



R Graphics

An Introduction to Graphical Representation of Data using R

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Introduction

One of the most tedious things for beginners using R is to find out how to **configure graphs** properly. This manual is intended as an introduction to the vast possibilities R offers for the professional display of data.

The material is structured in **three chapters**:

- (1) Standard graphs, which is what everybody is using all the time
- (2) Trellis displays, with which an in-depth "dissection" of datasets is possible; data that are conventionally treated as "multivariate" can be visualized very conveniently.
- (3) Some additional graphics features from interesting R libraries; examples are shown for barplots, 3D surface plots and for the display of meteorological data.

Christoph Scherber,
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1. Standard plots

```
x<-c(1,3,4,6,8,9,12)
y<-c(5,8,6,10,9,13,12)
```

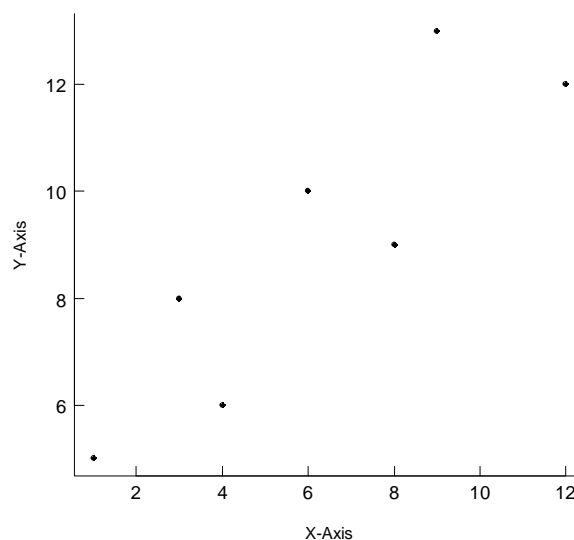
Quick plotting starts like this:
`plot(x,y)`

More advanced plotting, however, involves a step-wise approach:

- Set up an empty plotting region (`type="n"`)
- Draw the points
- Draw the axes
- Write the title(s)
- Add regression lines etc.

```
par(tck=0.02,las=1,cex.axis=1.3,cex.lab=1.2,mgp=c(3,0.5,0),bty="L")
```

```
x<-c(1,3,4,6,8,9,12)
y<-c(5,8,6,10,9,13,12)
plot(x,y,type="n",axes=F,xlab="",ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-Axis")
box()
```



Suppose now that we want to **add arrows to the axes**; there are, in principle, two possibilities for this. One is to use the

arrows() command (which doesn't look too good) - the second (a bit more complicated one) is to use polygon():

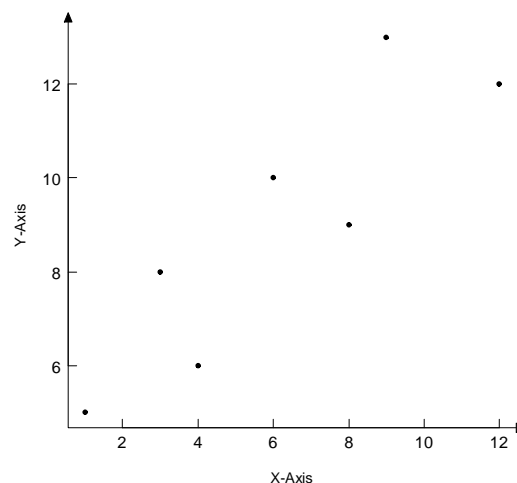
```
u <- par("usr")
u

# first possibility:
arrows(u[1], u[3], u[2], u[3], code = 2, xpd =
TRUE,angle=20,length=0.2)

arrows(u[1], u[3], u[1], u[4], code = 2, xpd =
TRUE,angle=20,length=0.2)

#second possibility:
polygon(c(u[2],u[2],u[2]+0.2),c(u[3]-
0.1,u[3]+0.1,u[3]),xpd=TRUE,col="black",lty=1)

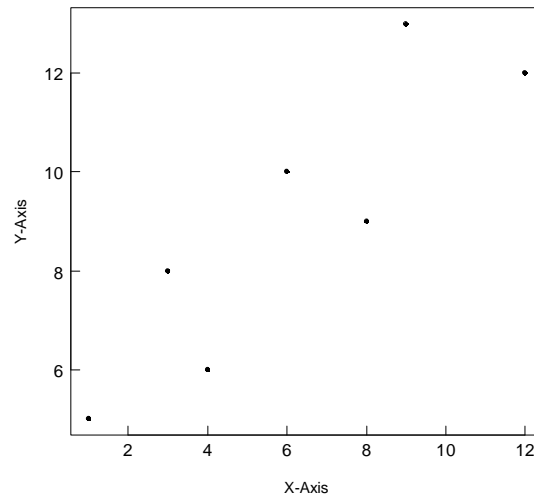
polygon(c(u[1]-
0.1,u[1]+0.1,u[1]),c(u[4],u[4],u[4]+0.2),xpd=TRUE,col="black",
lty=1)
```



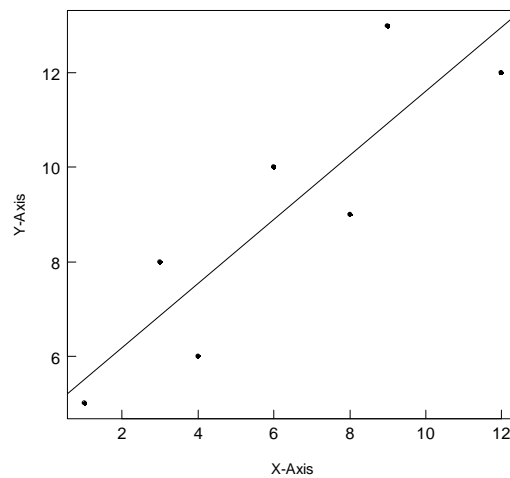
Usually, you wouldn't really need those arrowheads, so you just proceed using `par(bty="o")`:

```
par(tck=0.02,las=1,cex.axis=1.3,cex.lab=1.2,mgp=c(3,0.5,0),bty
="o")

x<-c(1,3,4,6,8,9,12)
y<-c(5,8,6,10,9,13,12)
plot(x,y,type="n",axes=F,xlab="",ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-Axis")
box()
```



You could then add a regression line to the graph by typing
`abline(lm(y~x))`



There is a lot more to play around; here are some of the most useful graphics parameters from the `par()` command:

Specification of graphics parameters using `par()`:

- **adj**

The value of `adj` determines the way in which text strings are justified. A value of 0 produces **left-justified** text, 0.5 **centered** text and 1 **right-justified** text. (Any value in `[0, 1]` is allowed, and on most devices values outside that interval will also work.) Note that the `adj` argument of `text` also allows `adj = c(x, y)` for different adjustment in x- and y- direction.

- **ann**

If set to `FALSE`, high-level plotting functions calling `plot.default` do not annotate the plots they produce with axis titles and overall titles. The default is to do annotation.

- **ask**

logical. If `TRUE` (and the R session is interactive) the user is asked for input, before a new figure is drawn. As this applies to the device, it also affects output by packages `grid` and `lattice`. It can be set even on non-screen devices.

- **bg**

The color to be used for the background of plots. When called from `par()` it also sets `new=FALSE`. A description of how colors are specified is given below. Note that some graphics functions such as `plot.default` and `points` have an argument of this name with a different meaning.

- **bty**

A character string which determined the type of box which is drawn about plots. If `bty` is one of "o", "l", "7", "c", "u", or "]" the resulting box resembles the corresponding upper case letter. A value of "n" suppresses the box.

- **cex**

A numerical value giving the amount by which plotting text and symbols should be scaled relative to the default. Note that some graphics functions such as `plot.default` have an argument of this name which multiplies this graphical parameter.

- **cex.axis**

The magnification to be used for axis annotation relative to the current setting of `cex`. (Some functions such as `points` accept a vector of values which are recycled. Other uses will take just the first value if a vector of length greater than one is supplied.)

- **cex.lab**

The magnification to be used for x and y labels relative to the current setting of `cex`.

- **cex.main**

The magnification to be used for main titles relative to the current setting of `cex`.

- **cex.sub**

The magnification to be used for sub-titles relative to the current setting of `cex`.

- **`col`**

A specification for the default plotting color. A description of how colors are specified is given below. (Some functions such as `lines` accept a vector of values which are recycled. Other uses will take just the first value if a vector of length greater than one is supplied.)

- **`col.axis`**

The color to be used for axis annotation.

- **`col.lab`**

The color to be used for x and y labels.

- **`col.main`**

The color to be used for plot main titles.

- **`col.sub`**

The color to be used for plot sub-titles.

- **`crt`**

A numerical value specifying (in degrees) how single characters should be rotated. It is unwise to expect values other than multiples of 90 to work. Compare with `srt` which does string rotation.

- **`family`**

The name of a font family for drawing text. The maximum allowed length is 200 bytes. This name gets mapped by each graphics device to device-specific font descriptions. The default value is "" which means that the default device fonts will be used. Standard values are "serif", "sans", "mono", and "symbol" and the Hershey font families are also available. (Different devices may define others, and some devices will ignore this setting completely.) This can be specified inline for text.

- **`fg`**

The color to be used for the foreground of plots. This is the default color used for things like axes and boxes around plots. When called from `par()` this also sets parameter `col` to the same value.

A description of how colors are specified is given below.

- **`fig`**

A numerical vector of the form `c(x1, x2, y1, y2)` which gives the (NDC) coordinates of the figure region in the display region of the device. If you set this, unlike `S`, you start a new plot, so to add to an existing plot use `new=TRUE` as well.

- **`fin`**

The figure region dimensions, `(width,height)`, in inches. If you set this, unlike `S`, you start a new plot.

- **`font`**

An integer which specifies which font to use for text. If possible, device drivers arrange so that 1 corresponds to plain text, 2 to bold face, 3 to italic and 4 to bold italic. Also, font 5 is expected to be the symbol font, in Adobe symbol encoding. On some devices font families can be selected by family to choose different sets of 5 fonts.

- **font.axis**

The font to be used for axis annotation.

- **font.lab**

The font to be used for x and y labels.

- **font.main**

The font to be used for plot main titles.

- **font.sub**

The font to be used for plot sub-titles.

- **gamma**

the gamma correction, see argument gamma to hsv. This is only accepted if the current device has support for changing the gamma correction: currently only windows and quartz do. (X11 has support for setting the gamma correction when opening the device, but not for changing it.)

- **lab**

A numerical vector of the form `c(x, y, len)` which modifies the default way that axes are annotated. The values of `x` and `y` give the (approximate) number of tickmarks on the `x` and `y` axes and `len` specifies the label length. The default is `c(5, 5, 7)`. Note that this only affects the way the parameters `xaxp` and `yaxp` are set when the user coordinate system is set up, and is not consulted when axes are drawn. `len` is unimplemented in R.

- **las**

numeric in `{0,1,2,3}`; the style of axis labels.

0: always parallel to the axis [default],

1: always horizontal,

2: always perpendicular to the axis,

3: always vertical.

Note that other string/character rotation (via argument `srt` to `par`) does not affect the axis labels.

- **lend**

The line end style. This can be specified as an integer or string:

0 and "round" mean rounded line caps [default];

1 and "butt" mean butt line caps;

2 and "square" mean square line caps.

- **lheight**

The line height multiplier. The height of a line of text (used to vertically space multi-line text) is found by multiplying the current font size both by the current character expansion and by the line height multiplier. Default value is 1.

- **ljoin**

The line join style. This can be specified as an integer or string:

0 and "round" mean rounded line joins [default];
1 and "mitre" mean mitred line joins;
2 and "bevel" mean bevelled line joins.

- **lmitre**

The line mitre limit. This controls when mitred line joins are automatically converted into bevelled line joins. The value must be larger than 1 and the default is 10. Not all devices will honour this setting.

- **lty**

The line type. Line types can either be specified as an integer (0=blank, 1=solid, 2=dashed, 3=dotted, 4=dotdash, 5=longdash, 6=twodash) or as one of the character strings "blank", "solid", "dashed", "dotted", "dotdash", "longdash", or "twodash", where "blank" uses 'invisible lines' (i.e., does not draw them).

Alternatively, a string of up to 8 characters (from c(1:9, "A":"F")) may be given, giving the length of line segments which are alternatively drawn and skipped. See section 'Line Type Specification' below.

Some functions such as lines accept a vector of values which are recycled. Other uses will take just the first value if a vector of length greater than one is supplied.

- **lwd**

The line width, a positive number, defaulting to 1. The interpretation is device-specific, and some devices do not implement line widths less than one. (See the help on the device for details of the interpretation.)

Some functions such as lines accept a vector of values which are recycled. Other uses will take just the first value if a vector of length greater than one is supplied.

- **mai**

A numerical vector of the form c(bottom, left, top, right) which gives the margin size specified in inches.

- **mar**

A numerical vector of the form c(bottom, left, top, right) which gives the number of lines of margin to be specified on the four sides of the plot. The default is c(5, 4, 4, 2) + 0.1.

- **mex**

mex is a character size expansion factor which is used to describe coordinates in the margins of plots. Note that this does not change the font size, rather specifies the size of font used to convert between mar and mai, and between oma and omi.

- **mfc**ol, **mfr**ow

A vector of the form `c(nr, nc)`. Subsequent figures will be drawn in an `nr`-by-`nc` array on the device by columns (`mfc`ol), or rows (`mfr`ow), respectively.

In a layout with exactly two rows and columns the base value of `"cex"` is reduced by a factor of 0.83: if there are three or more of either rows or columns, the reduction factor is 0.66.

Consider the alternatives, `layout` and `split.screen`.

- **mfg**

A numerical vector of the form `c(i, j)` where `i` and `j` indicate which figure in an array of figures is to be drawn next (if setting) or is being drawn (if enquiring). The array must already have been set by `mfc`ol or `mfr`ow.

For compatibility with `S`, the form `c(i, j, nr, nc)` is also accepted, when `nr` and `nc` should be the current number of rows and number of columns. Mismatches will be ignored, with a warning.

- **m**gp

The margin line (in `mex` units) for the axis title, axis labels and axis line. The default is `c(3, 1, 0)`.

- **m**kh

The height in inches of symbols to be drawn when the value of `pch` is an integer. Completely ignored currently.

- **n**ew

logical, defaulting to `FALSE`. If set to `TRUE`, the next high-level plotting command (actually `plot.new`) should not clean the frame before drawing "as if it was on a new device".

- **o**ma

A vector of the form `c(bottom, left, top, right)` giving the size of the outer margins in lines of text.

- **o**md

A vector of the form `c(x1, x2, y1, y2)` giving the outer margin region in NDC (= normalized device coordinates), i.e., as fraction (in `[0,1]`) of the device region.

- **o**mi

A vector of the form `c(bottom, left, top, right)` giving the size of the outer margins in inches.

- **p**ch

Either an integer specifying a symbol or a single character to be used as the default in plotting points. See `points` for possible values and their interpretation.

- **p**in

The current plot dimensions, `(width,height)`, in inches.

- **p**lt

A vector of the form `c(x1, x2, y1, y2)` giving the coordinates of the plot region as fractions of the current figure region.

- **ps**

integer; the pointsize of text (but not symbols). Unlike the pointsize argument of most devices, this does not change the relationship between `mar` and `mai` (nor `oma` and `omi`), but it does scale `cin` and `csi`.

What is meant by 'pointsize' is device-specific, but most devices mean a multiple of lbp, that is 1/72 of an inch.

- **pty**

A character specifying the type of plot region to be used; "s" generates a square plotting region and "m" generates the maximal plotting region.

- **srt**

The string rotation in degrees. See the comment about `crt`.

- **tck**

The length of tick marks as a fraction of the smaller of the width or height of the plotting region. If `tck` \geq 0.5 it is interpreted as a fraction of the relevant side, so if `tck` = 1 grid lines are drawn. The default setting (`tck` = NA) is to use `tcl` = -0.5 (see below).

- **tcl**

The length of tick marks as a fraction of the height of a line of text. The default value is -0.5; setting `tcl` = NA sets `tck` = -0.01 which is S' default.

- **usr**

A vector of the form `c(x1, x2, y1, y2)` giving the extremes of the user coordinates of the plotting region. When a logarithmic scale is in use (i.e., `par("xlog")` is true, see below), then the x-limits will be $10^{\text{par("usr")[1:2]}}$. Similarly for the y-axis.

- **xaxp**

A vector of the form `c(x1, x2, n)` giving the coordinates of the extreme tick marks and the number of intervals between tick-marks when `par("xlog")` is false. Otherwise, when log coordinates are active, the three values have a different meaning: For a small range, `n` is negative, and the ticks are as in the linear case, otherwise, `n` is in 1:3, specifying a case number, and `x1` and `x2` are the lowest and highest power of 10 inside the user coordinates, $10^{\text{par("usr")[1:2]}}$. (The "usr" coordinates are log₁₀-transformed here!)

- `n=1`
- will produce tick marks at 10^j for integer `j`,
- `n=2`
- gives marks `k 10^j` with `k` in `{1, 5}`,
- `n=3`

- gives marks $k \cdot 10^j$ with k in $\{1, 2, 5\}$.
- See `axTicks()` for a pure R implementation of this.

This parameter is reset when a user coordinate system is set up, for example by starting a new page or by calling `plot.window` or setting `par("usr")`: n is taken from `par("lab")`. It affects the default behaviour of subsequent calls to `axis` for sides 1 or 3.

- **xaxs**

The style of axis interval calculation to be used for the x-axis. Possible values are "r", "i", "e", "s", "d". The styles are generally controlled by the range of data or `xlim`, if given. Style "r" (regular) first extends the data range by 4 percent and then finds an axis with pretty labels that fits within the range. Style "i" (internal) just finds an axis with pretty labels that fits within the original data range. Style "s" (standard) finds an axis with pretty labels within which the original data range fits. Style "e" (extended) is like style "s", except that it also ensures that there is room for plotting symbols within the bounding box. Style "d" (direct) specifies that the current axis should be used on subsequent plots. (Only "r" and "i" styles are currently implemented)

- **xaxt**

A character which specifies the x axis type. Specifying "n" suppresses plotting of the axis. The standard value is "s": for compatibility with S values "l" and "t" are accepted but are equivalent to "s": any value other than "n" implies plotting.

- **xlog**

A logical value (see `log` in `plot.default`). If TRUE, a logarithmic scale is in use (e.g., after `plot(*, log = "x")`). For a new device, it defaults to FALSE, i.e., linear scale.

- **xpd**

A logical value or NA. If FALSE, all plotting is clipped to the plot region, if TRUE, all plotting is clipped to the figure region, and if NA, all plotting is clipped to the device region.

- **yaxp**

A vector of the form `c(y1, y2, n)` giving the coordinates of the extreme tick marks and the number of intervals between tick-marks unless for log coordinates, see `xaxp` above.

- **yaxs**

The style of axis interval calculation to be used for the y-axis. See `xaxs` above.

- **yaxt**

A character which specifies the y axis type. Specifying "n" suppresses plotting.

- **ylog**

A logical value; see `xlog` above.

- **Color Specification**

Colors can be specified in several different ways. The simplest way is with a character string giving the color name (e.g., "red"). A list of the possible colors can be obtained with the function `colors`. Alternatively, colors can be specified directly in terms of their RGB components with a string of the form "#RRGGBB" where each of the pairs RR, GG, BB consist of two hexadecimal digits giving a value in the range 00 to FF. Colors can also be specified by giving an index into a small table of colors, the palette. This provides compatibility with S. Index 0 corresponds to the background color.

Additionally, "transparent" or (integer) NA is transparent, useful for filled areas (such as the background!), and just invisible for things like lines or text. Semi-transparent colors are available for use on devices that support them.

The functions `rgb`, `hsv`, `hcl`, `gray` and `rainbow` provide additional ways of generating colors.

- **Line Type Specification**

Line types can either be specified by giving an index into a small built-in table of line types (1 = solid, 2 = dashed, etc, see `lty` above) or directly as the lengths of on/off stretches of line. This is done with a string of an even number (up to eight) of characters, namely non-zero (hexadecimal) digits which give the lengths in consecutive positions in the string. For example, the string "33" specifies three units on followed by three off and "3313" specifies three units on followed by three off followed by one on and finally three off. The 'units' here are (on most devices) proportional to `lwd`, and with `lwd = 1` are in pixels or points or 1/96 inch.

The five standard dash-dot line types (`lty = 2:6`) correspond to `c("44", "13", "1343", "73", "2262")`.

Note that NA is not a valid value for `lty`.

Prior to R 2.4.0, zero hex digits were accepted and handled in a device-dependent way, e.g. `X11()` mapped them to one, `pdf()` regarded zero as terminating the string and `windows()` gave a zero-length stretch.

Colors

Here is a list of colors that can be used with R; the list can also be reproduced using `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"	"chocolatel1"	"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"

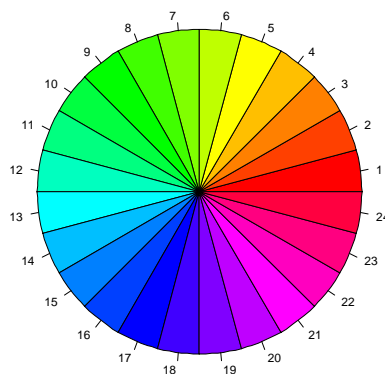

```

[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"

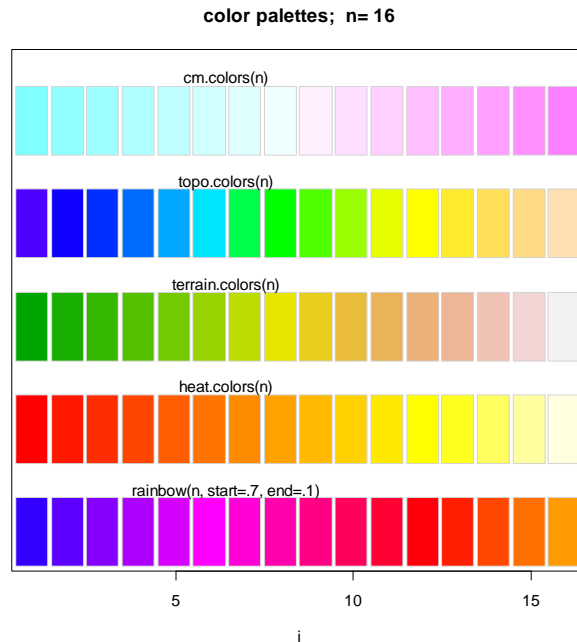
```

A nice example of color usage in R is the `rainbow()` color palette used for plotting a pie chart:

```
pie(rep(1, 24), col = rainbow(24), radius = 0.9)
```



In addition to the rainbow colors, there are several other built-in color palettes in R:



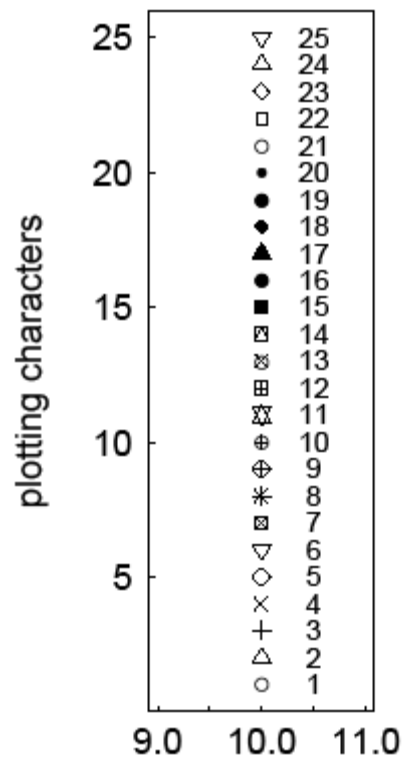
Just for background information, these palettes have been called using

```
demo.pal <-
  function(n, border = if (n<32) "light gray" else NA,
           main = paste("color palettes; n=",n),
           ch.col = c("rainbow(n, start=.7, end=.1)",
                     "heat.colors(n)", "terrain.colors(n)", "topo.colors(n)",
                     "cm.colors(n)"))
  {
    nt <- length(ch.col)
    i <- 1:n; j <- n / nt; d <- j/6; dy <- 2*d
    plot(i,i+d, type="n", yaxt="n", ylab="", main=main)
    for (k in 1:nt) {
      rect(i-.5, (k-1)*j+ dy, i+.4, k*j,
           col = eval(parse(text=ch.col[k])), border =
border)
      text(2*j, k * j +dy/4, ch.col[k])
    }
  }
n <- if(.Device == "postscript") 64 else 16

demo.pal(n)
```

Plotting characters

Here is an overview of R's default plotting characters.



This figure has been created using the `pdf()` device like this:

```
pdf(file="C:\\characters.pdf",pointsize=14,width=3,height=7)
```

```
par(mgp=c(3,0.5,0),tck=0.02,cex.axis=1.3,cex.lab=1.3,lwd=1,las=1)
```

```
plot(rep(10,25),1:25,pch=1:25,xlim=c(9,11),xlab="",ylab="plotting characters")
```

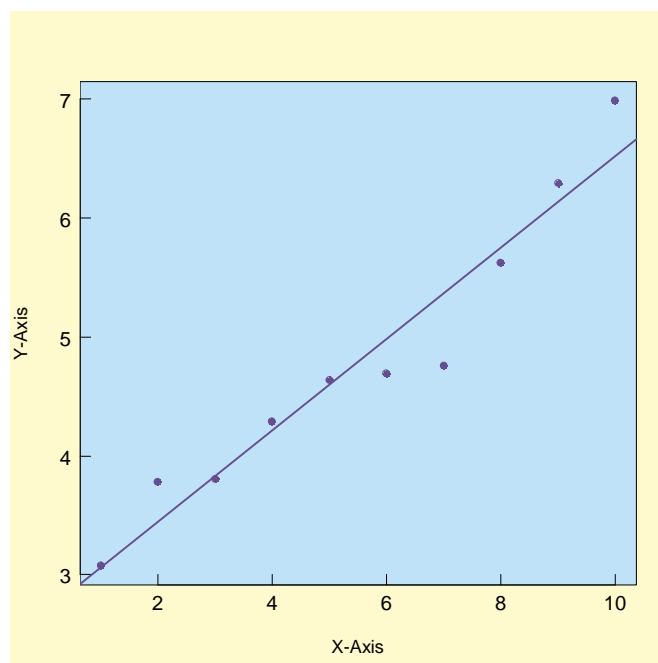
```
text(rep(10.5,25),1:25,1:25)
```

```
dev.off()
```

Let's use some of the above-mentioned commands to make our original plot look fancier:

```
par(tck=0.02,las=1,cex.axis=1.3,cex.lab=1.2,mgp=c(3,0.5,0),
    bty="o",bg="lemonchiffon")
```

```
plot(x,y,type="n",axes=F,xlab="",ylab="")
u <- par("usr")
rect(u[1],u[3],u[2],u[4],col=rgb(191,226,248,max=255))
points(x,y,pch=16,cex=1.3,col=rgb(102,78,145,max=255))
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-Axis")
abline(lm(y~x),lwd=2,col=rgb(102,78,145,max=255))
```

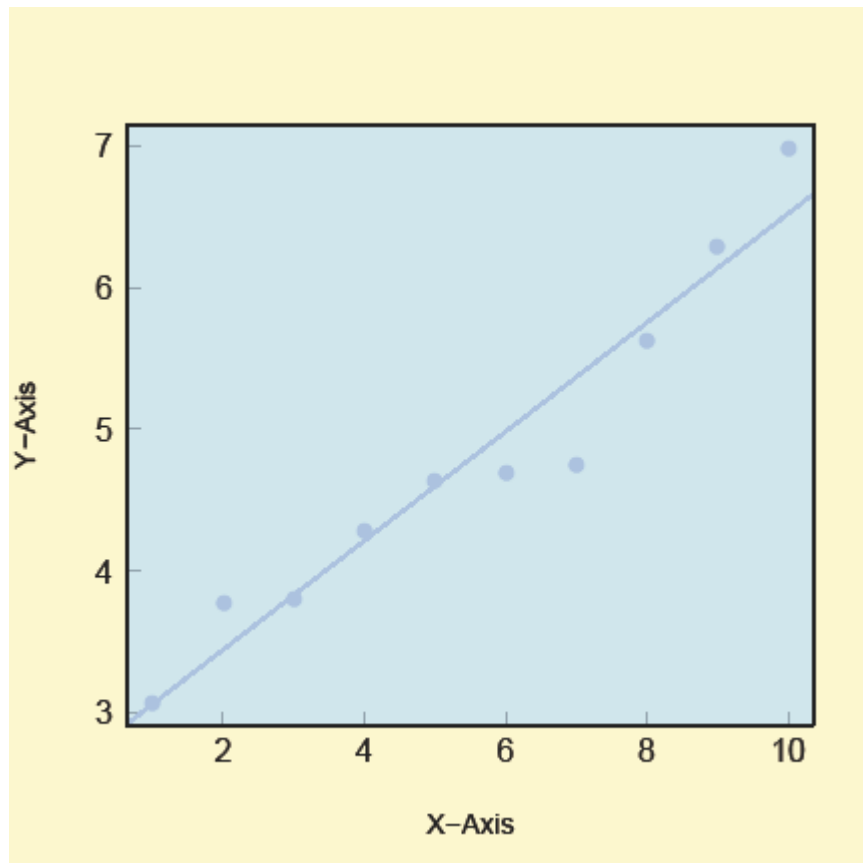


Here, colors are specified using `rgb()` values directly. **Transparency** of colored components of a plot can also be adjusted, but this is restricted to the `pdf()` device:

```
pdf("G:\\transparency.pdf",pointsize=12)
par(tck=0.02,las=1,cex.axis=1.3,cex.lab=1.2,mgp=c(3,0.5,0),
    bty="o",bg="lemonchiffon")
```

```
plot(x,y,type="n",axes=F,xlab="",ylab="")
u <- par("usr")
points(x,y,pch=16,cex=1.3,col=rgb(102,78,145,max=255))
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-Axis")
abline(lm(y~x),lwd=2,col=rgb(102,78,145,max=255))
```

```
rect(u[1],u[3],u[2],u[4],
col=rgb(191/255,226/255,248/255,alpha=0.8),
border=F)
box(lwd=2)
dev.off()
```



Now what most of you will basically struggle with when creating R graphics are some of the basic questions such as:

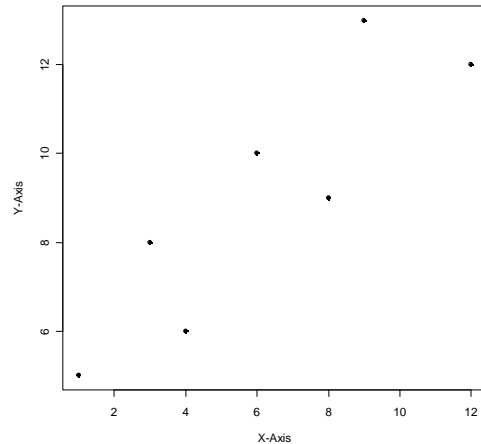
- How do I change the font size
- How do I make lines thicker or thinner, or change the line type
- How can I change plotting characters
- How do I change the distance between axis notations, axis title(s) and plot margins
- How do I control the size of the plot area?

All these questions are not difficult to answer, once you have carefully read through the `par()` help page.

Here are some solutions, again using our standard plot.

Here is the standard plot once more:

```
x<-c(1,3,4,6,8,9,12)
y<-c(5,8,6,10,9,13,12)
plot(x,y,type="n",axes=F,xlab
="",ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-
Axis")
box()
```



It has the basic look of any newly created R graphics object, so it is an ideal example of how to proceed when editing any R graph.

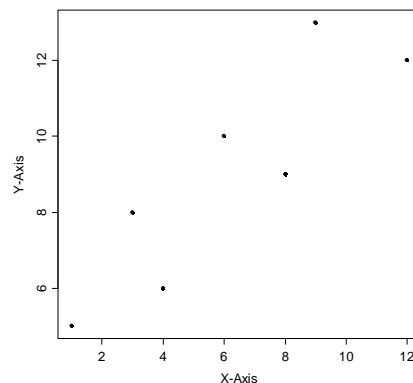
How do I change the font size?

Using `par()`, you can do this using the commands

```
cex.axis()
cex.lab() and
cex.sub()
```

"**cex**" means "character expansion"; by default, the value is 1, so if we want to increase our font sizes by 1.3, we write `cex(1.3)`, or, using our default graph again:

```
par(cex.axis=1.3,
cex.lab=1.3)
plot(x,y,type="n",axes=F,xlab
="",ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-
Axis")
box()
```

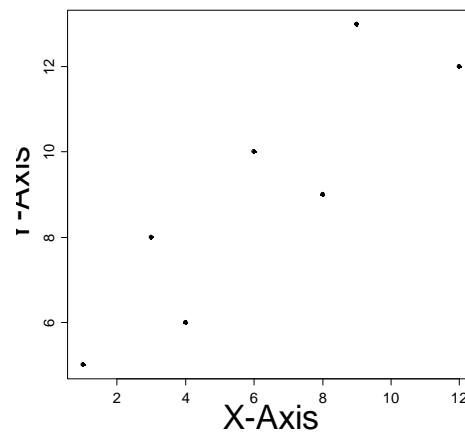


Let's see what happens if we exaggerate too much:

```

par(
  cex.axis=1.3,
  cex.lab=3)
plot(x,y,type="n",axes=F,xlab
     ="",ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-
Axis")
box()

```



Now the y axis label has vanished, and we need to account for this by changing the size of the plotting device area. You would usually adjust this using the `mar()` command:

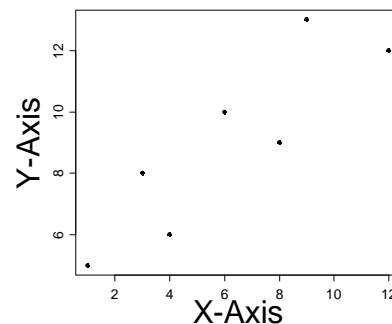
The standard values for `mar()` are `c(5, 4, 4, 2) + 0.1`, in the direction "bottom", "left", "top", "right".

Let's see what happens if we change these values:

```

par(mar=c(10,6,6,5),cex.axis=
  1.3,cex.lab=3)
plot(x,y,type="n",axes=F,xlab
     ="",ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-
Axis")
box()

```

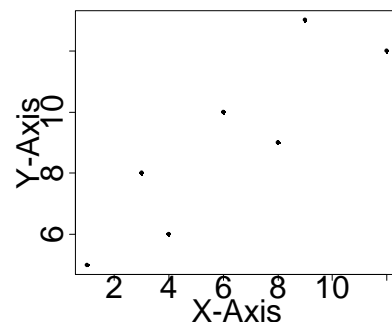


This looks alright now; but what if we changed the `cex.axis` to 3, also?

```

par(mar=c(10,6,6,5),cex.axis=
  3,cex.lab=3)
plot(x,y,type="n",axes=F,xlab
     ="",ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-
Axis")
box()

```

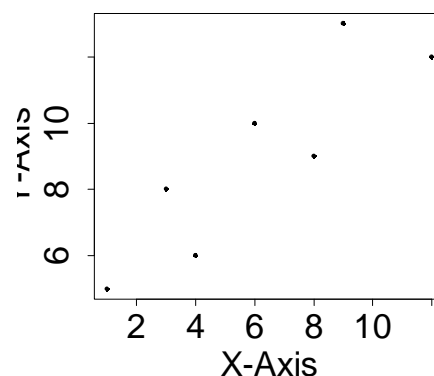


Well, now we need another command to change the distances between axis labels and axes. This is done using the `mgp()` command, which says: "How many lines of standard-sized text shall fit outside the plotting area?" Here, it is

- (1) How many lines underneath the axis tick labels?
- (2) How many lines between axis and tick label and
- (3) How many lines shall the axis line itself be away from the plot?

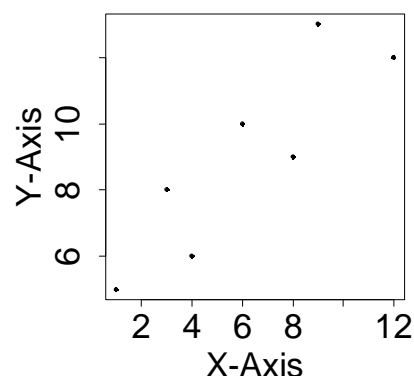
let us try the following:

```
par(mar=c(10,6,6,5),
    mgp=c(5,2,0),
    cex.axis=3,
    cex.lab=3)
plot(x,y,type="n",axes=F,xlab=""
     ,ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-
Axis")
box()
```



Now, again, we need to adjust the plotting area:

```
par(mar=c(10,10,6,5),
    mgp=c(5,2,0),
    cex.axis=3,
    cex.lab=3)
plot(x,y,type="n",axes=F,xlab=""
     ,ylab="")
points(x,y,pch=16)
axis(1)
axis(2)
title(xlab="X-Axis",ylab="Y-
Axis")
box()
```

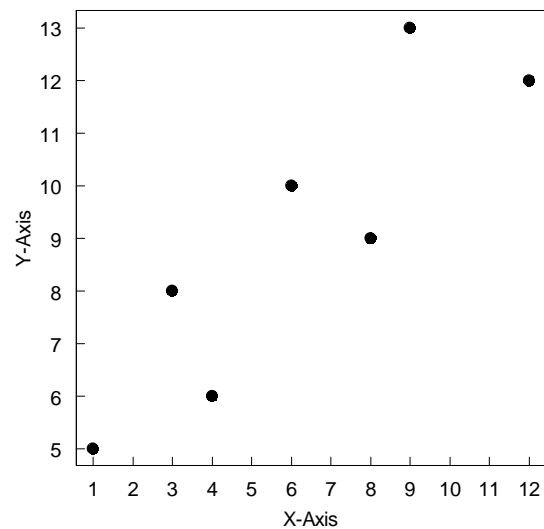


If you are not satisfied with the way the axis ticks are written, you can also change this; let us further change several other things such as the rotation of the tickmark labels using `las()`, the orientation of the tickmarks using `tck()`, and the sizes of the plotting characters using `cex()` inside the `points()` command:


```

par(mar=c(6,6,2,1),
    mgp=c(3,1,0),
    cex.axis=1.5,
    cex.lab=1.5,
    las=1, tck=0.02)
plot(x,y,type="n",axes=F,xlab=""
     ,ylab="")
points(x,y,pch=16,cex=2)
axis(1,at=1:12)
axis(2,at=1:14)
title(xlab="X-Axis",ylab="Y-
Axis")
box()

```

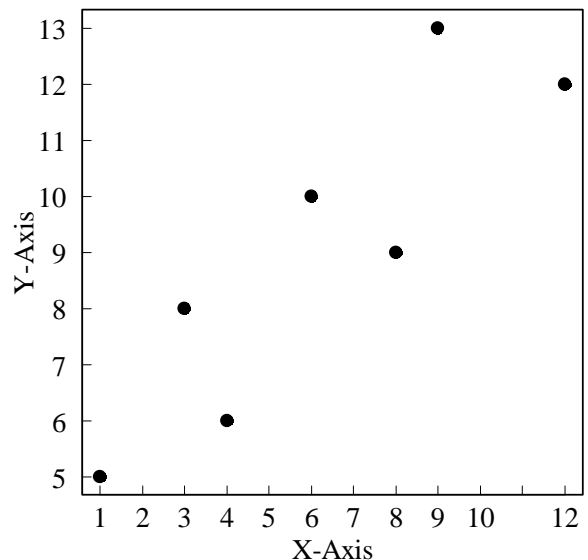


Now, finally, let us change the fonts used for overall labelling (`family()` and `font()`), the overall font sizes onscreen more using `cex.axis()` and `cex.lab()`, and the line widths using `lwd()`:

```

par(mar=c(6,6,2,1),
    mgp=c(3,1,0),
    cex.axis=2,
    cex.lab=2,
    las=1,
    font=2,
    family="serif",
    lwd=2,tck=0.02
)
plot(x,y,type="n",axes=F,xlab=""
     ,ylab="")
points(x,y,pch=16,cex=2)
axis(1,at=1:12)
axis(2,at=1:14)
title(xlab="X-Axis",ylab="Y-
Axis")
box()

```



Another way of dealing with all these above-mentioned things quite conveniently is to use the `pdf` device and directly use the command:

```
pdf(file="C:\\sample.pdf",pointsize=12)
followed by additional par() commands and in the end a
dev.off() command like written below:
```

```

par(mar=c(6,6,2,1),mgp=c(3,1,0),las=1,lwd=2)
plot(x,y,type="n",axes=F,xlab="" ,ylab="")
points(x,y,pch=16,cex=2)

```

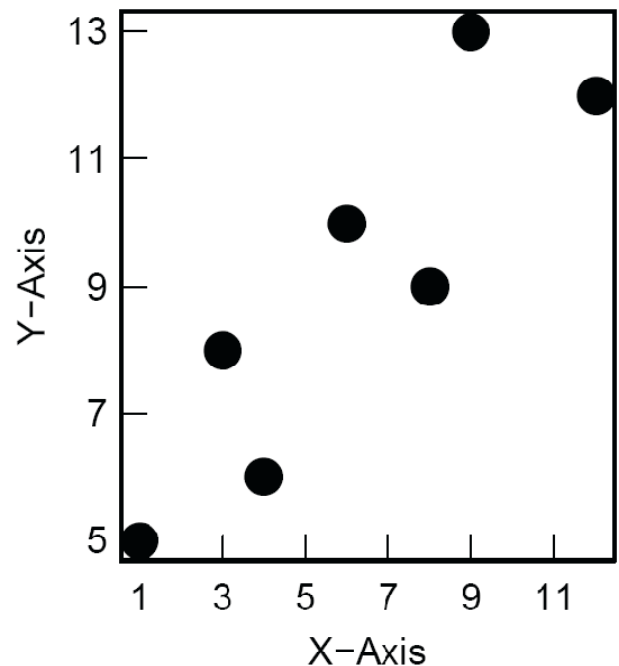
```
axis(1,at=1:12)
axis(2,at=1:14)
title(xlab="X-Axis",ylab="Y-Axis")
box()
dev.off()
```

where `pointsize()` gives the size of the font to be used.

Using the pdf device also gives you the option to specify the font family to be used, and the size you would like to have for your document; usually, it is also useful to state explicitly what character encoding you would wish to use. To create our following last figure, several other options have been adjusted, because the sizes of the figure were explicitly given to be 4 inches wide and 5 inches in height:

```
pdf("G:\\sample.pdf",width=4,
height=5,
pagecentre=T,pointsize=14,
fonts=c("Helvetica"),encoding
="WinAnsi")

par(las=1,lwd=2,tck=0.05,mgp=
c(1.5,0.3,0))
plot(x,y,type="n",axes=F,xlab
="",ylab="")
points(x,y,pch=16,cex=2)
axis(1,at=seq(1,12,2))
axis(2,at=seq(1,14,2))
title(xlab="X-Axis",ylab="Y-
Axis")
box()
dev.off()
```



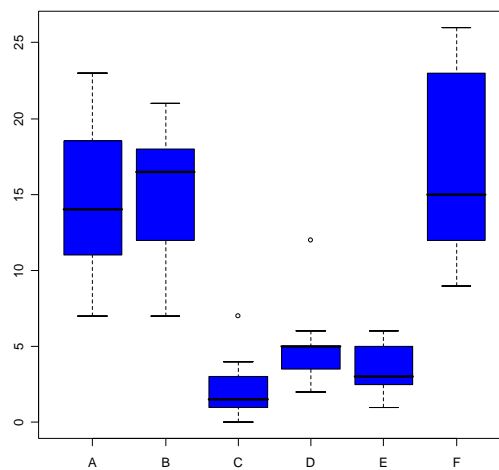
That's all for now. Let's now switch to other plot types.

Boxplots can be drawn using several options. One of the most interesting ones is `notch=T`, which draws notches that show confidence intervals for the medians:

```
par(tck=0.02, las=1, cex.axis=1.3, cex.lab=1.2, mgp=c(3, 0.5, 0),  
    bty="o", bg="lemonchiffon")
```

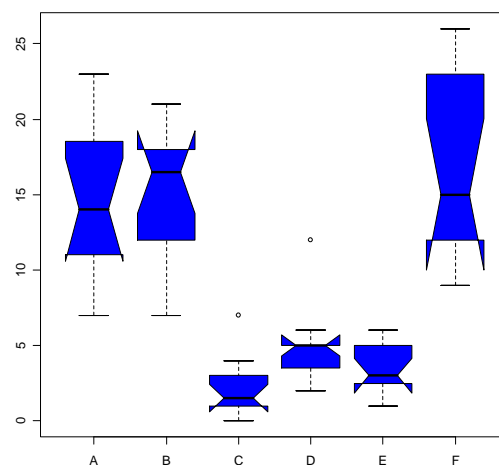
a) standard boxplot:

```
boxplot(count ~ spray, data = InsectSprays, notch = F,  
        col = "blue")
```



b) with notches:

```
boxplot(count ~ spray, data = InsectSprays, notch = TRUE,  
        col = "blue")
```



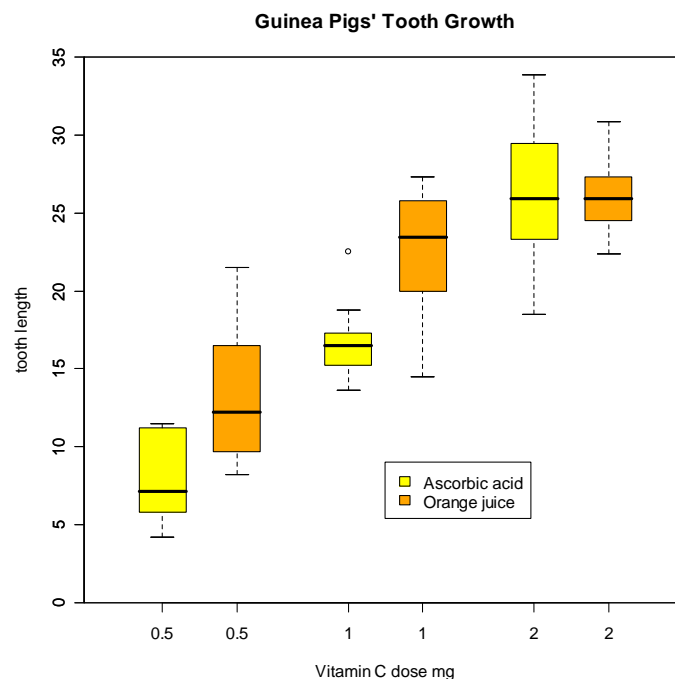
Another useful feature of boxplots is to have several boxes grouped at the x axis; this is something that most other stats packages will fail to produce:

```
boxplot(len ~ dose, data = ToothGrowth,
        boxwex = 0.25, at = 1:3 - 0.2,
        subset = supp == "VC", col = "yellow",
        main = "Guinea Pigs' Tooth Growth",
        xlab = "Vitamin C dose mg",
        ylab = "tooth length", ylim = c(0, 35), yaxs = "i")
boxplot(len ~ dose, data = ToothGrowth, add = TRUE,
        boxwex = 0.25, at = 1:3 + 0.2,
        subset = supp == "OJ", col = "orange")
legend(2, 9, c("Ascorbic acid", "Orange juice"),
       fill = c("yellow", "orange"))
```

The trick here is to use the

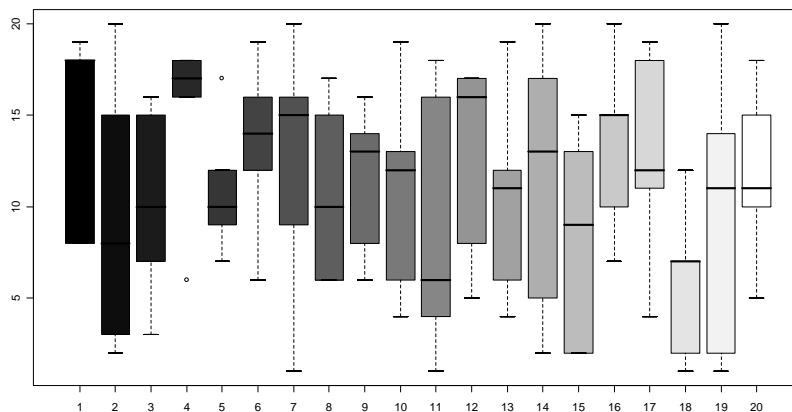
- "add" command, the
- "at" command and the
- "subset"

command when specifying boxplots.



Working with different shadings of grey can be shown using a quick example dataset:

```
boxplot(
  sample(1:20,100,replace=T)~
  sort(rep(1:20,5)),
  col=gray(seq(from=0,to=1,length=20)))
```



2. Trellis plots

Here are some introductory examples of what can be done using trellis plots:

```
require(lattice)

sundar.theme <- function() {
  par <- col.whitebg()
  par$strip.background$col <- rep("#000099", 7)
  par$add.text$col <- "#eeeeaa"
  par$add.text$font <- 2
  par$background$col <- "#ffffff"
  par$superpose.line$lty <- rep(1, 7)
  par$superpose.line$col[1:2] <- c("#880000",
"#008800")
  par$superpose.symbol$col[1:2] <- c("#880000",
"#008800")
  par
}

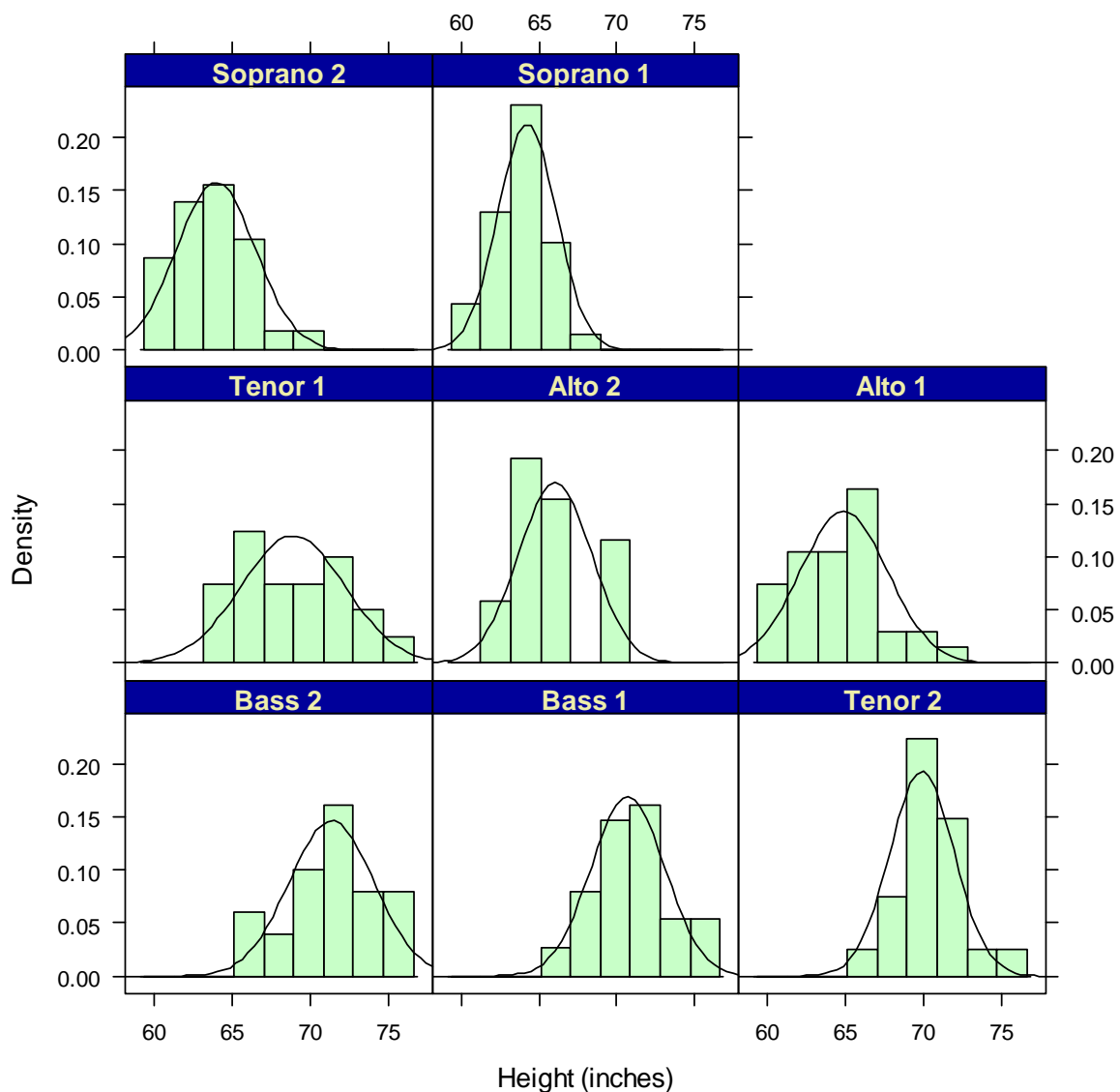
trellis.par.set(sundar.theme())

print( # necessary if the file is source()'d
```

```

histogram( ~ height | voice.part, data = singer,
           xlab = "Height (inches)", type = "density",
           panel = function(x, ...) {
             panel.histogram(x, ...)
             panel.mathdensity(dmath = dnorm, col =
"black",
                               args =
list(mean=mean(x),sd=sd(x)))
           } )
)

```



Conditioning plots (coplots):

```

library(MASS)

with(crabs,
     coplot(FL ~ RW | sp * sex,
           subscripts = T,

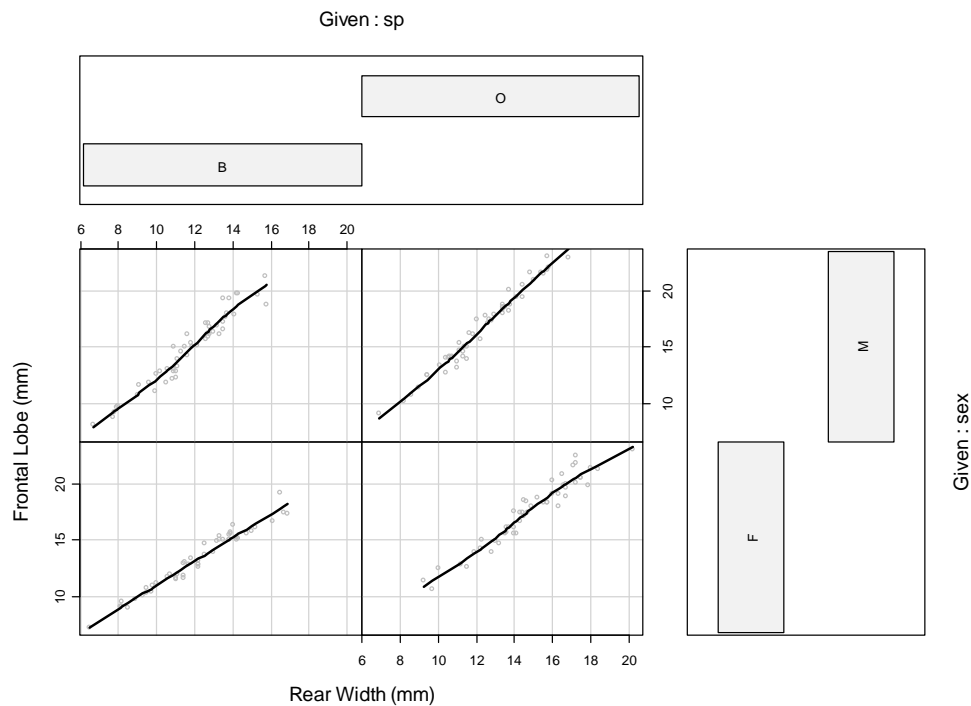
```

```

      xlab = "Rear Width (mm)",
      ylab = "Frontal Lobe (mm)",
      panel = function(x, y, subscripts, ...){
        points(x, y, col = "gray75")

        lines(smooth.spline(RW[subscripts], FL[subscripts]), lwd
              = 2,col = "black")
      }
    )
  )

```

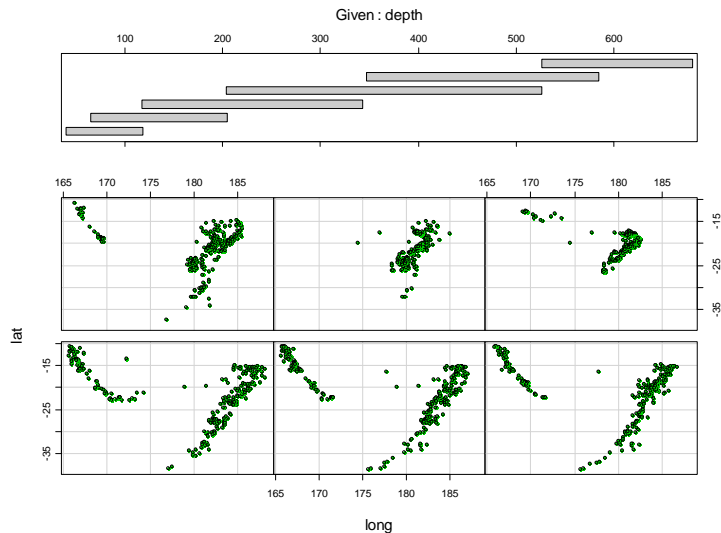


Coplot() can also be used with numerical conditioning variables:

```

coplot(lat ~ long | depth, data = quakes, pch = 21, bg =
"green3")

```



Here is another example showing the usage of `xyplot()`:

```
library(lattice)

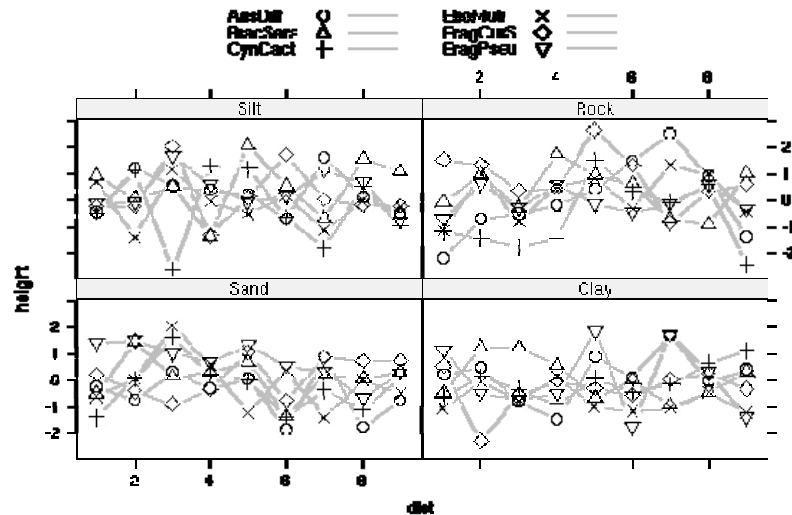
lattice.options(default.theme = canonical.theme(color =
FALSE))

tmp <- expand.grid(geology =
c("Sand","Clay","Silt","Rock"),species = c("ArisDiff",
"BracSera", "CynDact", "ElioMuti", "EragCurS",
"EragPseu"),dist = seq(1,9,1) )

tmp$height <- rnorm(216)

sp <- list(superpose.symbol = list(pch = 1:6, cex = 1.2),
superpose.line = list(col = "grey", lty = 1))

print(xyplot(height ~ dist | geology, data = tmp,
groups = species,
layout = c(2,2),
panel = function(x, y, type, ...) {
panel.superpose(x, y, type="l", ...)
lpoints(x, y, pch=16, col="white", cex=2)
panel.superpose(x, y, type="p",...)
},
par.settings = sp,
auto.key = list(columns = 2, lines = TRUE)))
```

3. Examples from special graphics packages

The examples below are from the "R Graph Gallery" maintained by Francois Romain at

<http://addictedtor.free.fr/graphiques/>

The Hmisc package

To create the following graph (showing variations of boxplots), several libraries will need to be downloaded and installed:

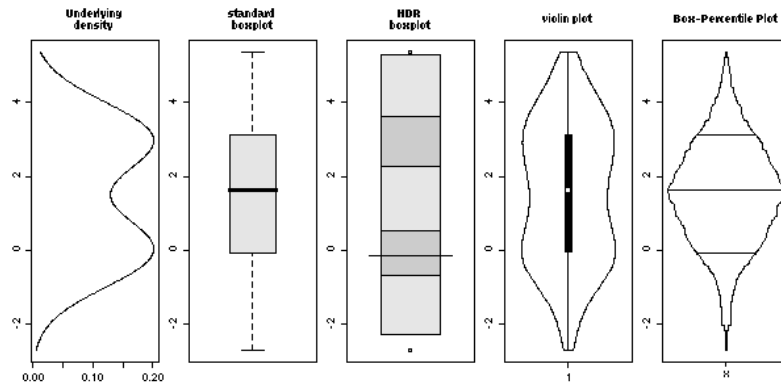
```
require(hdrcde)
require(vioplot)
require(Hmisc)

x <- c(rnorm(100,0,1), rnorm(100,3,1))
opar <- par(mfrow=c(1,5), mar=c(3,2,4,1))

xxx <- seq(min(x), max(x), length=500)
yyy <- dnorm(xxx)/2 + dnorm(xxx, mean=3)/2

plot(yyy, xxx, type="l", main="Underlying\ndensity")

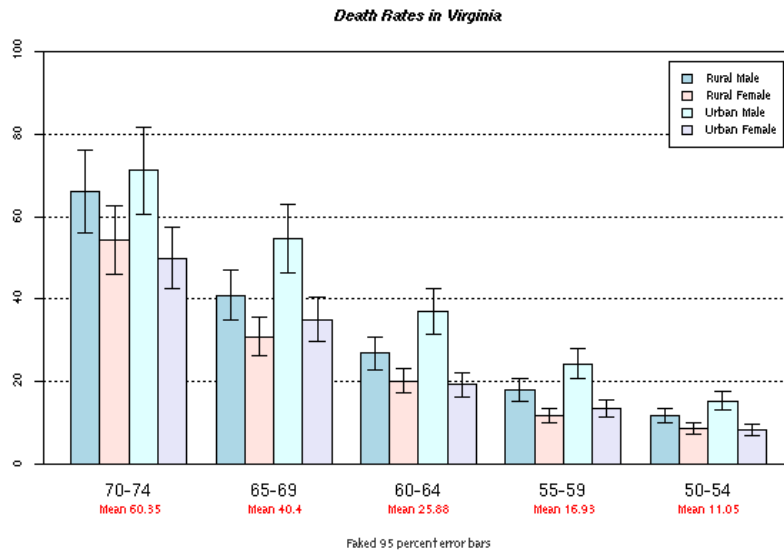
boxplot(x, col="gray90", main="standard\nboxplot")
hdr.boxplot(x, main="HDR\nboxplot")
vioplot(x)
title("violin plot")
bplot(x)
par(opar)
```



gplots package

This package is a fusion of conventional and lattice graphics. It contains very useful features, but some are still in development.

```
library(gplots)
# Example with confidence intervals and grid
hh <- t(VADeaths)[, 5:1]
mybarcol <- "gray20"
ci.l <- hh * 0.85
ci.u <- hh * 1.15
mp <- barplot2(hh, beside = TRUE,
               col = c("lightblue", "mistyrose",
                       "lightcyan", "lavender"),
               legend = colnames(VADeaths), ylim = c(0, 100),
               main = "Death Rates in Virginia", font.main = 4,
               sub = "Faked 95 percent error bars", col.sub =
mybarcol,
               cex.names = 1.5, plot.ci = TRUE, ci.l = ci.l, ci.u =
ci.u,
               plot.grid = TRUE)
mtext(side = 1, at = colMeans(mp), line = 2,
      text = paste("Mean", formatC(colMeans(hh))), col =
"red")
box()
```



Similarly, plotting a two-dimensional histogram is also possible using the `gplots` library:

```
require(gplots)
```

```
# example data, bivariate normal, no correlation
```

```
x <- rnorm(2000, sd=4)
```

```
y <- rnorm(2000, sd=1)
```

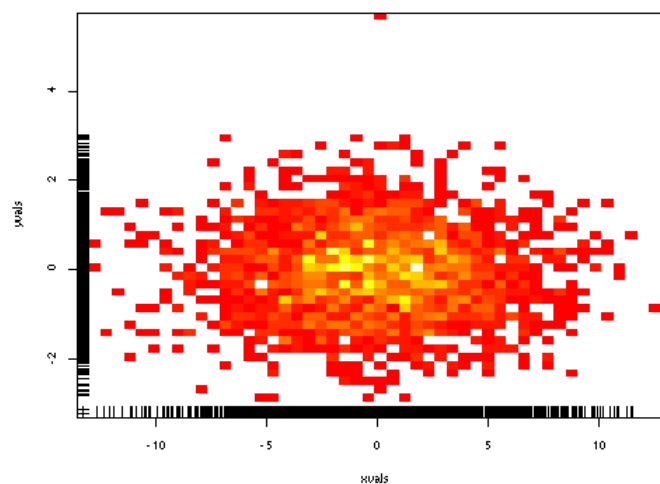
```
# separate scales for each axis, this looks circular
```

```
hist2d(x,y, nbins=50, col = c("white",heat.colors(16)))
```

```
rug(x,side=1)
```

```
rug(y,side=2)
```

```
box()
```

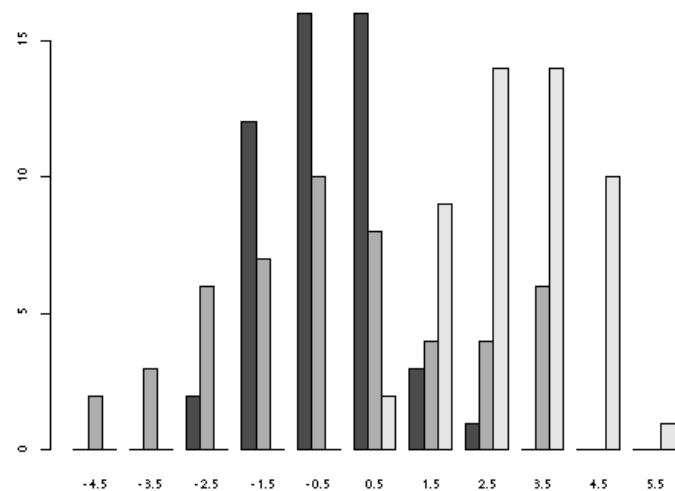


plotrix package

The following short command allows the drawing of a double histogram side-by-side; the only thing required is to have the different datapoints concatenated in a list with as many sub-lists as there shall be histograms:

```
require(plotrix)
```

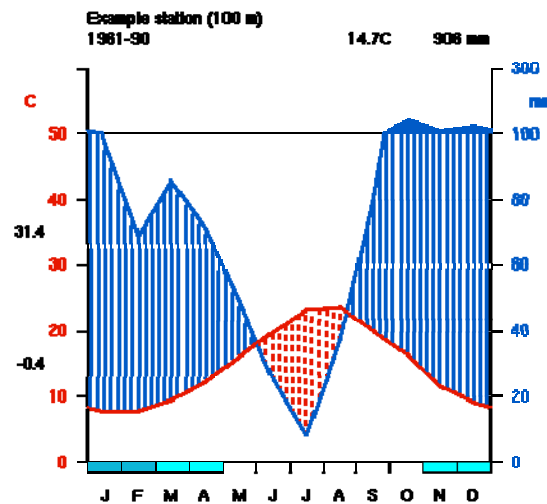
```
l <- list(rnorm(50),rnorm(50,sd=2),rnorm(50,mean=3))
multhist(l)
```

**Climatol package**

This package allows the drawing of classical climate diagrams.

```
require(climatol)
```

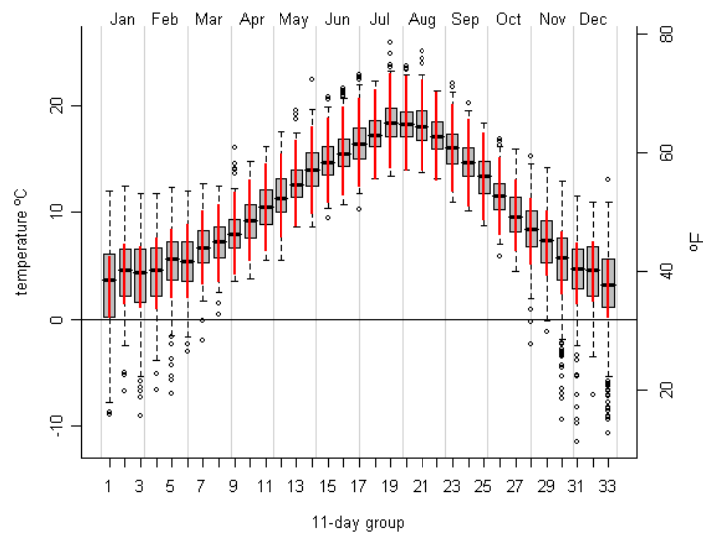
```
data(cli.dat)
diagwl(cli.dat,est="Example station",alt=100,per="1961-90",mlab="en")
```



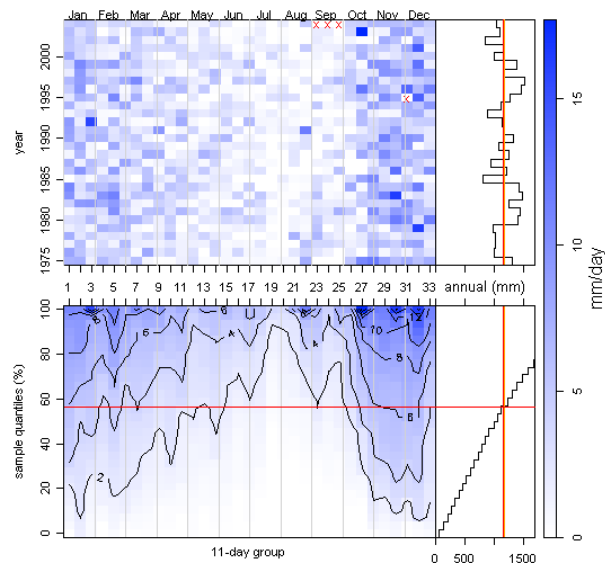
Seas package

This package provides plotting options for seasonal data.

```
require(seas)
data(mscdata)
par(cex=0.8)
plot.seas.temp(mscdata,id="1108447",add.alt=TRUE,style=c(0,1))
```



```
require(seas)
data(mscdata)
par(cex=0.8)
image(seas.sum(mscdata,id="1108447"),style=c(0,1))
```



Lastly, here are is a 3D surface plot that have also been plotted using R:

```

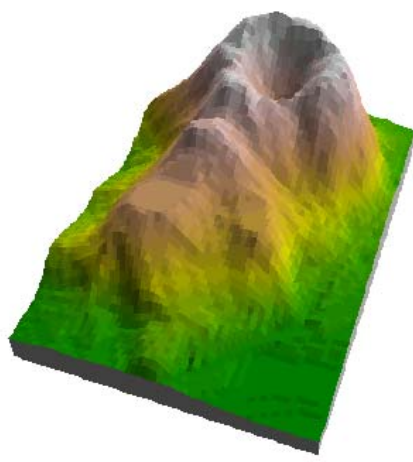
z <- 2 * volcano          # Exaggerate the relief
x <- 10 * (1:nrow(z))    # 10 meter spacing (S to N)
y <- 10 * (1:ncol(z))   # 10 meter spacing (E to W)

z0 <- min(z) - 20
z <- rbind(z0, cbind(z0, z, z0), z0)
x <- c(min(x) - 1e-10, x, max(x) + 1e-10)
y <- c(min(y) - 1e-10, y, max(y) + 1e-10)

fill <- matrix("green3", nr = nrow(z)-1, nc = ncol(z)-1)
fill[ , i2 <- c(1,ncol(fill))] <- "gray"
fill[i1 <- c(1,nrow(fill)) , ] <- "gray"

fcol <- fill
zi <- volcano[ -1,-1] + volcano[ -1,-61] +
      volcano[-87,-1] + volcano[-87,-61] ## / 4
fcol[-i1,-i2] <-
  terrain.colors(20)[cut(zi, quantile(zi, seq(0,1, len =
    21)), include.lowest = TRUE)]
par(mar=rep(.5,4))
persp(x, y, 2*z, theta = 110, phi = 40, col = fcol, scale =
FALSE, ltheta = -120, shade = 0.4, border = NA, box = FALSE)

```



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