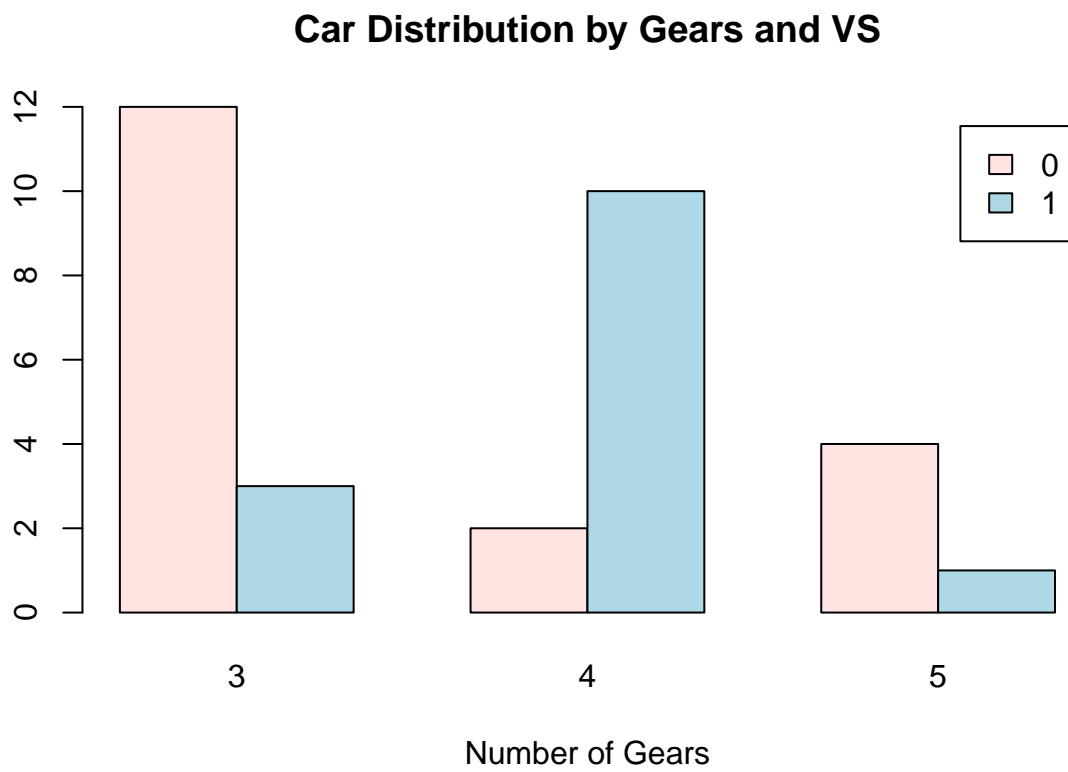
# Bar plot
barplot(VADeaths, col = my_colors, beside = TRUE)

# Add legend
legend("topleft", legend = rownames(VADeaths),
       fill = my_colors, box.lty = 0, cex = 0.8)
```



Example-3:

```
# Grouped Bar Plot
counts <- table(mtcars$vs, mtcars$gear)
barplot(counts, main="Car Distribution by Gears and VS",
        xlab="Number of Gears", col=c("mistyrose","lightblue"),
        legend = rownames(counts), beside=TRUE)
```



Note: If you need more formatting, see the **3-Standard R graphs formatting**.

Chapter 5

Default Histogram and Density Plots in R

A Histogram is NOT a Bar Chart:

Preference: <https://www.edrawsoft.com/histogram-vs-bar-chart.php>

- Histograms VS. Bar Charts
- A Histogram is NOT a Bar Chart

Simple Histograms:

- x: a numeric vector
- breaks: breakpoints between histogram cells.

```
# Simple Histogram
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0))
hist(mtcars$mpg,col="mistyrose")
hist(mtcars$mpg, breaks =30,col="pink")
```

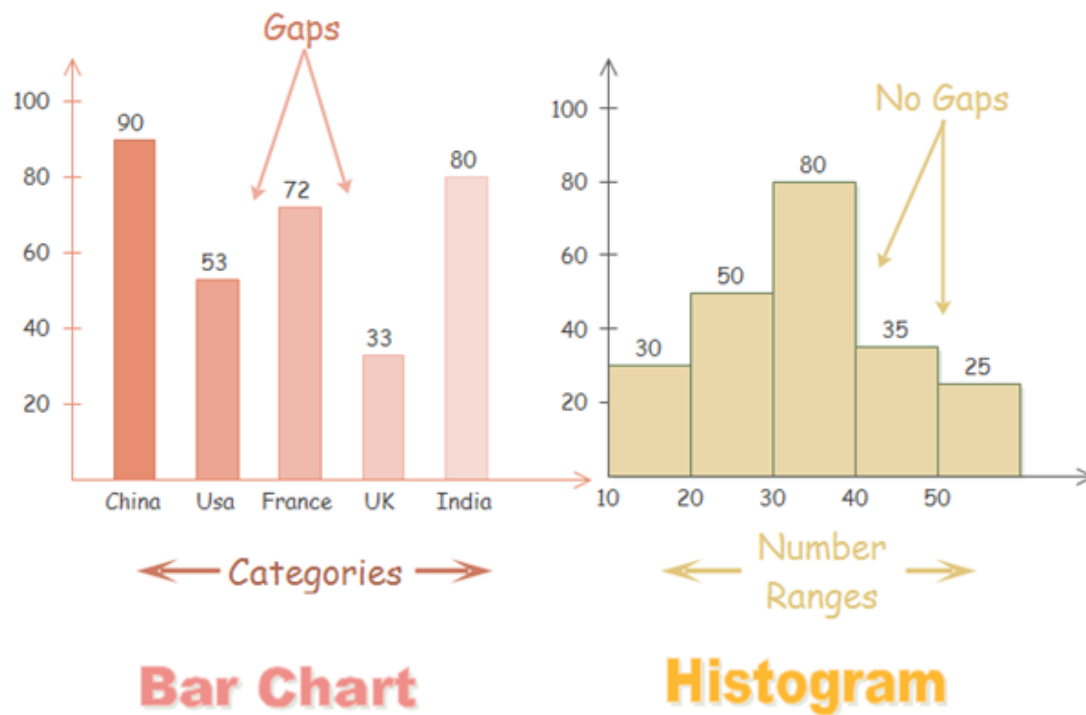
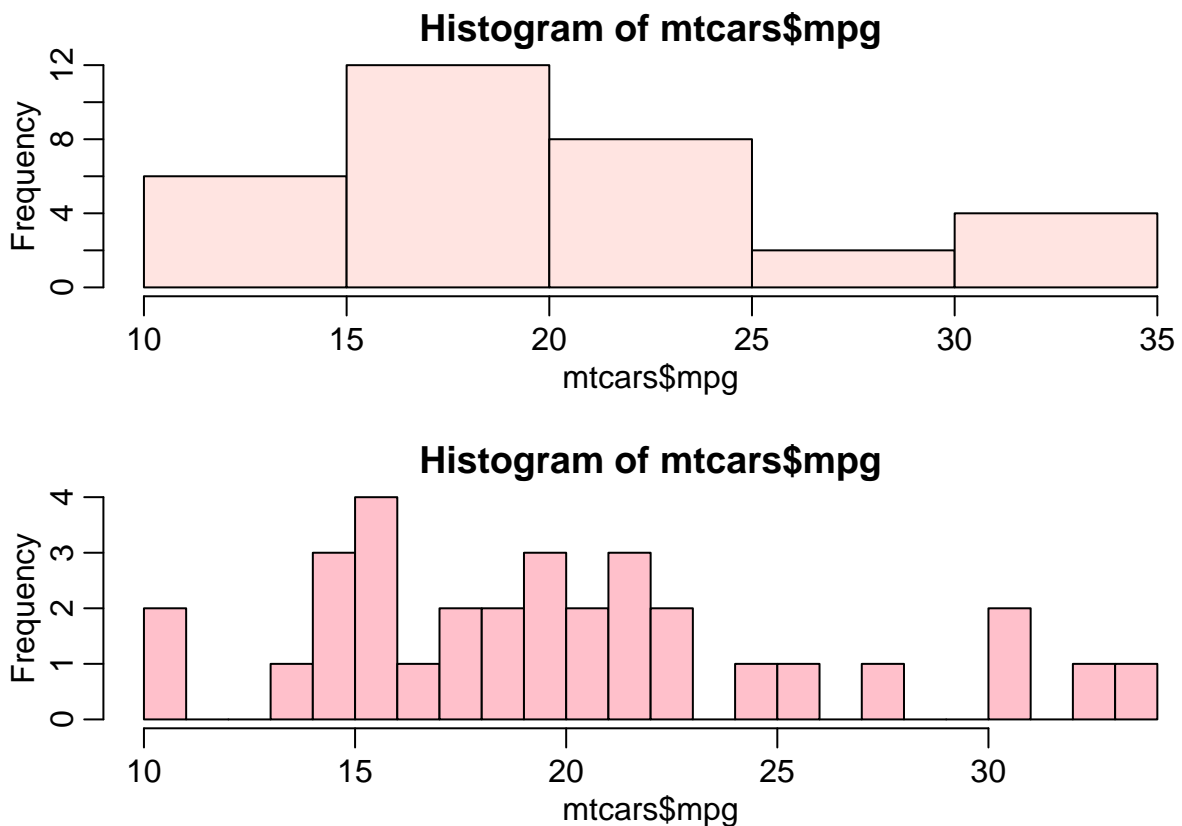
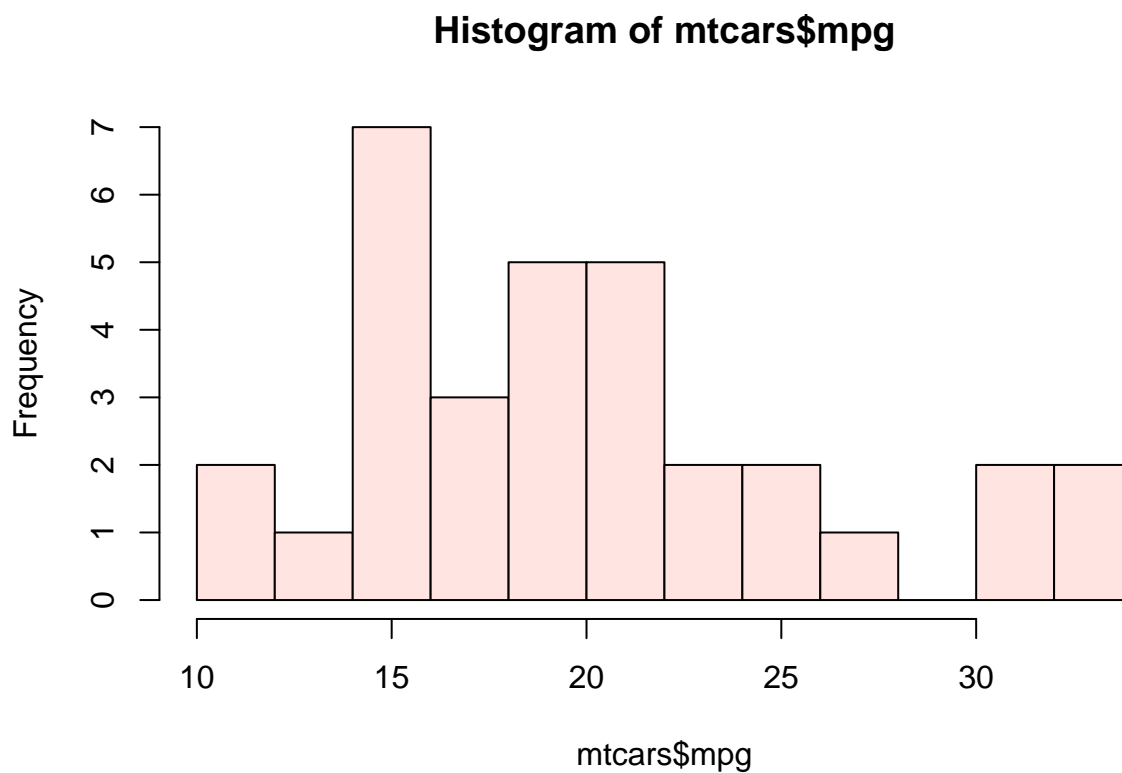


Figure 5.1:



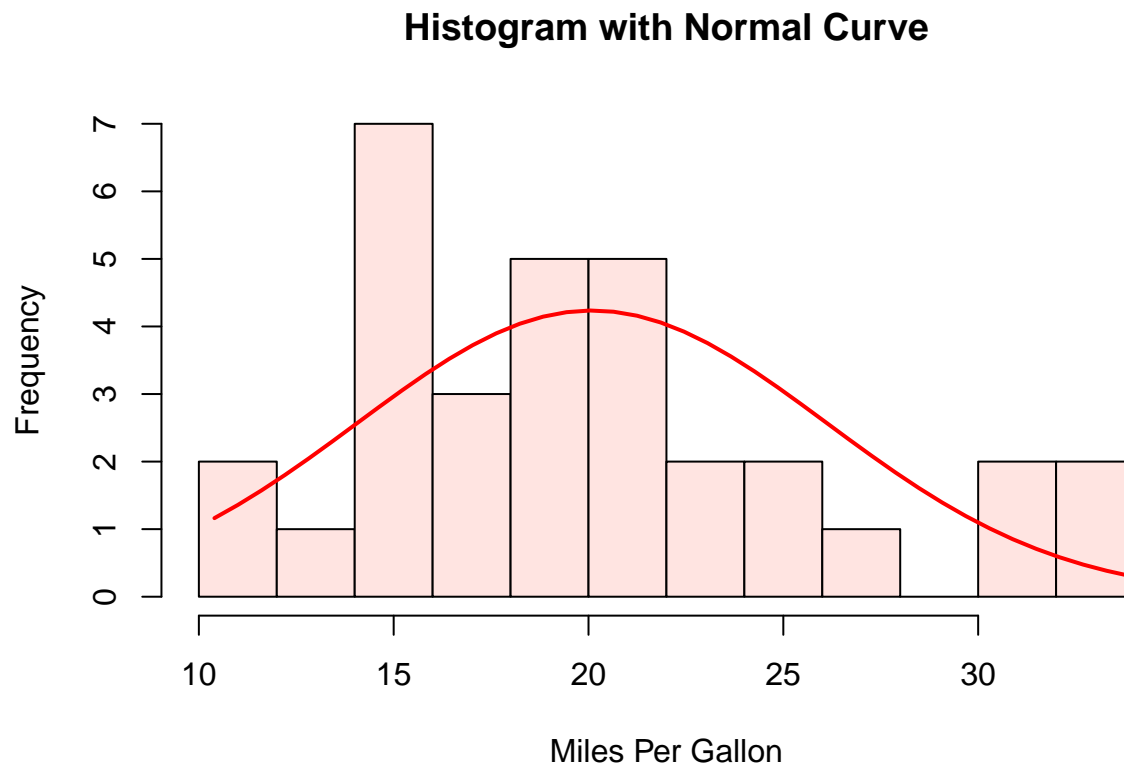
Colored Histogram:

```
hist(mtcars$mpg, breaks=12, col="mistyrose")
```



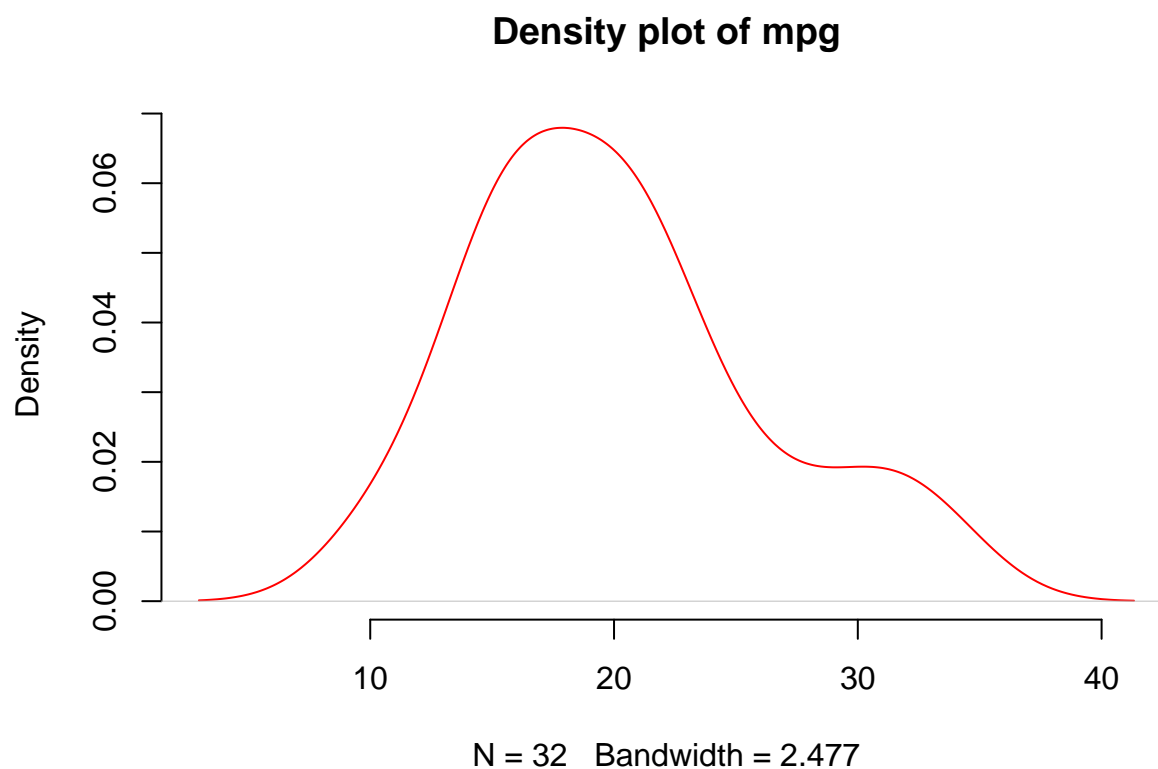
Add a Normal Curve:

```
x <- mtcars$mpg
h<-hist(x, breaks=10, col="mistyrose", xlab="Miles Per Gallon",
      main="Histogram with Normal Curve")
xfit<-seq(min(x),max(x),length=40)
yfit<-dnorm(xfit,mean=mean(x),sd=sd(x))
yfit <- yfit*diff(h$mids[1:2])*length(x)
lines(xfit, yfit, col="red", lwd=2)
```



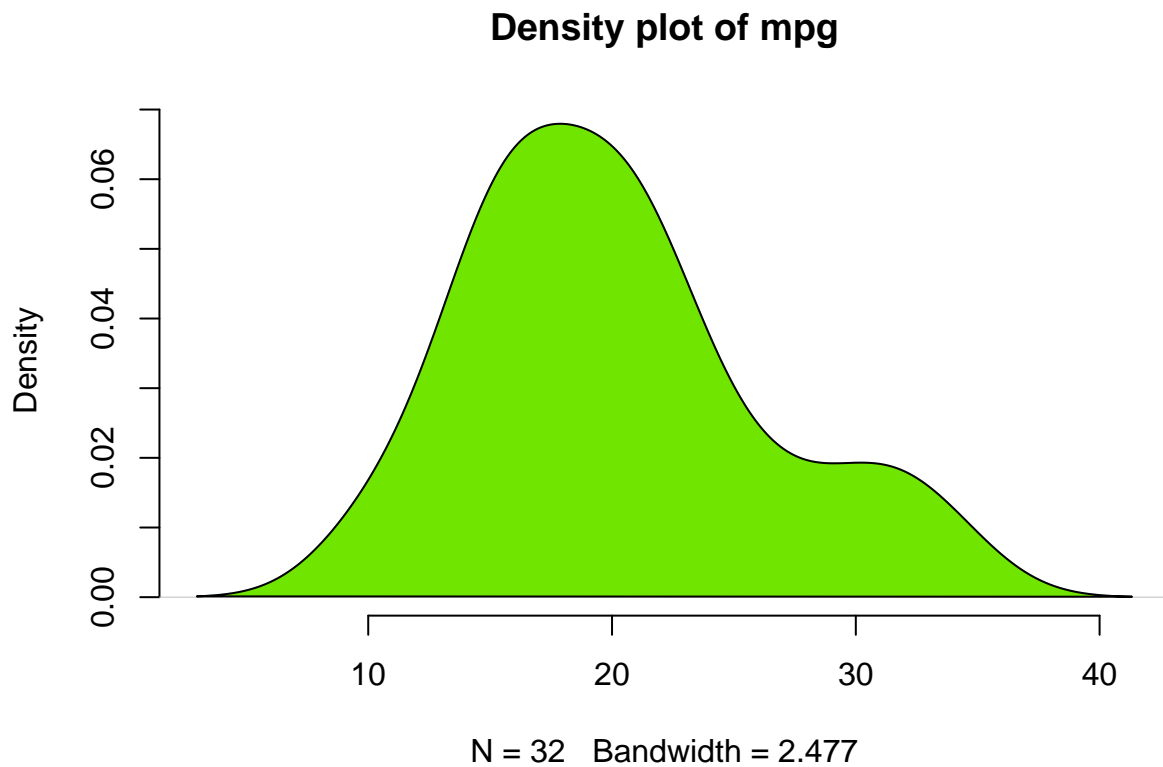
Density plots: `density()`:

```
# The density data
dens <- density(mtcars$mpg)
# plot density
plot(dens, frame = FALSE, col = "red",
     main = "Density plot of mpg")
```

Density plot using `polygon()`:

```
plot(dens, frame = FALSE, col = "lightblue",  
     main = "Density plot of mpg")  
polygon(dens, col = "#70E500")
```



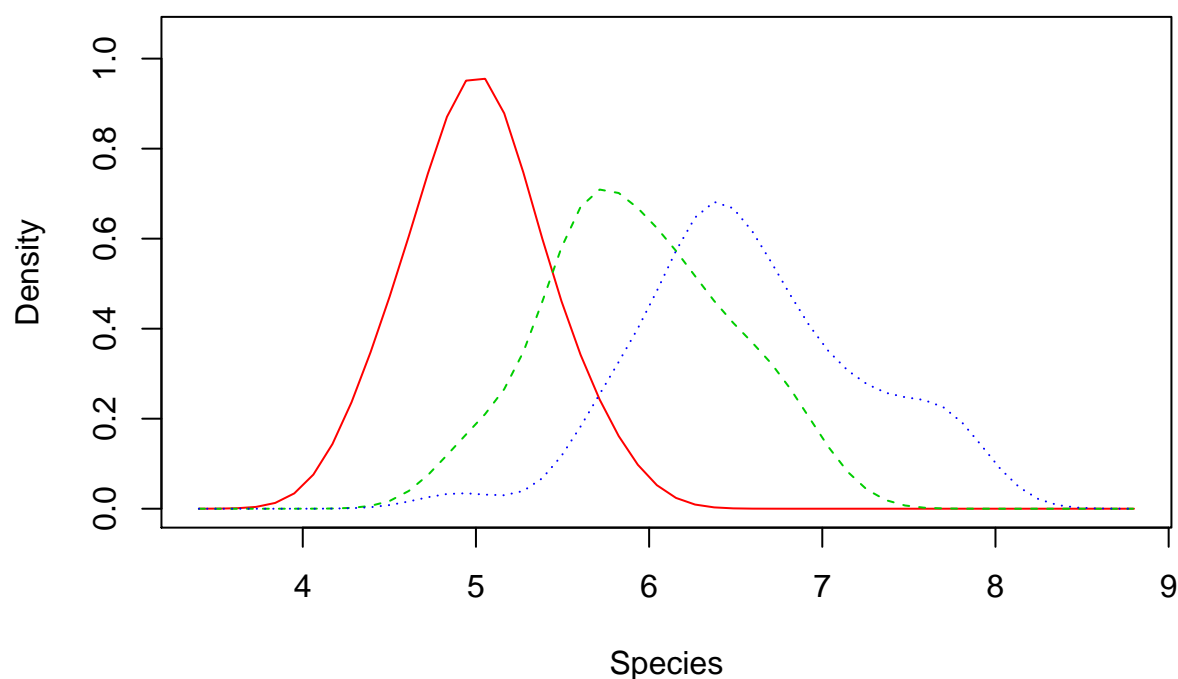
Additional->Comparing density Plots:

Example-1:

```
#install.packages("sm")  
library(sm)
```

```
## Package 'sm', version 2.2-5.6: type help(sm) for summary information  
sm.density.compare(iris$Sepal.Length, iris$Species, xlab="Species")  
title(main="Distributions of Species")
```

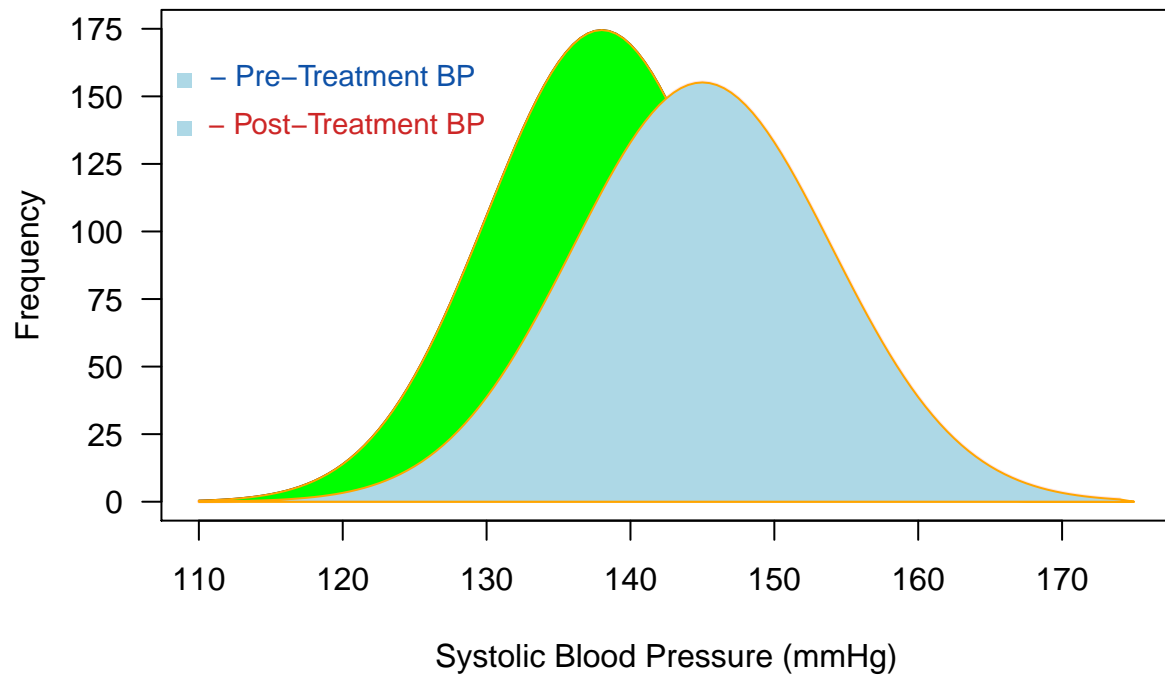
Distributions of Species



Example-2:

```
x <- seq(from = 110, to = 174, by = 0.5)
y1 <- dnorm(x, mean = 145, sd = 9)
y2 <- dnorm(x, mean = 138, sd = 8)
plot(x, y1, type="l", lwd=2, col="mistyrose",
     main="Systolic Blood Pressure Before and After Treatment",
     xlab = "Systolic Blood Pressure (mmHg)",
     ylab = "Frequency", yaxt="n",
     xlim = c(110, 175), ylim = c(0, 0.05))
lines(x, y2)
polygon(c(110,x,175),c(0,y2,0), col="green",
       border = "orange")
polygon(c(117,x,175),c(0,y1,0), col="lightblue",
       border = "orange")
ylab=c(seq(from=0, to=175, by=25))
y=c(seq(from=0, to=0.05, length.out = 8))
axis(2,at=y,labels=ylab, las=1)
text(x = 120, y = 0.045, "- Pre-Treatment BP", col = "#0E51A7", cex = 0.9)
text(x = 120, y = 0.04, " - Post-Treatment BP", col = "firebrick3", cex = 0.9)
points(109, 0.0445, pch = 15, col = "lightblue")
points(109, 0.0395, pch = 15, col = "lightblue")
```

Systolic Blood Pressure Before and After Treatment



Chapter 6

Default Line Plots in R

Note that the function `lines()` need a `plot()`.

Basic line plots:

Example-1:

```
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0))
x <- 1:10
y1 <- x*x
y2 <- 3*y1
# Basic stair steps plot
plot(x, y1, type = "S", plot="pink")
```

```
## Warning in plot.window(...): "plot" is not a graphical parameter
```

```
## Warning in plot.xy(xy, type, ...): "plot" is not a graphical parameter
```

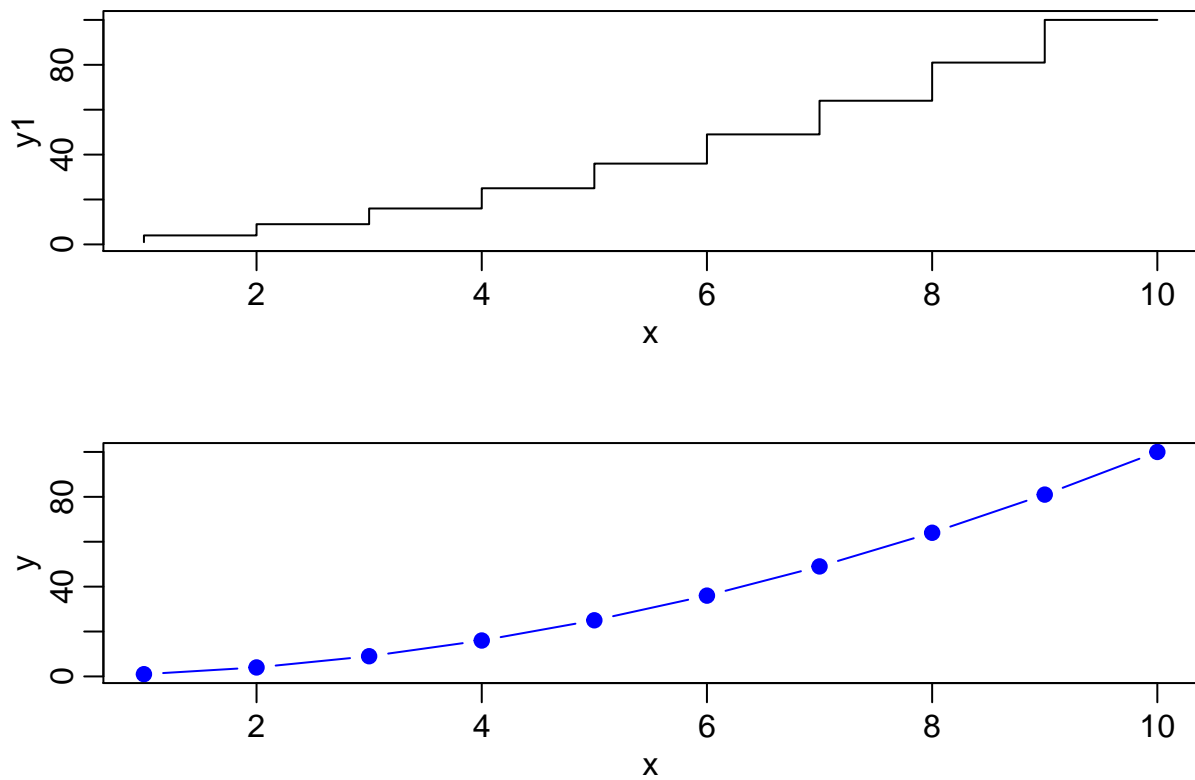
```
## Warning in axis(side = side, at = at, labels = labels, ...): "plot" is not
## a graphical parameter
```

```
## Warning in axis(side = side, at = at, labels = labels, ...): "plot" is not
## a graphical parameter
```

```
## Warning in box(...): "plot" is not a graphical parameter
```

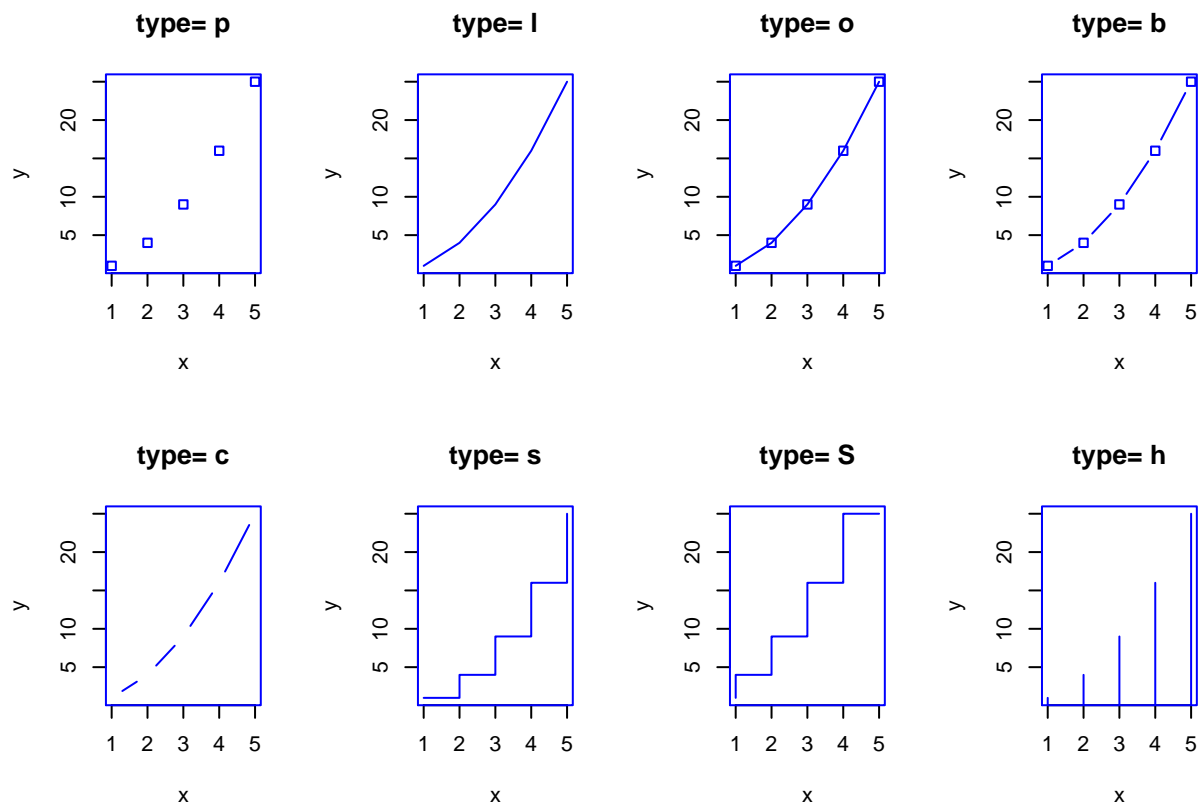
```
## Warning in title(...): "plot" is not a graphical parameter
```

```
# Points and line
plot(x, y1, type = "b", pch = 19,
     col = "blue", xlab = "x", ylab = "y")
```



Example-2: As functions for different types

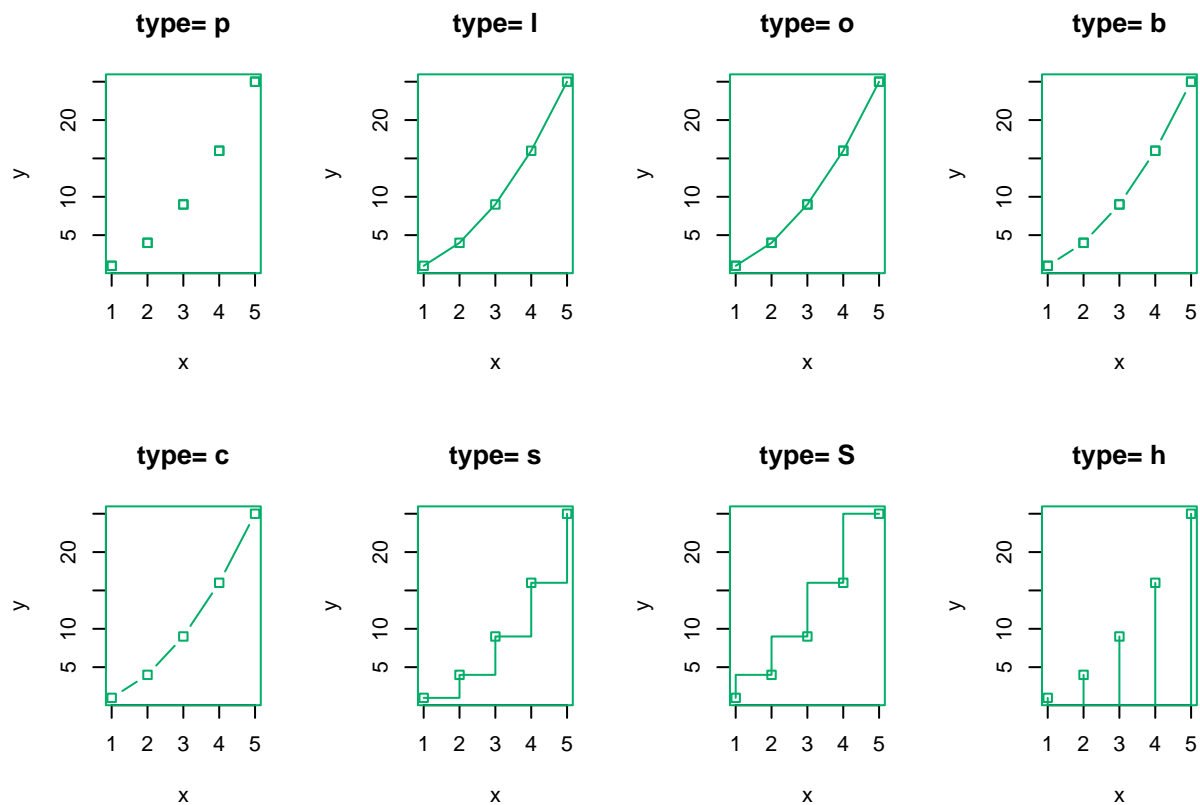
```
x <- c(1:5); y <- x*x # create example data
par(pch=22, col="blue") # plotting symbol and color
par(mfrow=c(2,4)) # all plots on one page
opts = c("p", "l", "o", "b", "c", "s", "S", "h")
for(i in 1:length(opts)){
  heading = paste("type=",opts[i])
  plot(x, y, type="n", main=heading)
  lines(x, y, type=opts[i])
}
```



Example-3: Plot the points type=""

```
x <- c(1:5); y <- x*x # create example data
par(pch=22, col="#00AE68") # plotting symbol and color
par(mfrow=c(2,4)) # all plots on one page
opts = c("p","l","o","b","c","s","S","h")

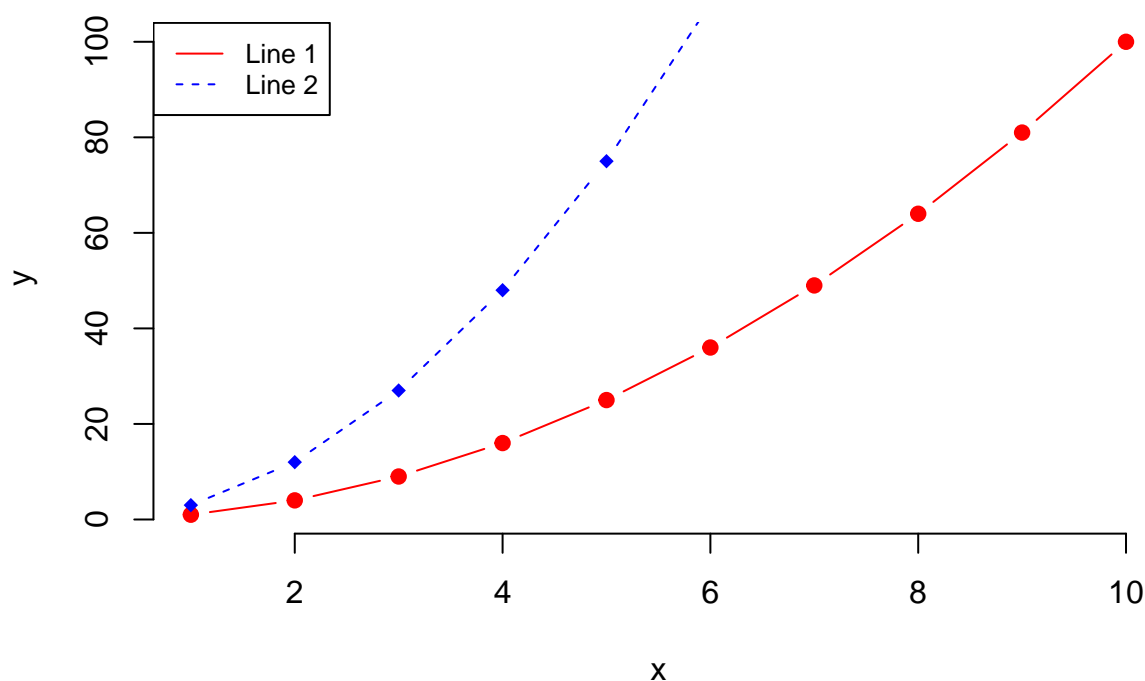
for(i in 1:length(opts)){
  heading = paste("type=",opts[i])
  plot(x, y, main=heading)
  lines(x, y, type=opts[i]) }
```



Multiple lines plots:

Example-1:

```
# First line
x <- 1:10
y1 <- (x*x)
plot(x, y1, type = "b", frame = FALSE, pch = 19,
     col = "red", xlab = "x", ylab = "y")
# A second line
lines(x, y2, pch = 18, col = "blue", type = "b", lty = 2)
# Add a legend to the plot
legend("topleft", legend=c("Line 1", "Line 2"),
     col=c("red", "blue"), lty = 1:2, cex=0.8)
```

Example-2: As functions for different types

```
# Create Line Chart
# convert factor to numeric for convenience
Orange$Tree <- as.numeric(Orange$Tree)
ntrees <- max(Orange$Tree)

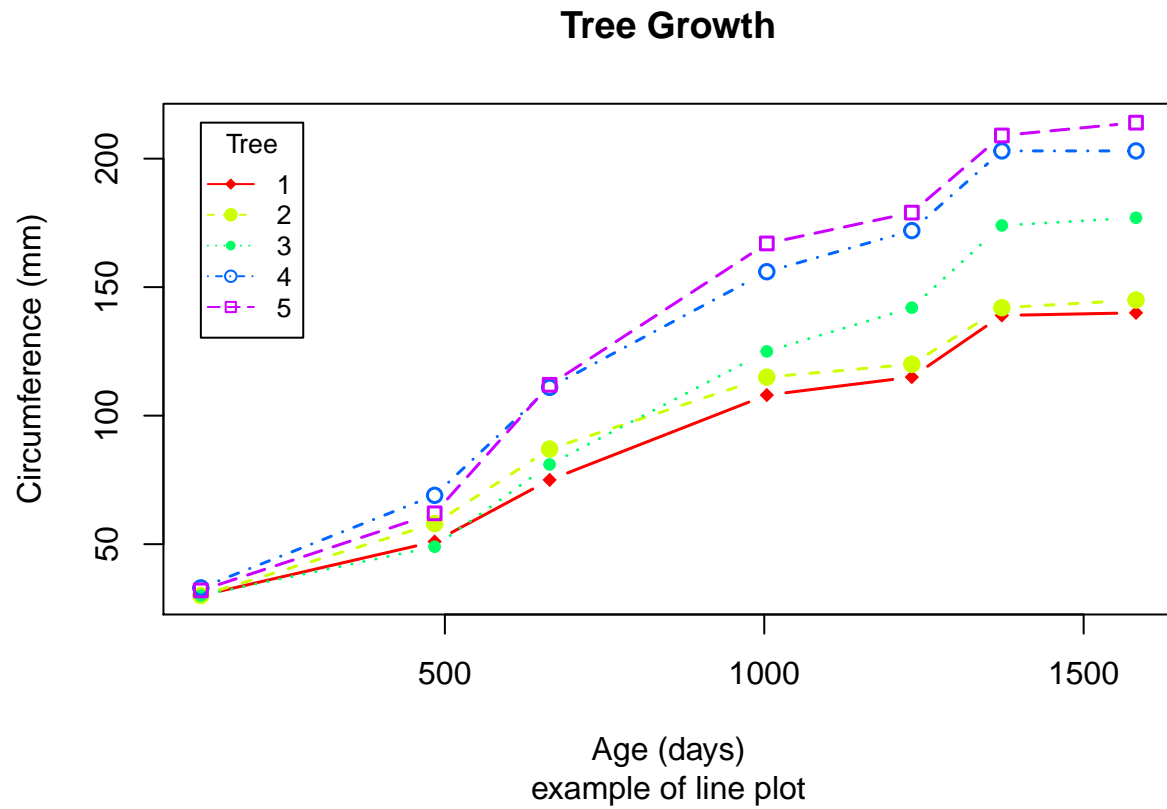
# get the range for the x and y axis
xrange <- range(Orange$age)
yrange <- range(Orange$circumference)

# set up the plot
plot(xrange, yrange, type="n", xlab="Age (days)",
     ylab="Circumference (mm)" )
colors <- rainbow(ntrees)
linetype <- c(1:ntrees)
plotchar <- seq(18,18+ntrees,1)

# add lines
for (i in 1:ntrees) {
  tree <- subset(Orange, Tree==i)
  lines(tree$age, tree$circumference, type="b", lwd=1.5,
        lty=linetype[i], col=colors[i], pch=plotchar[i])
}

# add a title and subtitle
title("Tree Growth", "example of line plot")
```

```
# add a legend  
legend(xrange[1], yrange[2], 1:ntrees, cex=0.8, col=colors,  
       pch=plotchar, lty=linetype, title="Tree")
```



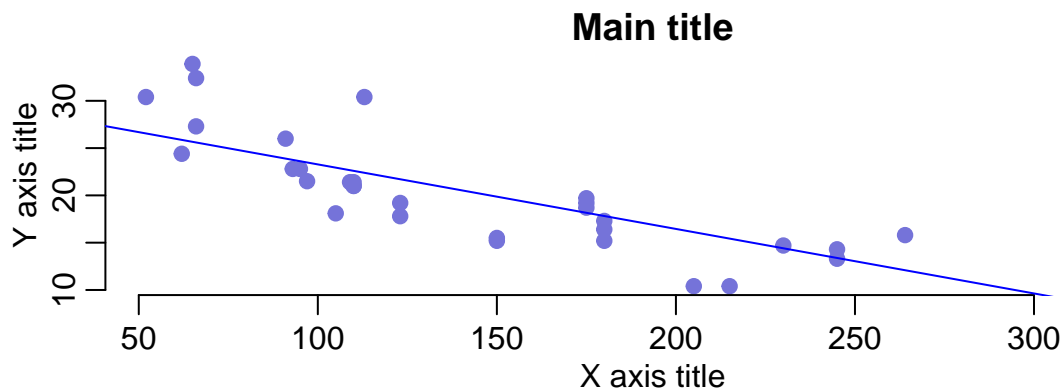
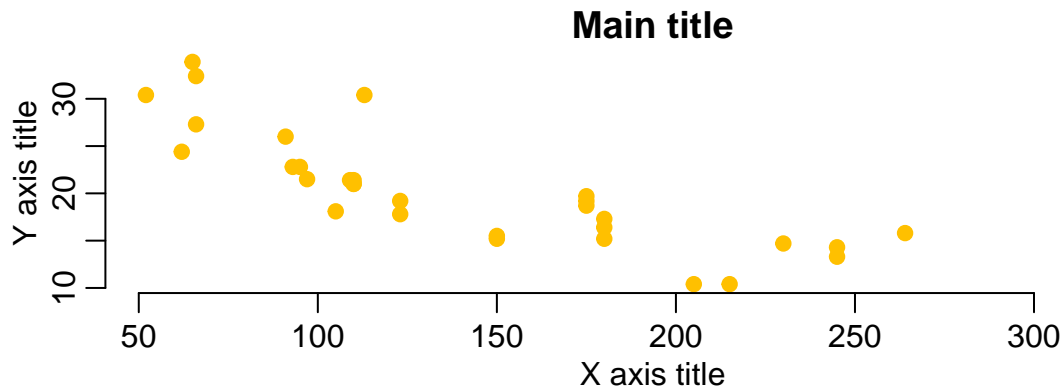
Chapter 7

Default Scatter Plots in R

R base scatter plot: `plot()`:

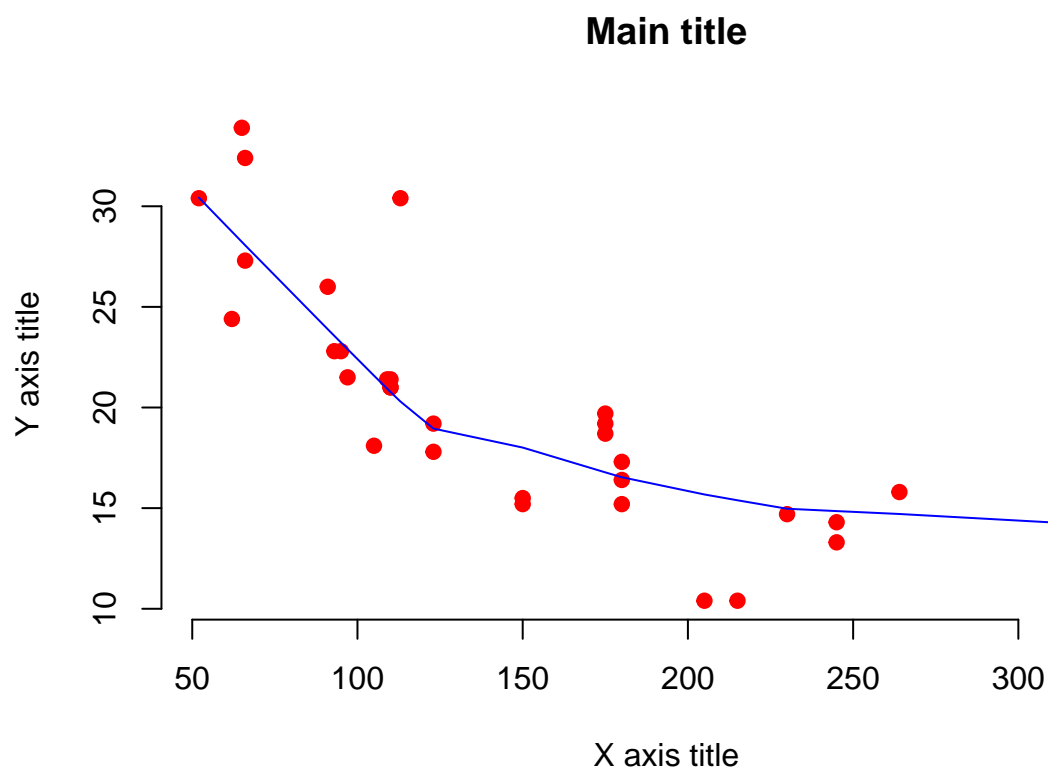
Example-1:

```
par(mfrow=c(2,1), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0))
x <- mtcars$hp
y <- mtcars$mpg
# Plot with main and axis titles
# Change point shape (pch = 19) and remove frame.
plot(x, y, main = "Main title",
      xlab = "X axis title", ylab = "Y axis title",
      pch = 19, frame = FALSE,col="#FFC000")
# Add regression line
plot(x, y, main = "Main title",
      xlab = "X axis title", ylab = "Y axis title",
      pch = 19, frame = FALSE,col="#7573D9")
abline(lm(y ~ x, data = mtcars), col = "blue")
```



Example-2:

```
x <- mtcars$hp
y <- mtcars$mpg
# Add loess fit
plot(x, y, main = "Main title",
      xlab = "X axis title", ylab = "Y axis title",
      pch = 19, frame = FALSE, col = "red")
lines(lowess(x, y), col = "blue")
```



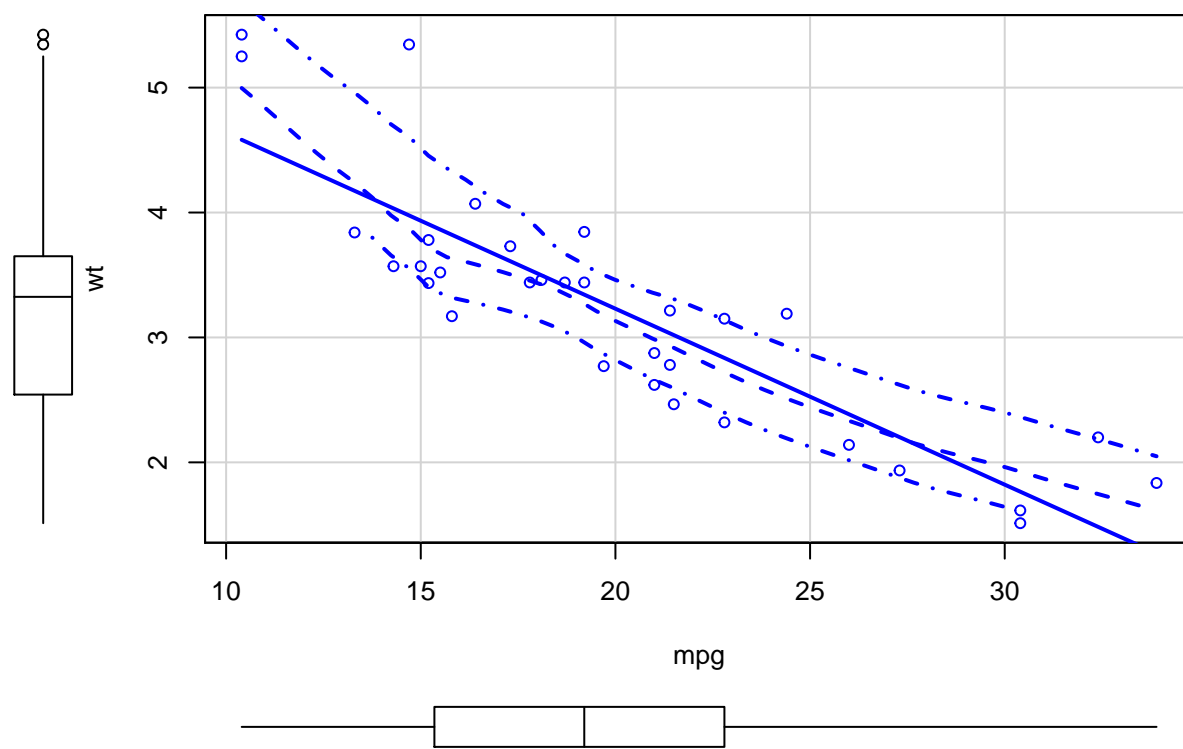
Additional->Enhanced scatter plots:

The components of the plot contains:

- the points
- the regression line (in green)
- the smoothed conditional spread (in red dashed line)
- the non-parametric regression smooth (solid line, red)

```
#install.packages("car")
library("car")
```

```
## Loading required package: carData
scatterplot(wt ~ mpg, data = mtcars)
```

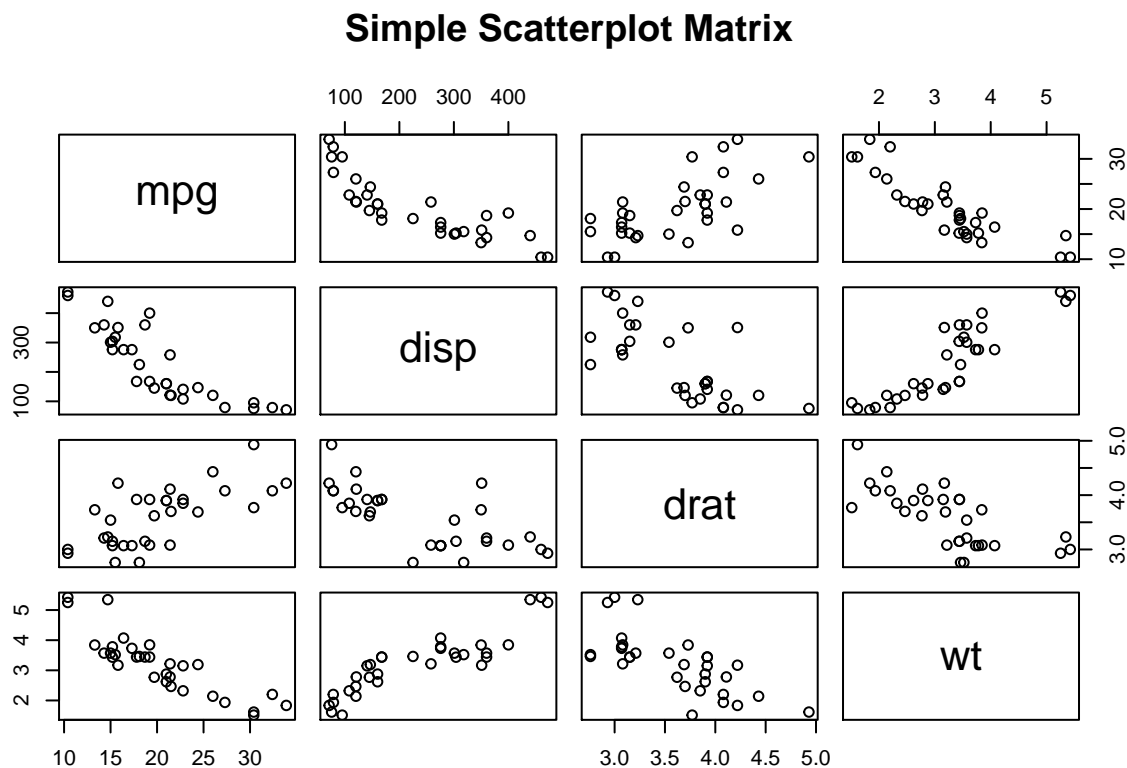


Chapter 8

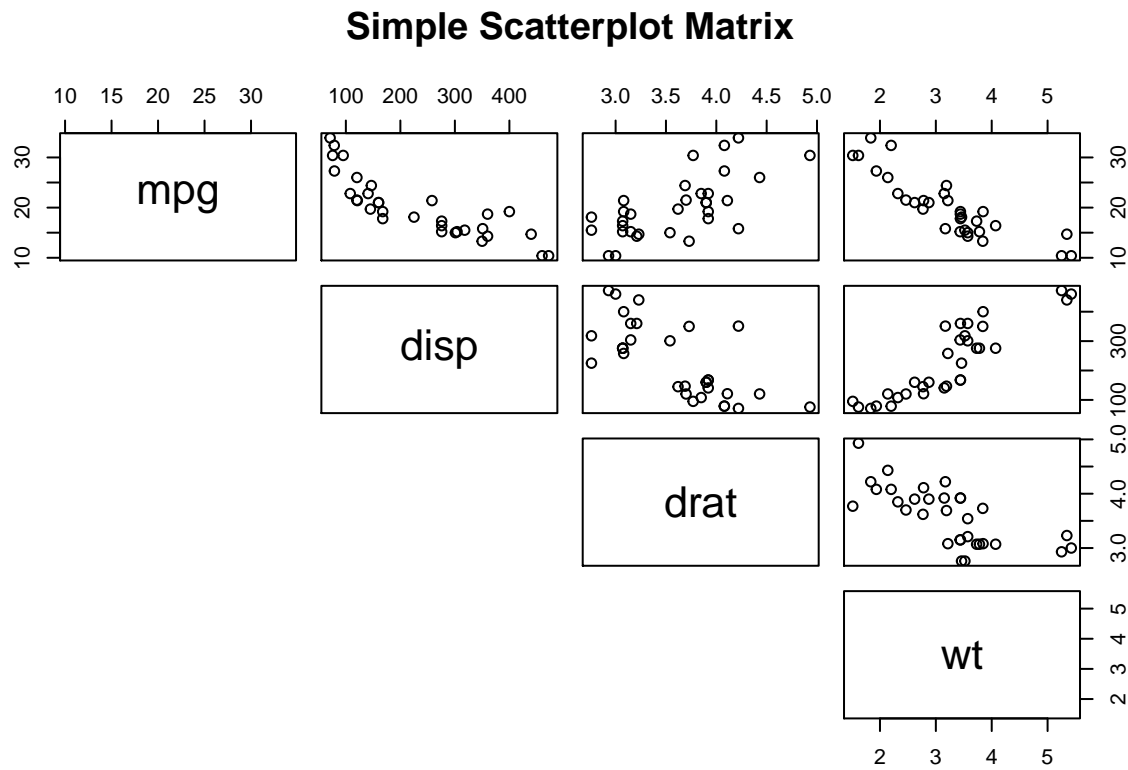
Default Scatter Plot Matrices in R

Scatter Plot Matrices:

```
par(mfrow=c(2,1))  
pairs(~mpg+disp+drat+wt,data=mtcars,  
      main="Simple Scatterplot Matrix")
```

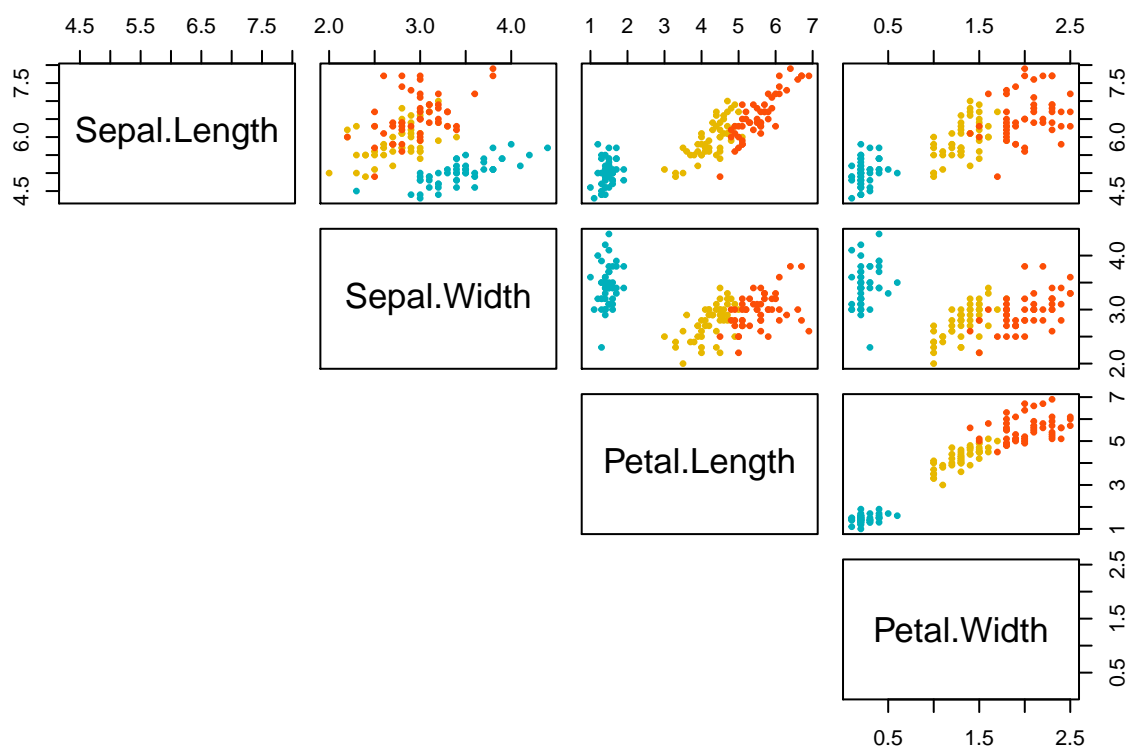


```
pairs(~mpg+disp+drat+wt,data=mtcars,lower.panel = NULL,  
      main="Simple Scatterplot Matrix")
```



Color points by groups Scatter plots:

```
my_col <- c("#00AFBB", "#E7B800", "#FC4E07")
pairs(iris[,1:4], pch = 19, cex = 0.5,
      col = my_col[iris$Species],
      lower.panel=NULL)
```

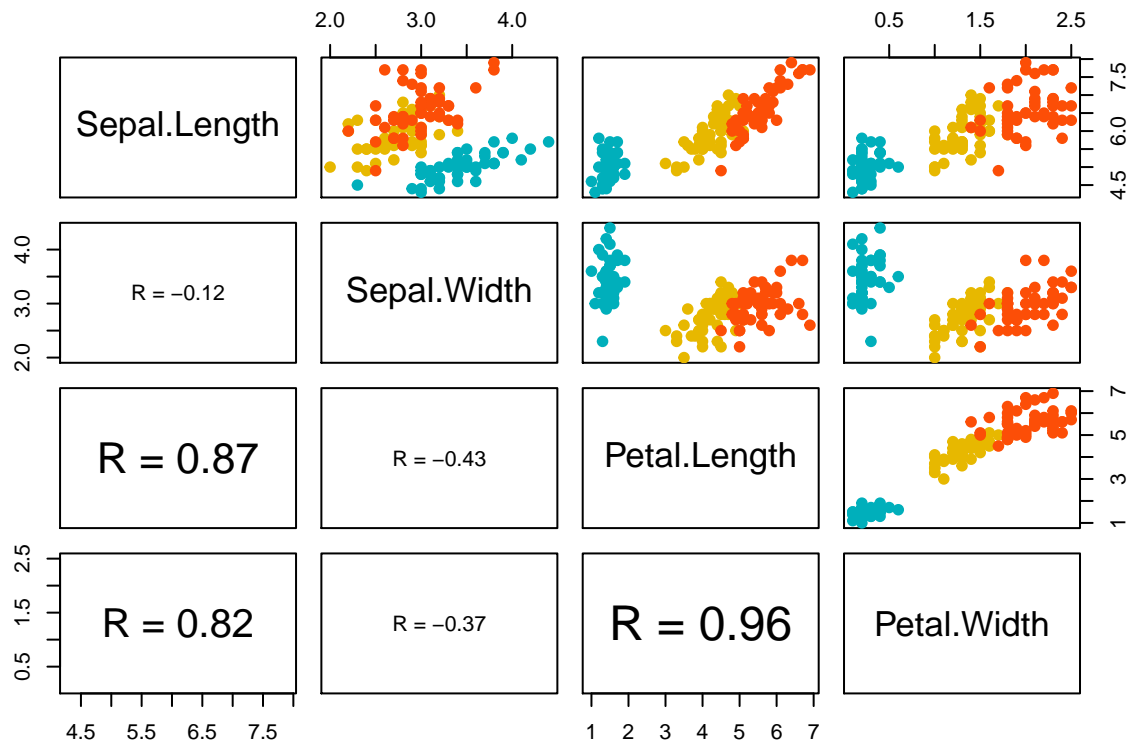



Add correlations on the lower panels:

```
# Correlation panel
my_col <- c("#00AFBB", "#E7B800", "#FC4E07")
panel.cor <- function(x, y){
  usr <- par("usr"); on.exit(par(usr))
  par(usr = c(0, 1, 0, 1))
  r <- round(cor(x, y), digits=2)
  txt <- paste0("R = ", r)
  cex.cor <- 0.8/strwidth(txt)
  text(0.5, 0.5, txt, cex = cex.cor * r)
}

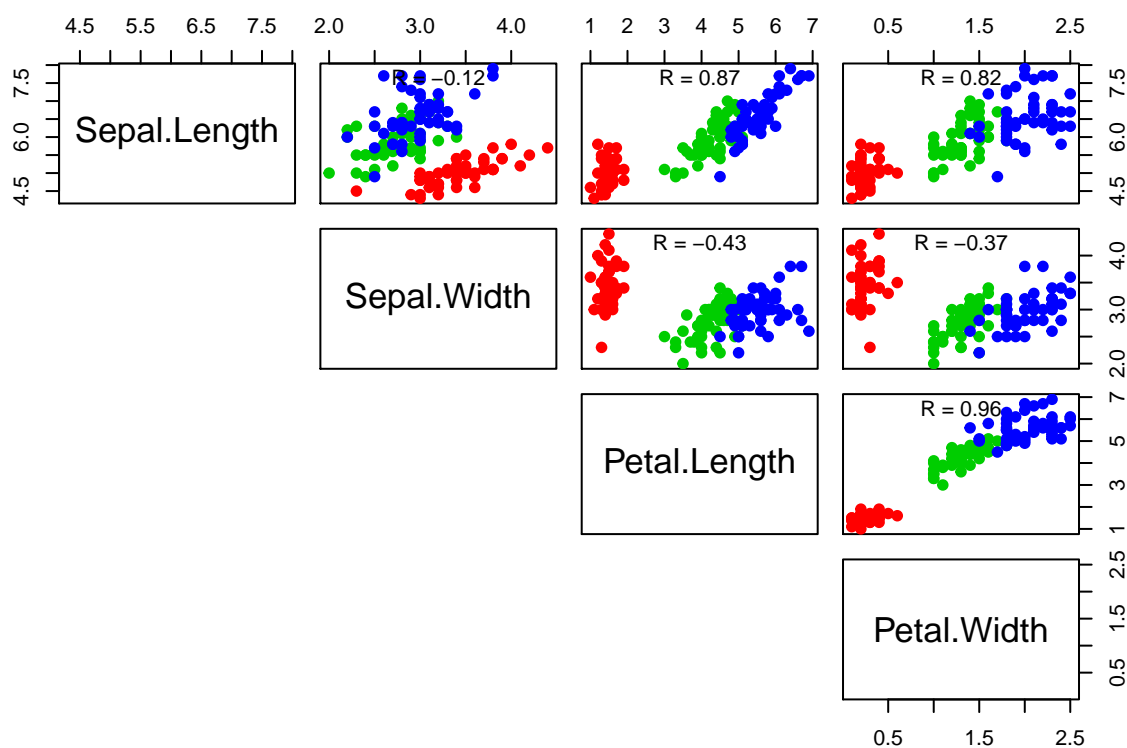
# Customize upper panel
upper.panel<-function(x, y){
  points(x,y, pch = 19, col = my_col[iris$Species])
}

# Create the plots
pairs(iris[,1:4],
      lower.panel = panel.cor,
      upper.panel = upper.panel)
```



Add correlations on the scatter plots:

```
my_col <- c("#00AFBB", "#E7B800", "#FC4E07")
# Customize upper panel
upper.panel<-function(x, y){
  points(x,y, pch=19, col=c("red", "green3", "blue")[iris$Species])
  r <- round(cor(x, y), digits=2)
  txt <- paste0("R = ", r)
  usr <- par("usr"); on.exit(par(usr))
  par(usr = c(0, 1, 0, 1))
  text(0.5, 0.9, txt)
}
pairs(iris[,1:4], lower.panel = NULL,
      upper.panel = upper.panel)
```



Additional-> the R package psych:

```
#install.packages("psych")
library(psych)
```

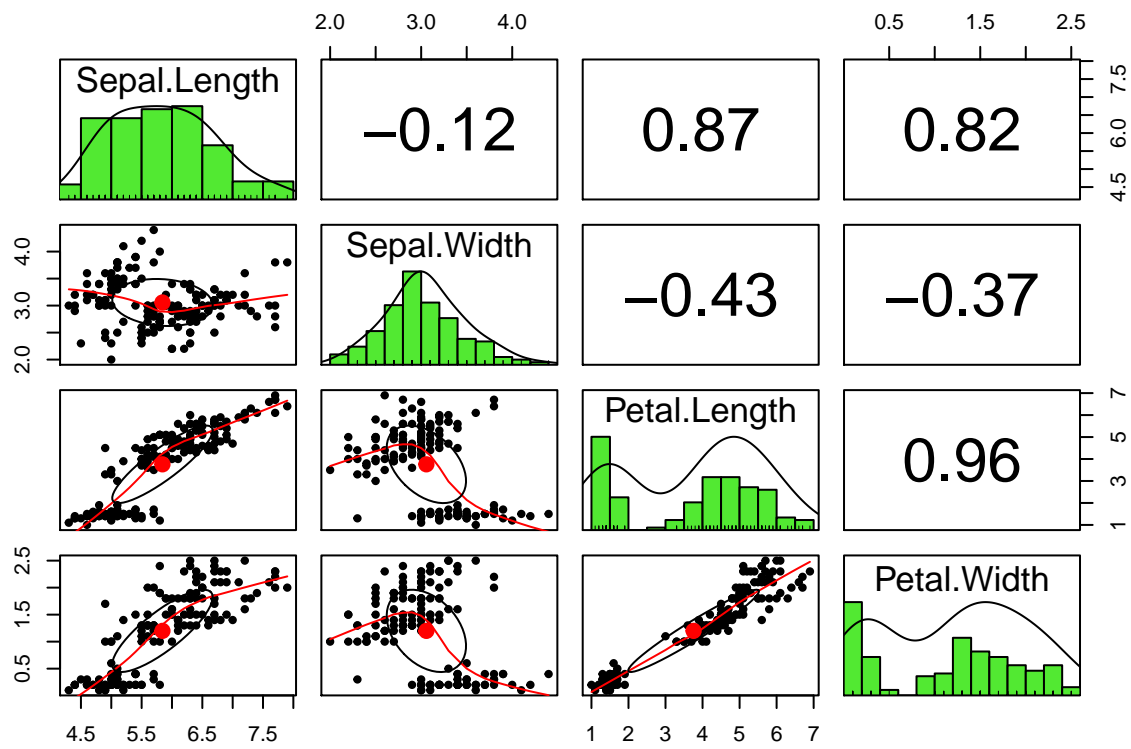
```
##
## Attaching package: 'psych'
```

```
## The following object is masked from 'package:car':
```

```
##
```

```
## logit
```

```
pairs.panels(iris[,-5],
  method = "pearson", # correlation method
  hist.col = "#52E932",
  density = TRUE, # show density plots
  ellipses = TRUE # show correlation ellipses
)
```



Chapter 9

Strip charts:1-D scatter plots

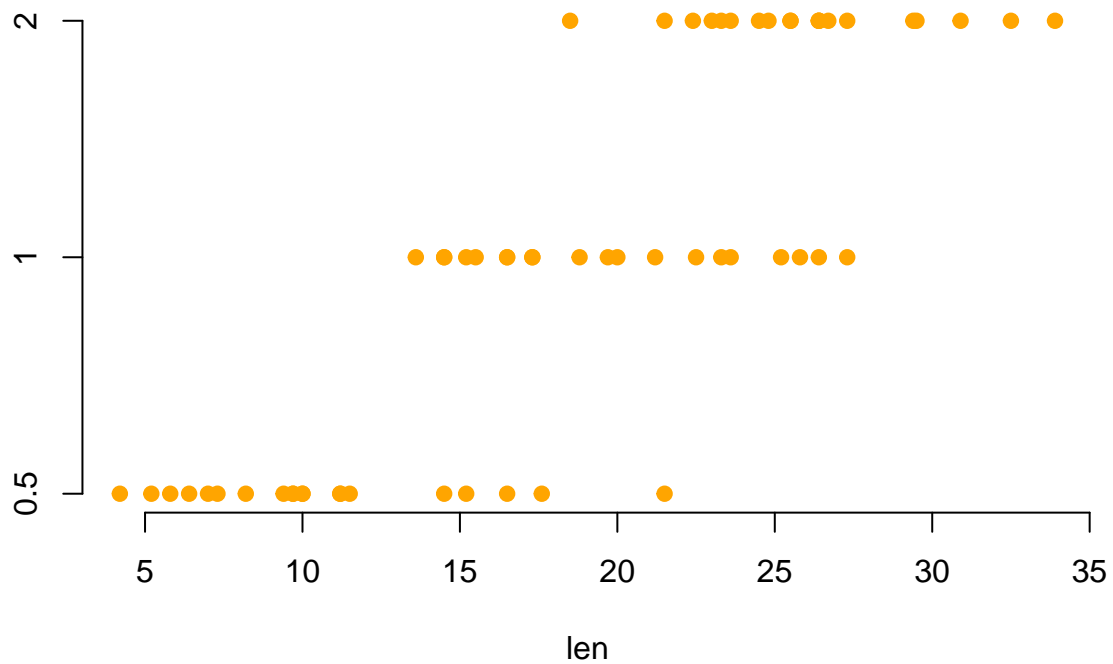
Formula:

```
stripchart(x, data = NULL method = "overplot", jitter = 0.1)
```

- x: the data from which the plots are to be produced. Allowed values are one or a list of numeric vector, each corresponding to a component plot.
- data: a data.frame (or list) from which the variables in x should be taken.
- Method: the method to be used to separate coincident points. Allowed values are one of "overplot", "jitter" or "stack".
- jitter: when method = "jitter" is used, jitter gives the amount of jittering applied.

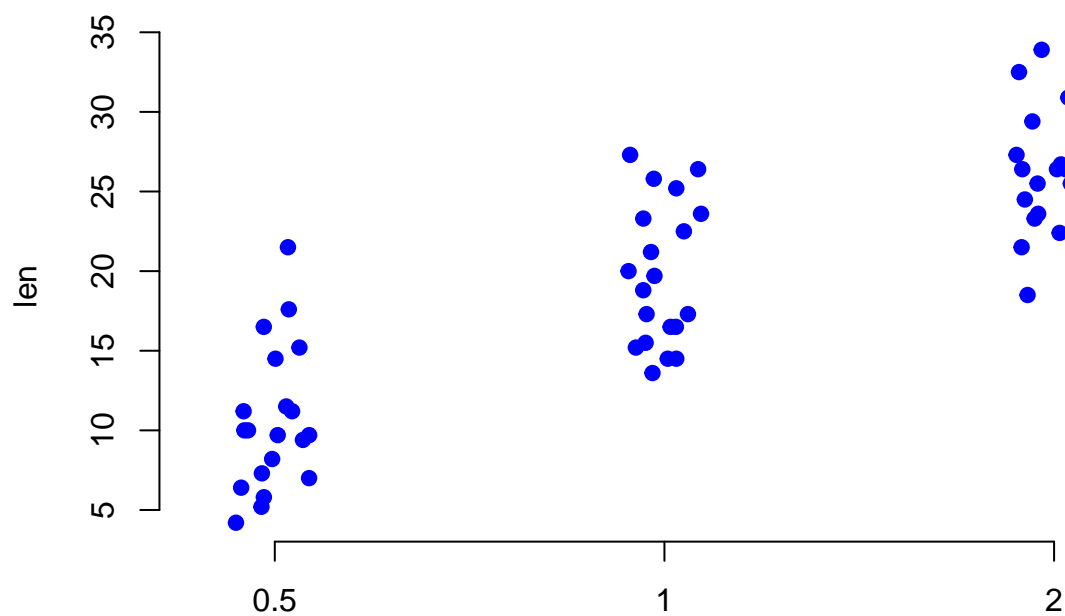
Example-1:

```
# Plot len by dose
stripchart(len ~ dose, data = ToothGrowth,
           pch = 19, frame = FALSE,col="orange")
```



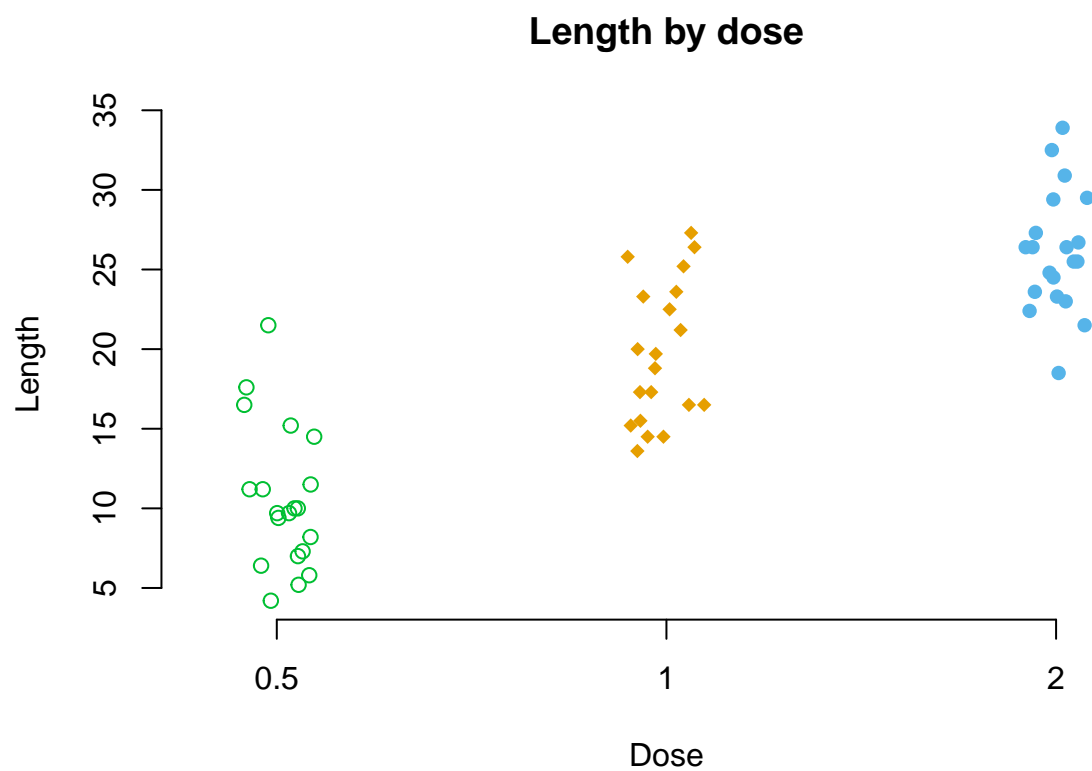
Example-2:

```
# Vertical plot using method = "jitter"
stripchart(len ~ dose, data = ToothGrowth,
  pch = 19, frame = FALSE, vertical = TRUE,
  method = "jitter", col="blue")
```



Example-3:

```
# Change point shapes (pch) and colors by groups
# add main title and axis labels
stripchart(len ~ dose, data = ToothGrowth,
  frame = FALSE, vertical = TRUE,
  method = "jitter", pch = c(21, 18, 16),
  col = c("#00BF32", "#E69F00", "#56B4E9"),
  main = "Length by dose", xlab = "Dose", ylab = "Length")
```



Chapter 10

Default Dot Plots in R

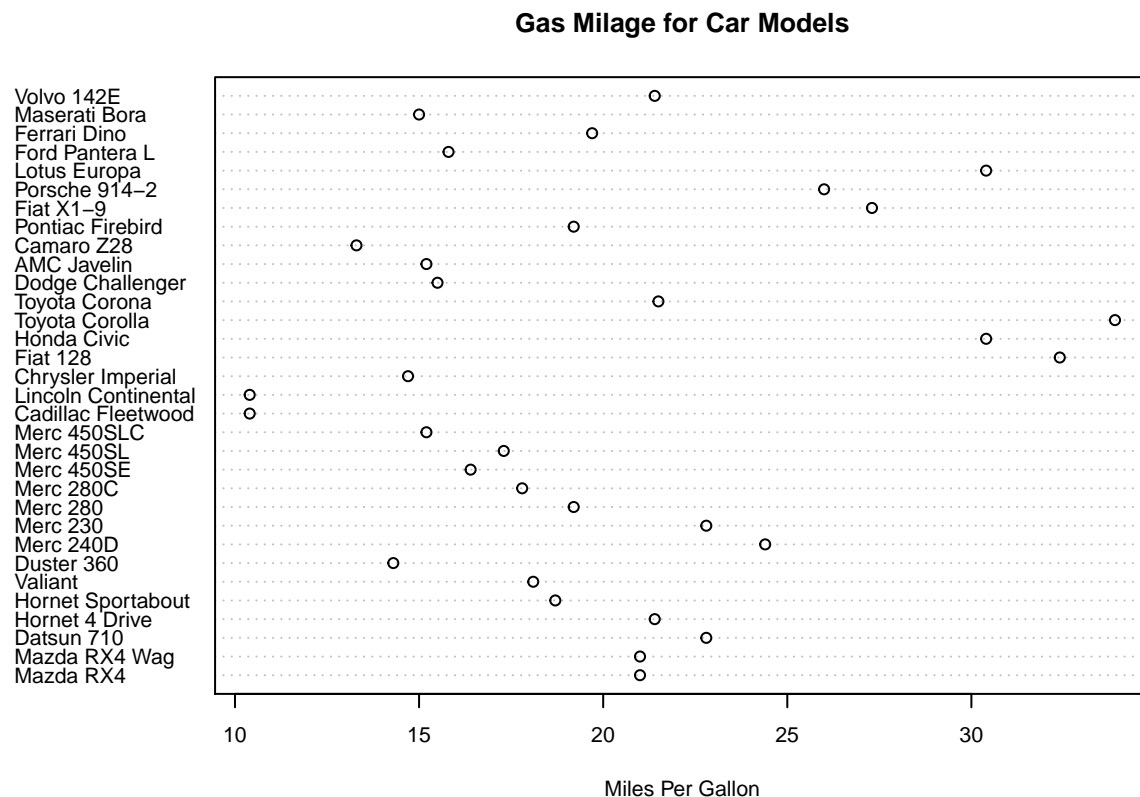
Formula:

```
dotchart(x, labels = NULL, groups = NULL, gcolor = par("fg"), color = par("fg"))
```

Simple Dot Plots numeric vectors:

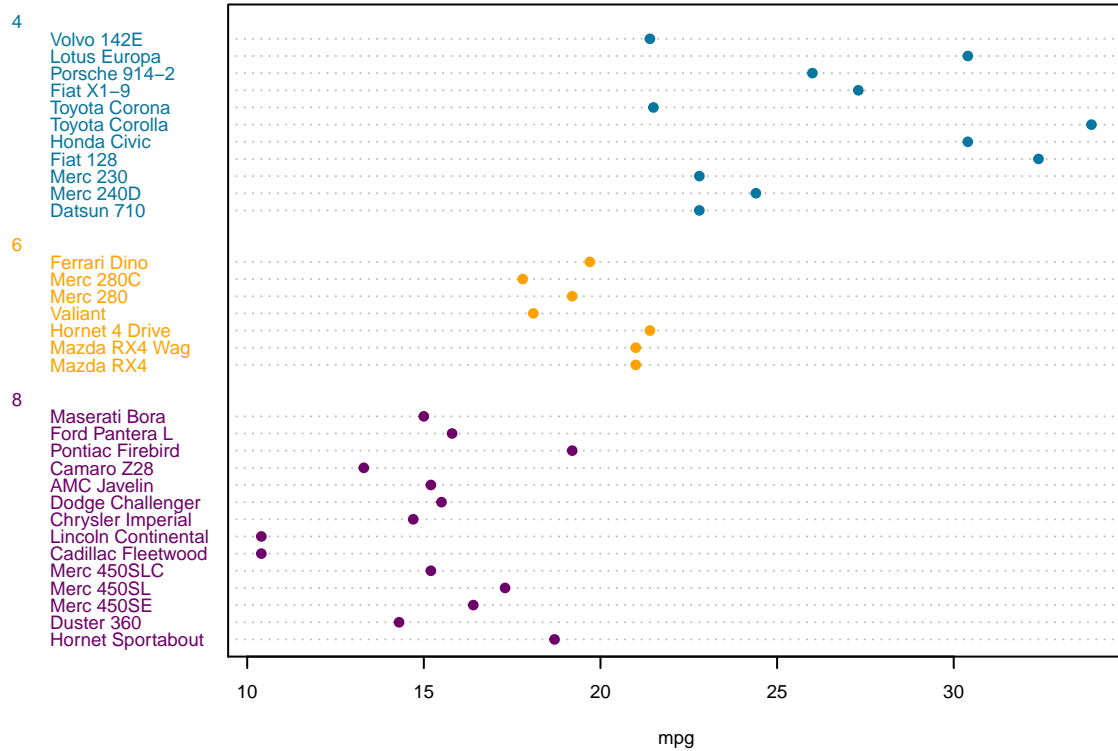
Example-1:

```
# Simple Dot Plot
dotchart(mtcars$mpg, labels=row.names(mtcars), cex=.7,
  main="Gas Milage for Car Models",
  xlab="Miles Per Gallon")
```



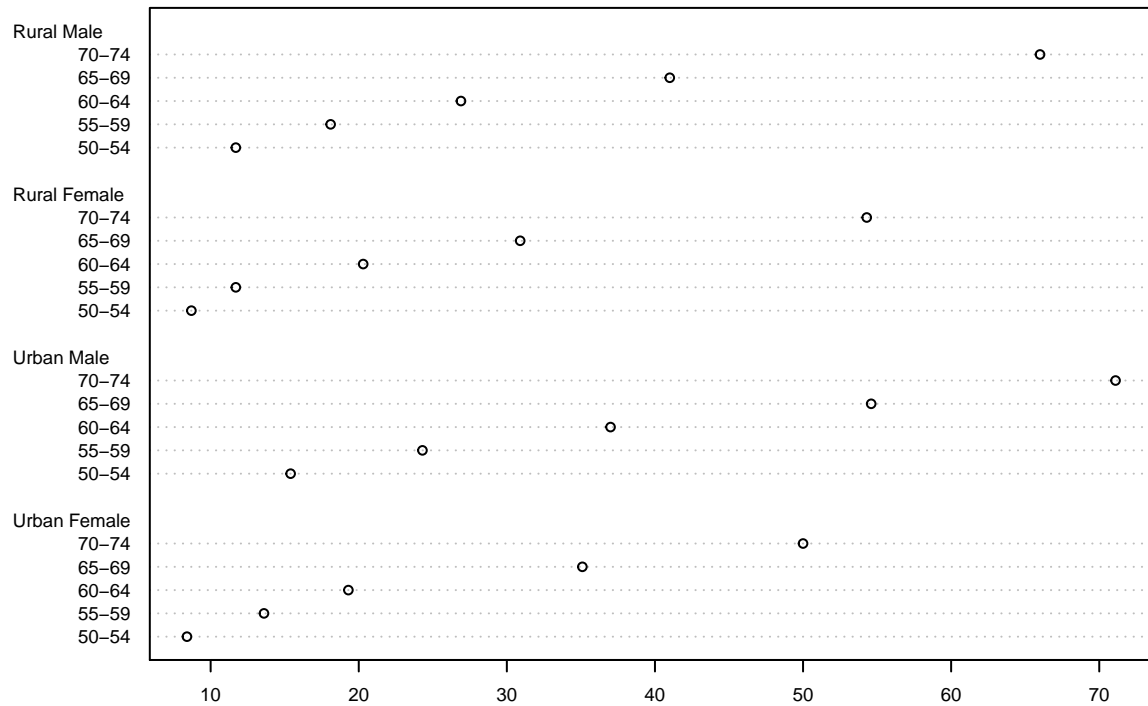
Example-2:

```
# Plot and color by groups cyl
grps <- as.factor(mtcars$cyl)
my_cols <- c("#0776A0", "#FFA100", "#6E0069")
dotchart(mtcars$mpg, labels = row.names(mtcars),
         groups = grps, gcolor = my_cols,
         color = my_cols[grps],
         cex = 0.6, pch = 19, xlab = "mpg")
```



Dot Plot matrix:

```
dotchart(VADeaths, cex = 0.6,
         main = "Death Rates in Virginia - 1940")
```

Death Rates in Virginia – 1940

Chapter 11

Default Pie Charts in R

Formula:

```
pie(x, labels = names(x), radius = 0.8)
```

- x: a vector of non-negative numerical quantities. The values in x are displayed as the areas of pie slices.
- labels: character strings giving names for the slices.
- radius: radius of the pie circle. If the character strings labeling the slices are long it may be necessary to use a smaller radius.

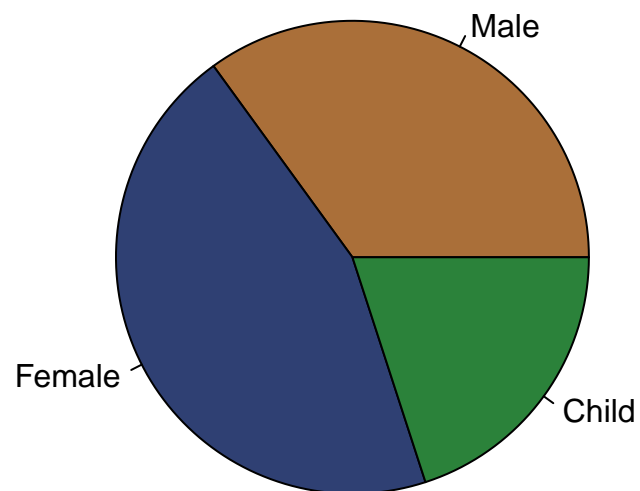
Basic pie charts: -> `pie()`

Example-1:

```
df <- data.frame(  
  group = c("Male", "Female", "Child"),  
  value = c(35, 45, 20)  
)  
df
```

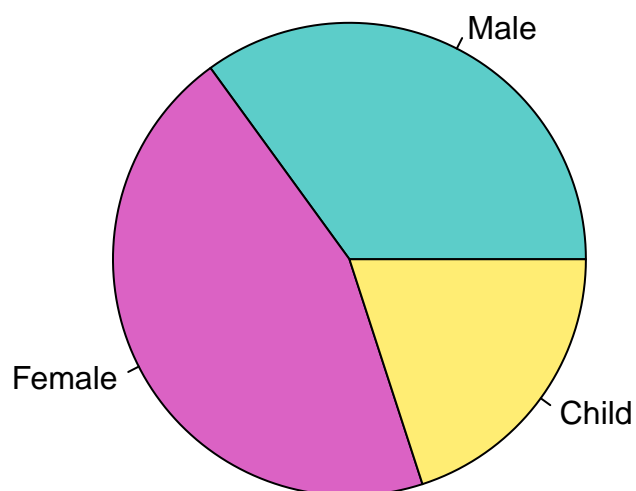
```
##      group value  
## 1   Male     35  
## 2 Female     45  
## 3  Child     20
```

```
pie(df$value, labels = df$group, radius = 1, col=c("#AA6F39", "#2F4073", "#2B823A"))
```



Example-2:

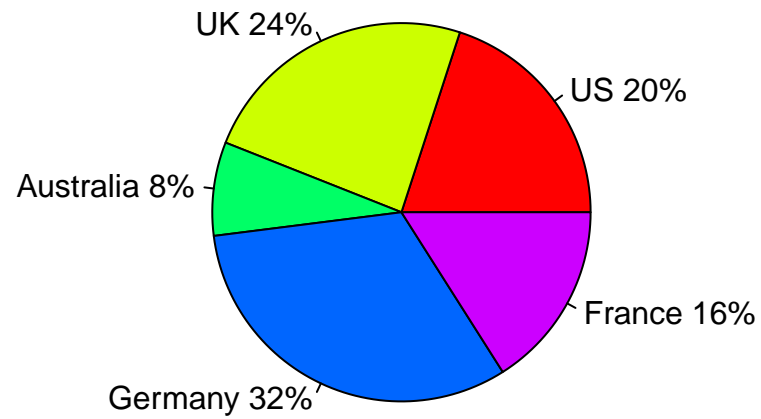
```
# Change colors
pie(df$value, labels = df$group, radius = 1,
    col = c("#5CCDC9", "#DB62C4", "#FFED73"))
```



Example-3:

```
# Pie Chart with Percentages
slices <- c(10, 12, 4, 16, 8)
lbls <- c("US", "UK", "Australia", "Germany", "France")
pct <- round(slices/sum(slices)*100)
lbls <- paste(lbls, pct) # add percents to labels
lbls <- paste(lbls,"%",sep="") # ad % to labels
pie(slices,labels = lbls, col=rainbow(length(lbls)),
    main="Pie Chart of Countries")
```

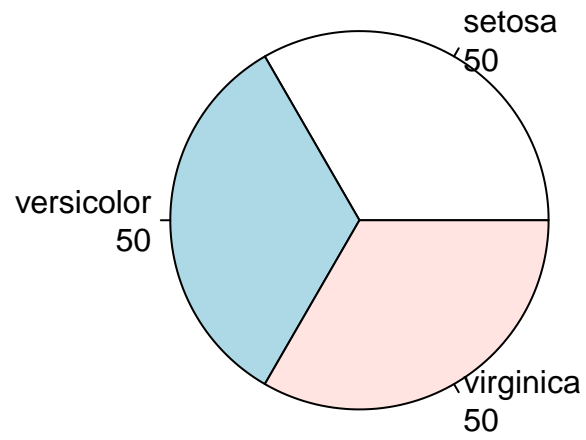
Pie Chart of Countries



Example-4:

```
# Pie Chart from data frame with Appended Sample Sizes
mytable <- table(iris$Species)
lbls <- paste(names(mytable), "\n", mytable, sep="")
pie(mytable, labels = lbls,
    main="Pie Chart of Species\n (with sample sizes)")
```


**Pie Chart of Species
(with sample sizes)**



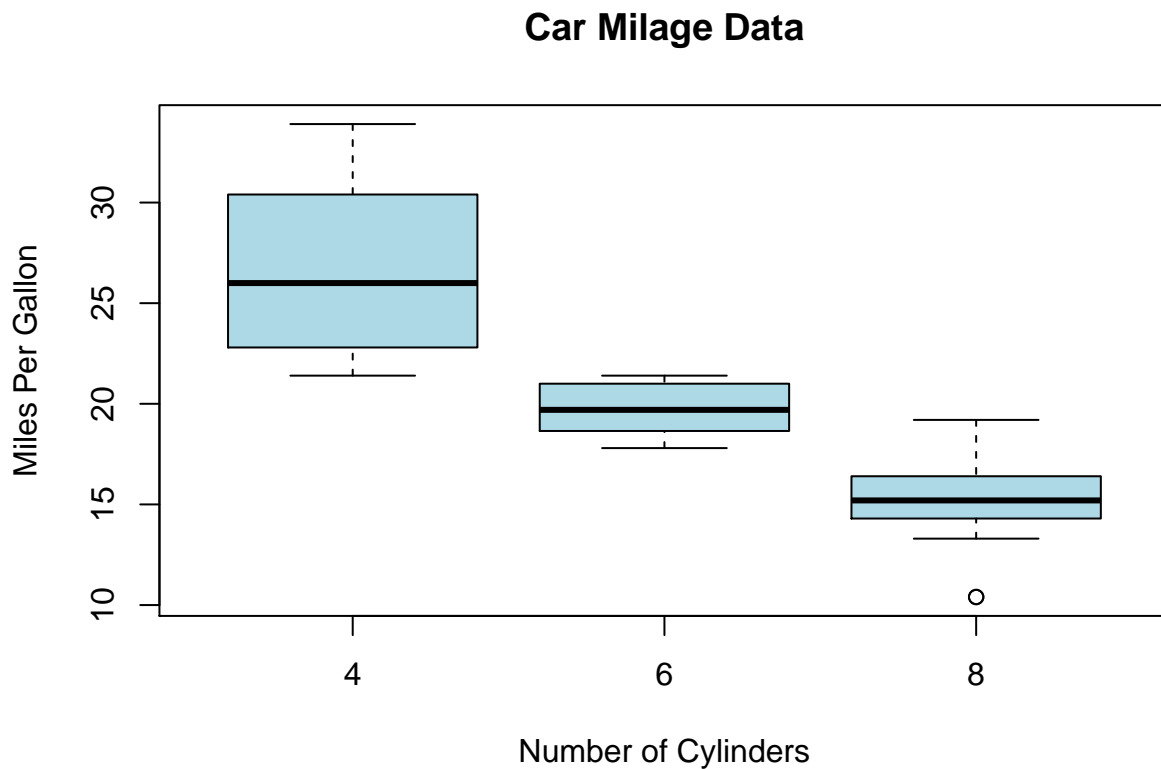
Chapter 12

Default Box Plots in R

Basic box plots:

Example-1:

```
# Boxplot of MPG by Car Cylinders  
boxplot(mpg~cyl,data=mtcars, main="Car Milage Data",  
        xlab="Number of Cylinders", ylab="Miles Per Gallon",col="lightblue")
```



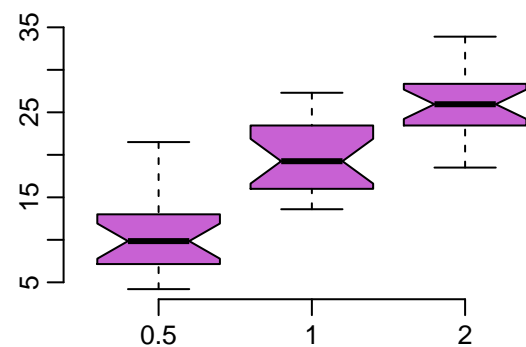
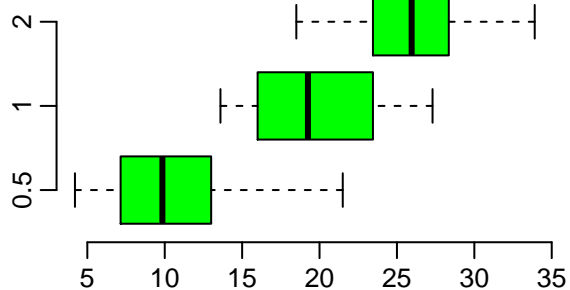
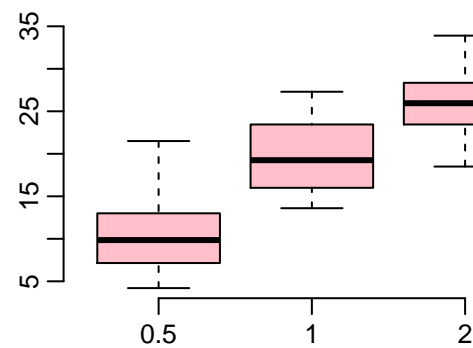
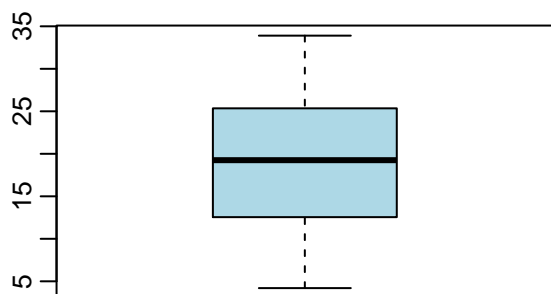
Example-2:

```
par(mfrow=c(2,2), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0))  
# Box plot of one variable
```

```

boxplot(ToothGrowth$len,col="lightblue")
# Box plots by groups (dose)
# remove frame
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,col="pink")
# Horizontal box plots
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        horizontal = TRUE,col="green")
# Notched box plots
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        notch = TRUE,col="#C861D3")

```

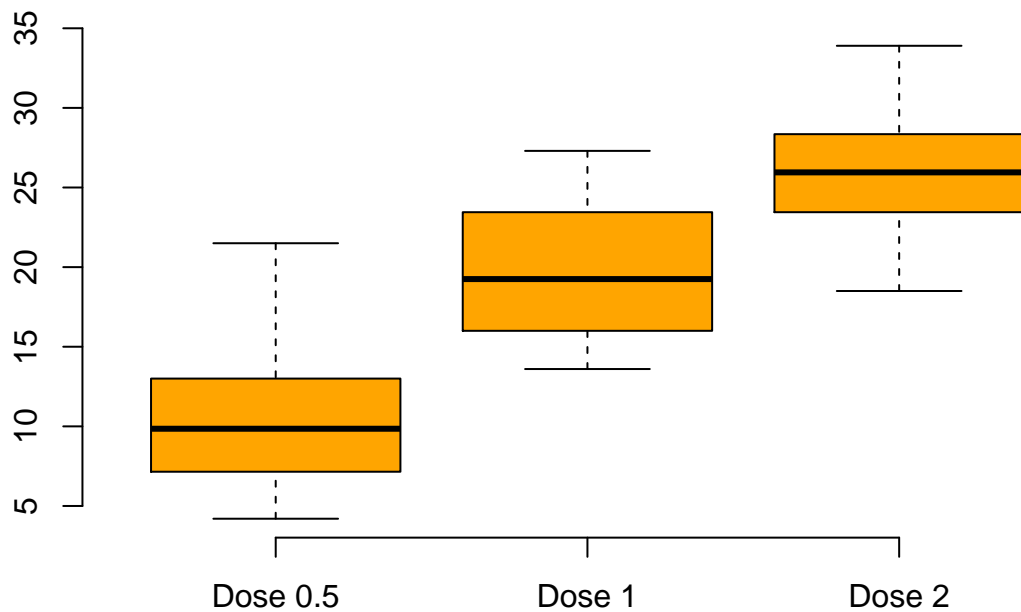


Change group names:

```

boxplot(ToothGrowth$len ~ ToothGrowth$dose, data = ToothGrowth, frame = FALSE,
        names = c("Dose 0.5", "Dose 1", "Dose 2"),col="orange")

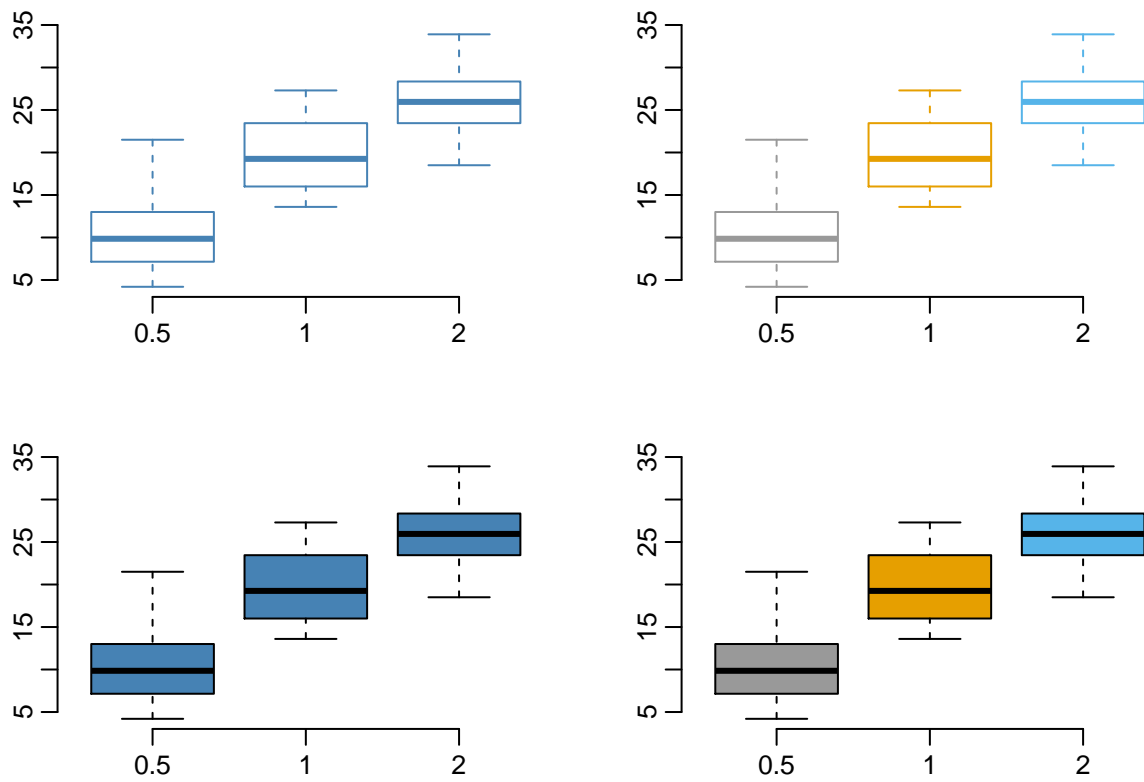
```



Change color:

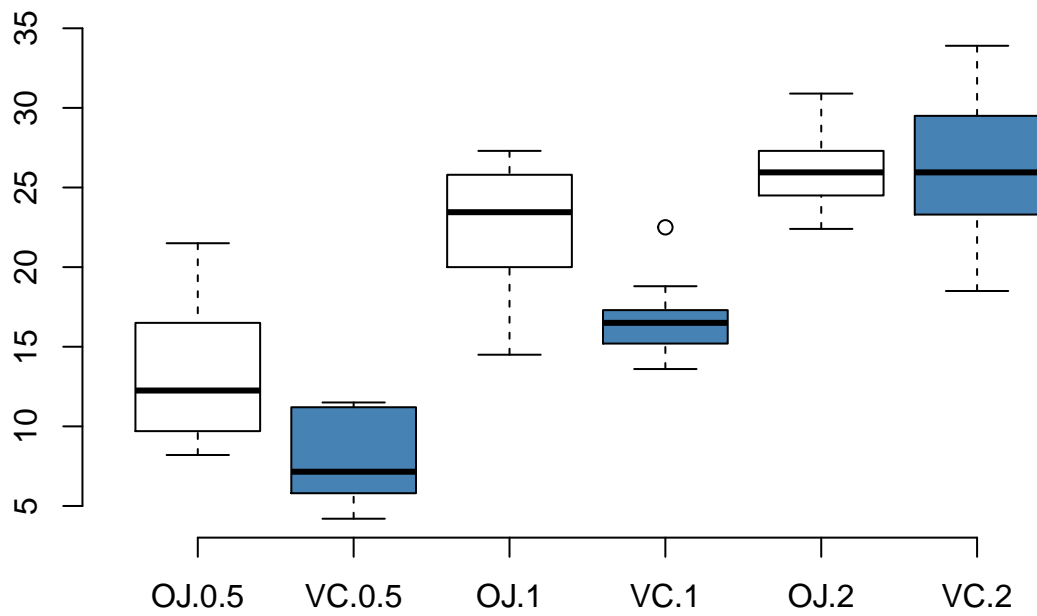
```
par(mfrow=c(2,2), mar=c(3,3,1,0)+.5, mgp=c(1.6,.6,0))

# Change the color of border using one single color
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        border = "steelblue")
# Change the color of border.
# Use different colors for each group
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        border = c("#999999", "#E69F00", "#56B4E9"))
# Change fill color : single color
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        col = "steelblue")
# Change fill color: multiple colors
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        col = c("#999999", "#E69F00", "#56B4E9"))
```



Multiple box plots:

```
boxplot(len ~ supp*dose, data = ToothGrowth,
        col = c("white", "steelblue"), frame = FALSE)
```

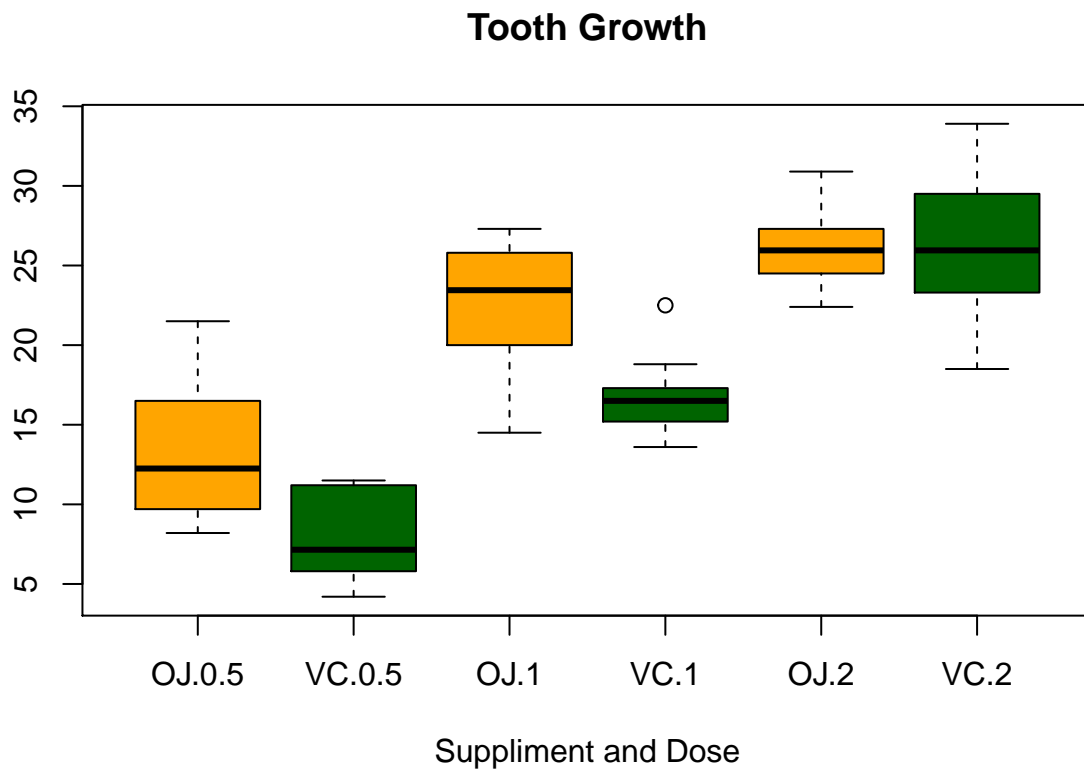


Main title and axis labels:

```
# Change axis titles
# Change color (col = "gray") and remove frame
# Create notched box plot
boxplot(len ~ dose, data = ToothGrowth,
        main = "Plot of length by dose",
        xlab = "Dose (mg)", ylab = "Length",
        col = "lightgray", frame = FALSE)
```

**Notched Boxplot:**

```
# Notched Boxplot of Tooth Growth Against 2 Crossed Factors  
# boxes colored for ease of interpretation  
boxplot(len~supp*dose, data=ToothGrowth, notch=FALSE,  
        col=c("orange","darkgreen")),  
        main="Tooth Growth", xlab="Suppliment and Dose")
```

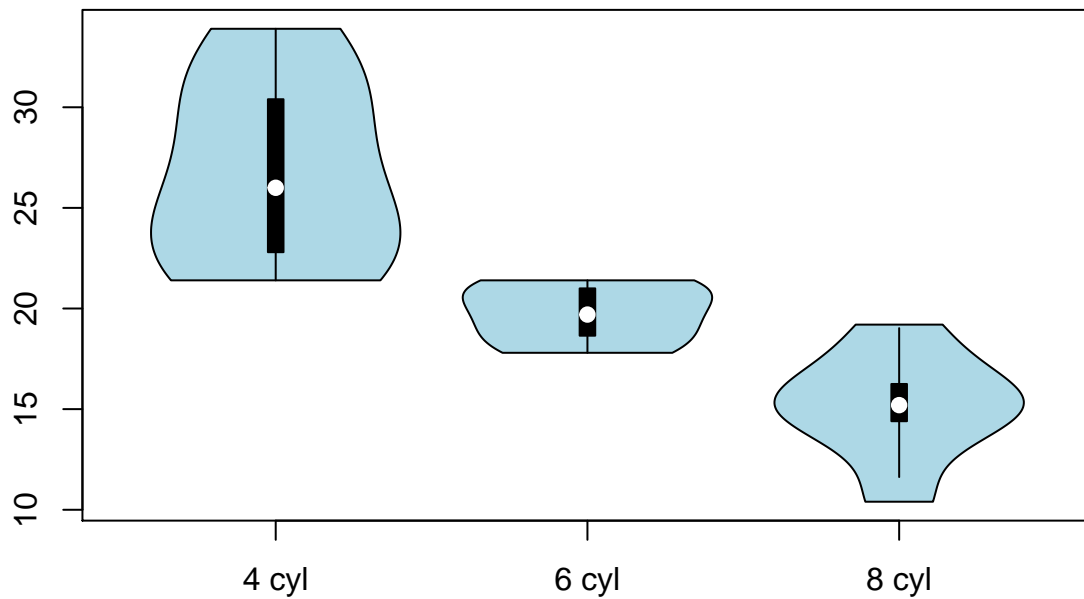
Additional box plots->

- `vioplot` function from **vioplot** package
- `bagplot(x, y)` function from **aplpack** package

Example-1:

```
# Violin Plots
#install.packages("vioplot")
library(vioplot)
x1 <- mtcars$mpg[mtcars$cyl==4]
x2 <- mtcars$mpg[mtcars$cyl==6]
x3 <- mtcars$mpg[mtcars$cyl==8]
vioplot(x1, x2, x3, names=c("4 cyl", "6 cyl", "8 cyl"),
        col="lightblue")
title("Violin Plots of Miles Per Gallon")
```

Violin Plots of Miles Per Gallon

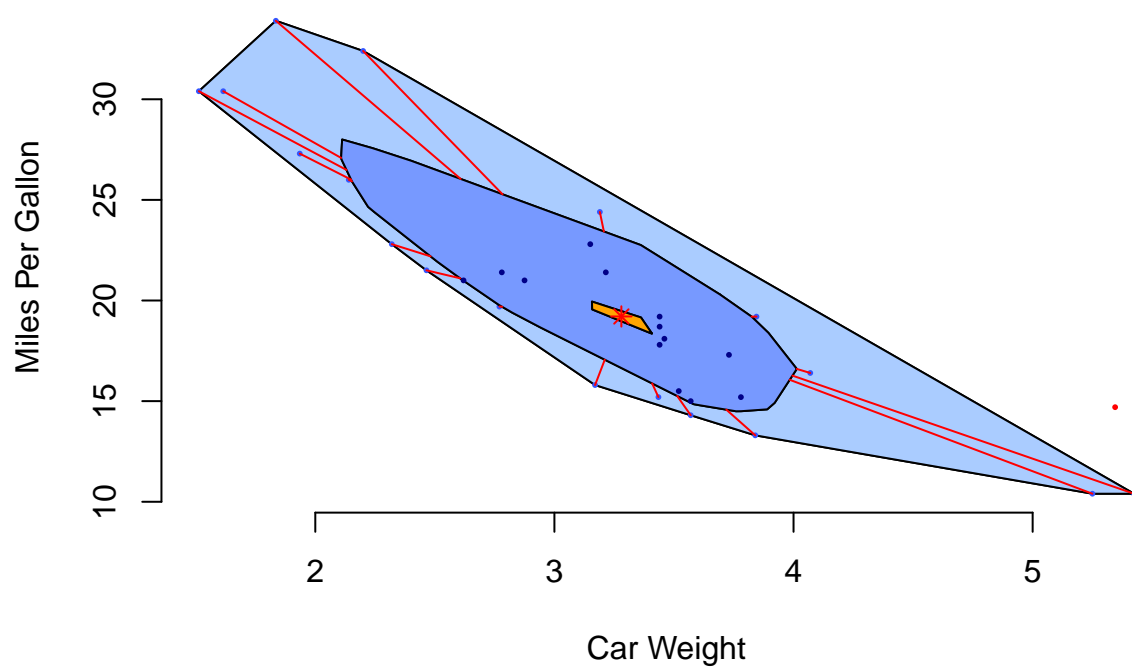


Example-2:

```
# Example of a Bagplot  
#install.packages("aplpack")  
library(aplpack)
```

```
## Loading required package: tcltk  
attach(mtcars)  
bagplot(wt,mpg, xlab="Car Weight", ylab="Miles Per Gallon",  
        main="Bagplot Example")
```

Bagplot Example

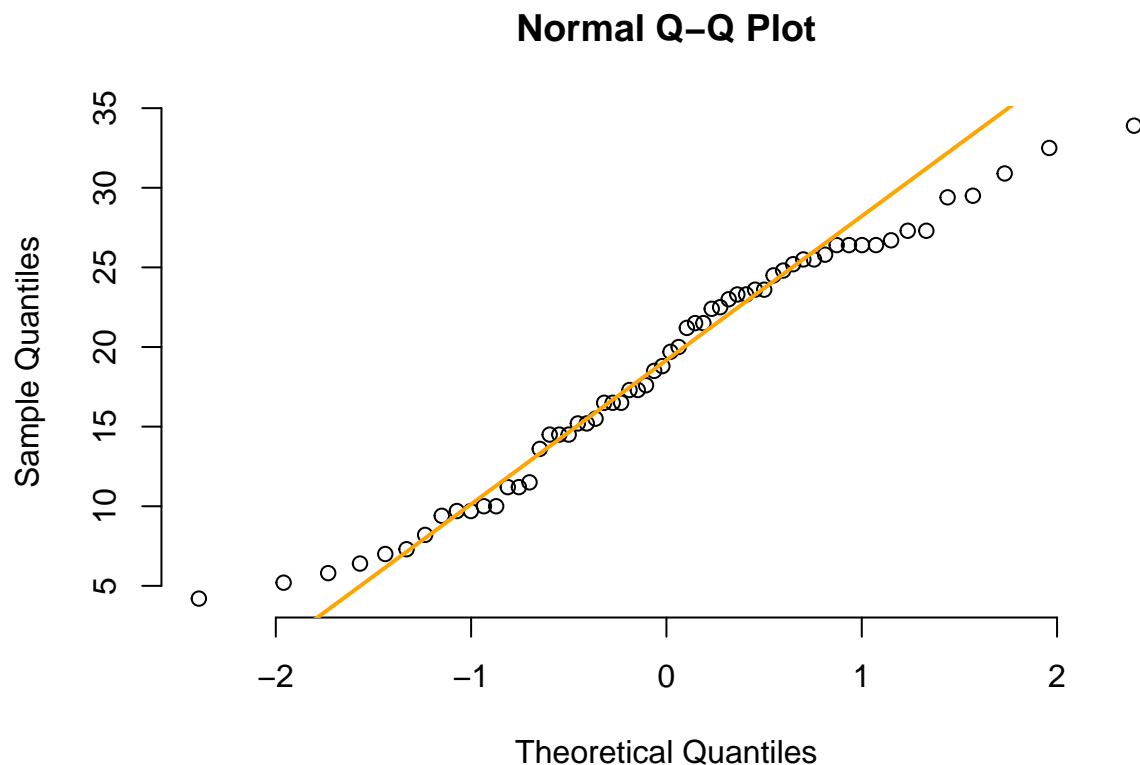


Chapter 13

QQ-Plots: Quantile-Quantile Plots

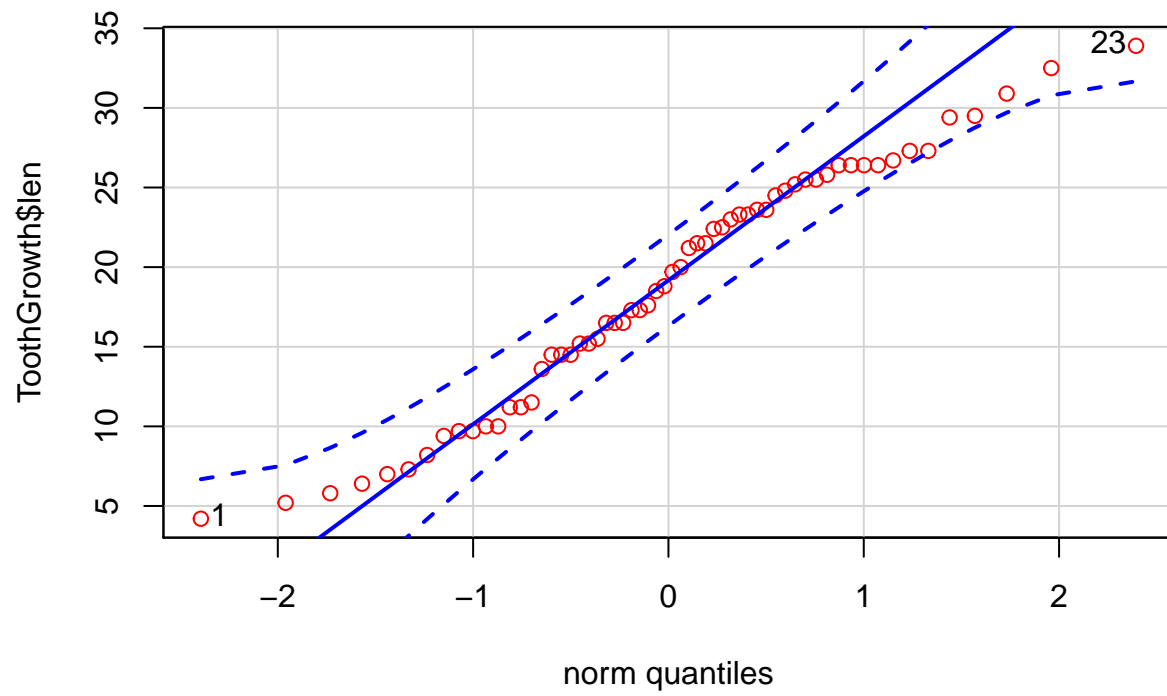
- `qqnorm()`: produces a normal QQ plot of the variable
- `qqline()`: adds a reference line

```
qqnorm(ToothGrowth$len, pch = 1, frame = FALSE)  
qqline(ToothGrowth$len, col = "orange", lwd = 2)
```



Additional -> `qqPlot()` in `car` package:

```
#install.packages("car")  
library("car")  
qqPlot(ToothGrowth$len, col="red")
```



```
## [1] 23 1
```

Chapter 14

Means and Confidence Intervals

With `plotmeans()` function from **gplots** package

```
#install.packages("gplots")  
library(gplots)
```

```
##  
## Attaching package: 'gplots'
```

```
## The following object is masked from 'package:stats':  
##  
##      lowess
```

```
# Plot the mean of teeth length by dose groups  
plotmeans(len ~ dose, data = ToothGrowth, frame = FALSE)
```

```
## Warning in plot.xy(xy.coords(x, y), type = type, ...): "frame" is not a  
## graphical parameter
```

```
## Warning in axis(1, at = 1:length(means), labels = legends, ...): "frame" is  
## not a graphical parameter
```

```
## Warning in plot.xy(xy.coords(x, y), type = type, ...): "frame" is not a  
## graphical parameter
```

