# Online Academic Data Analysis Bootcamp Using Open-Access Program R

# Base Plots in R

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Creating a Graph

Histograms and Density Plots

Dot Plots

Bar Plots

Line Charts

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Boxplots

Scatterplots

Advanced Graphics: graphical parameters, axes and text, combining plots

# Introduction: generic plot types in R

**plot()** function is the generic function for **plotting** in R. It can be used to create basic **graphs**.

A simplified format of the function is

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

**x** and **y**: the coordinates of points to plot

**type** : the type of graph to create; Possible values are :

type="p": for **p**oints (by default)

type="l": for **l**ines

type="b": for **b**oth; points are connected by a line

type="o": for both '**o**verplotted';

type="h": for '**h**istogram' like vertical lines

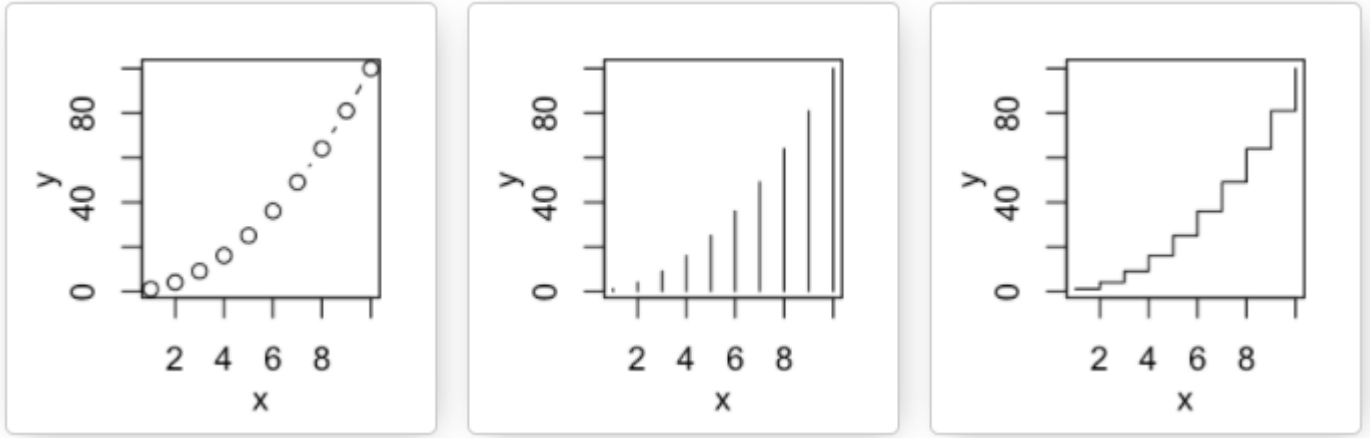
type="s": for **s**tair steps

type="n": for **n**o plotting

# Introduction: generic plot types in R

Examples

```
x<-1:10;
y=x*x
plot(x, y, type="b")
plot(x, y, type="h")
plot(x,y, type="s")
```



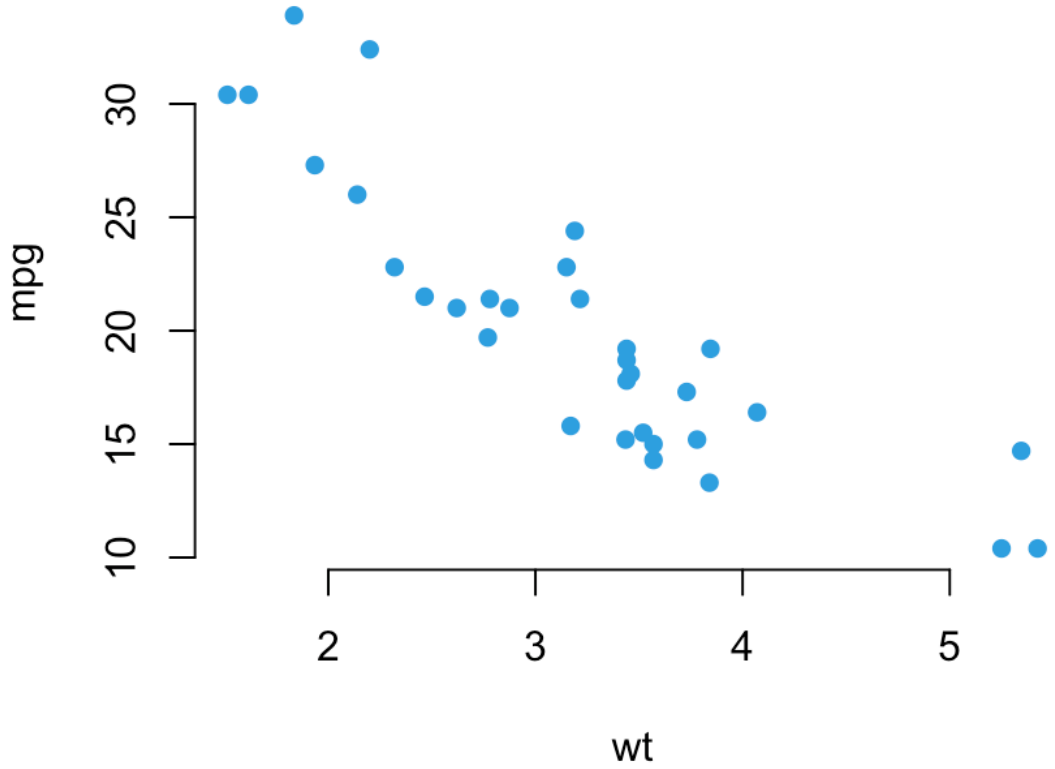
# Creating a Graph

The R base function **plot**() can be used to create graphs.

In R, graphs are typically created interactively.

# Creating a Graph
plot(x = mtcars$wt, y = mtcars$mpg,
pch = 16, frame = FALSE, xlab = "wt",
ylab = "mpg", col = "#2E9FDF")

The plot( ) function opens a graph window and plots weight vs. miles per gallon.

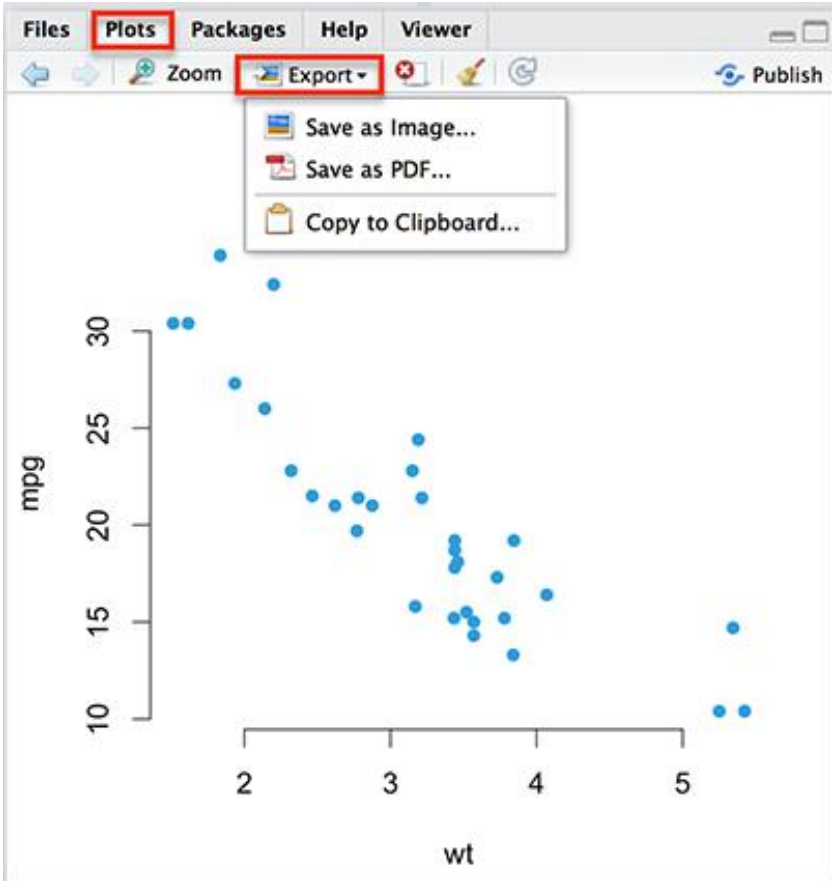


# Saving graphs: RStudio Plots Panel

If you are working with RStudio, the plot can be exported from menu in plot panel (lower right-pannel).

**Plots panel –> Export –> Save as Image or Save as PDF**

The choose directory and change file name



# Saving graphs: R codes to redirect graphs

It’s also possible to save the graph using R codes as follow:

1. Specify files to save your image using a function such as **jpeg**(), **png**(), **svg**() or **pdf**(). Additional argument indicating the width and the height of the image can be also used.

2. Create the plot

3. Close the file with **dev.off**()

Example 1: saving as pdf

```
# 1. Open a pdf file

 pdf("rplot.pdf")

# 2. Create a plot

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

# 3. Close the pdf file

dev.off()
```

# Saving graphs: R codes to redirect graphs

Example 2: saving as jpeg file

```
# 1. Open jpeg file
jpeg("rplot.jpg", width = 350, height = "350")
 # 2. Create the plot
plot(x = mtcars$wt, y = mtcars$mpg, pch = 16, frame = FALSE, xlab = "wt", ylab = "mpg", col = "#2E9FDF")
# 3. Close the file
dev.off()
```

The R codes in the previous 2 slides saves the files in the current working directory.

# Saving graphs: R codes to redirect graphs

To redirect graphic output use one of the following functions. Use **dev.off( )** to return output to the terminal.

| Function | Output to |
|---|---|
| pdf("mygraph.pdf") | pdf file |
| win.metafile("mygraph.wmf") | windows metafile |
| png("mygraph.png") | png file |
| jpeg("mygraph.jpg") | jpeg file |
| bmp("mygraph.bmp") | bmp file |
| postscript("mygraph.ps") | postscript file |

# Create histogram plots: hist()

Histograms display the distribution of a continuous variable by dividing up the range of scores into a specified number of bins on the x-axis and displaying the frequency of scores in each bin on the y-axis.

A histogram can be created using the function hist(), which simplified format is as follow:

hist(x, breaks = "Sturges")

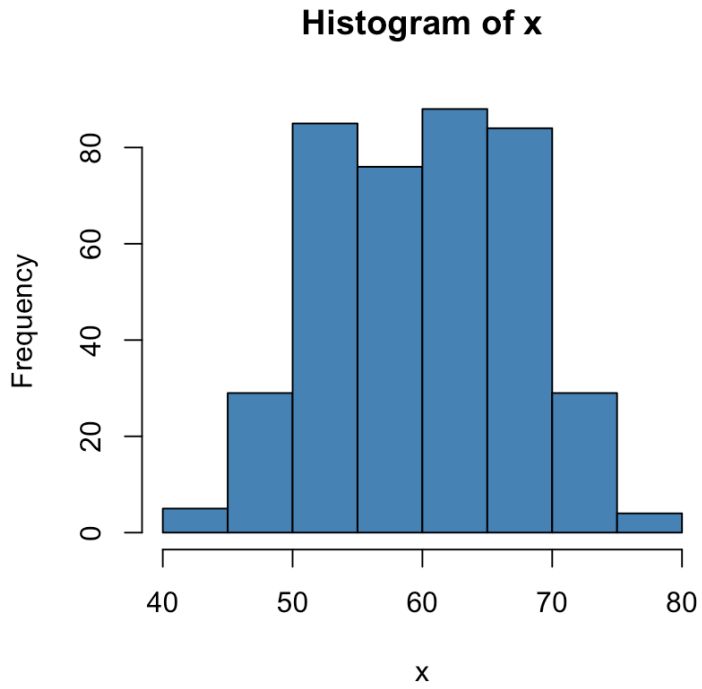
Where:

**x**: a is numeric vector

**breaks**: breakpoints between histogram cells

Example:

```
x <- mtcars$mpg

hist(x, col = "steelblue", frame = FALSE)
```

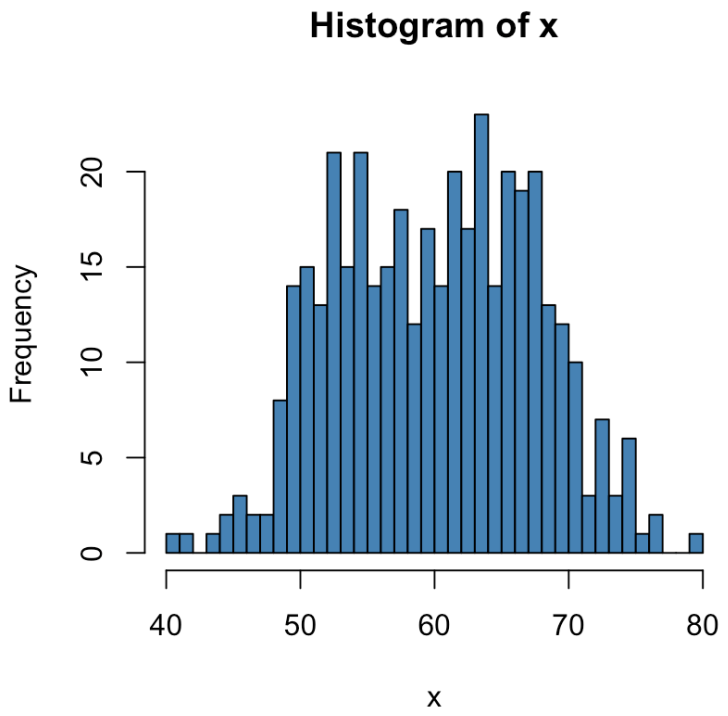


# Create histogram plots: hist()

# Change the number of breaks

```
hist(x, col = "steelblue", frame = FALSE, breaks = 30)
```

Histograms can be a poor method for determining the shape of a distribution – they are strongly affected by the number of bins used.



# Kernel Density Plots: density()

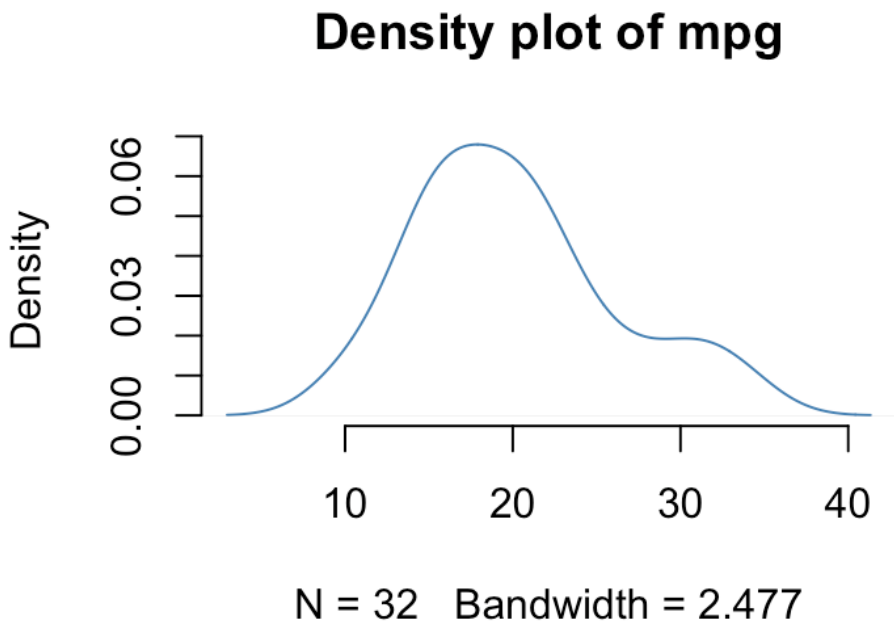
**Kernel density plots** are usually a much more effective than **histograms** way to view the distribution of a variable.

Kernel density estimation is a nonparametric method for estimating the probability density function of a random variable.

The function **density**() is used to estimate kernel density.

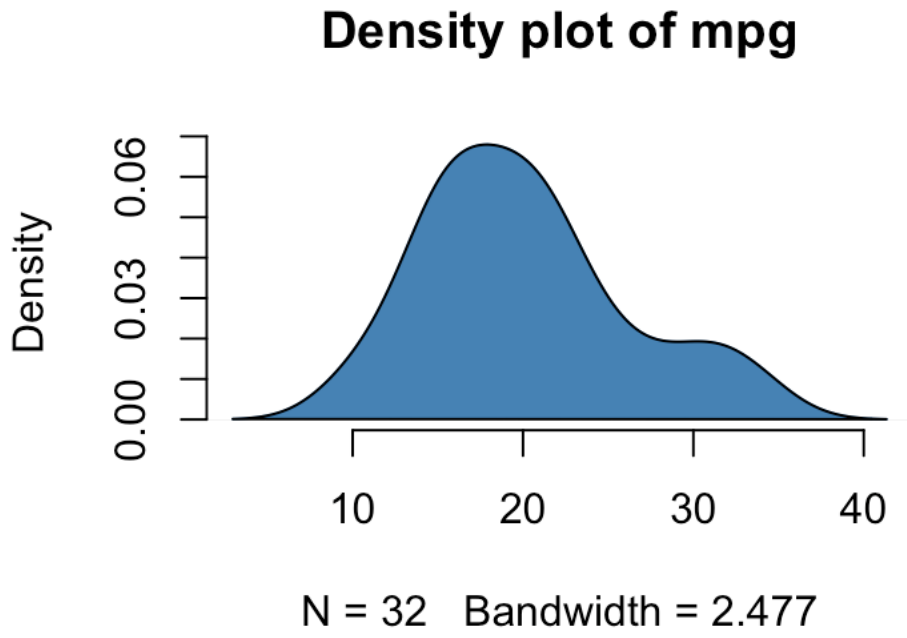
Create the plot using **plot(density(*x*))** where *x* is a numeric vector.

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



# Kernel Density Plots: density()

```
# Fill the density plot using polygon()

plot(dens, frame = FALSE, col = "steelblue",
main = "Density plot of mpg")

polygon(dens, col = "steelblue")
```



# Dot Plots: dotchart()

The function **dotchart**() is used to draw a cleveland dot plot.

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

**x**: numeric vector or matrix

**labels**: a vector of labels for each point.

**groups**: a grouping variable indicating how the elements of x are grouped.

**gcolor**: color to be used for group labels and values.

**color**: the color(s) to be used for points and labels.

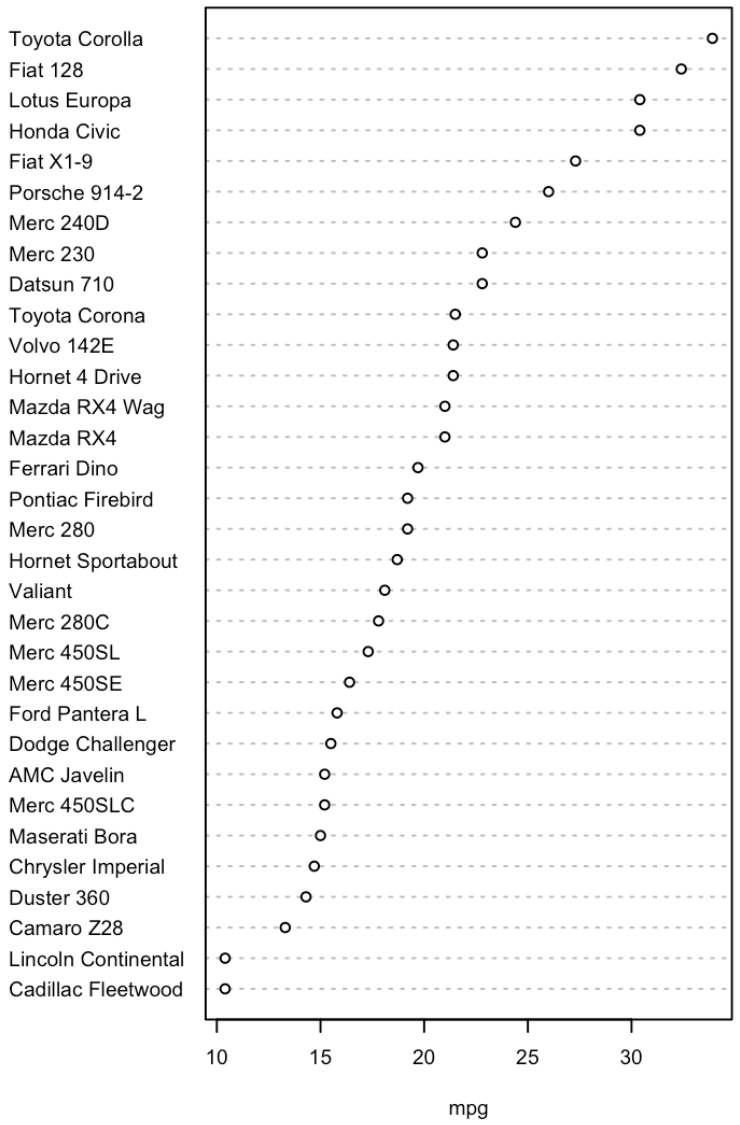
**cex:** controls the size of the labels.

# Dot Plots: dotchart()

**Dot chart of one numeric vector**

# Dot chart of a single numeric vector

```
dotchart(mtcars$mpg, labels =
row.names(mtcars),  cex = 0.6, xlab = "mpg")
```



# Dot Plots: dotchart()

**Dot chart of one numeric**

Plot and color by groups cyl

#Groups

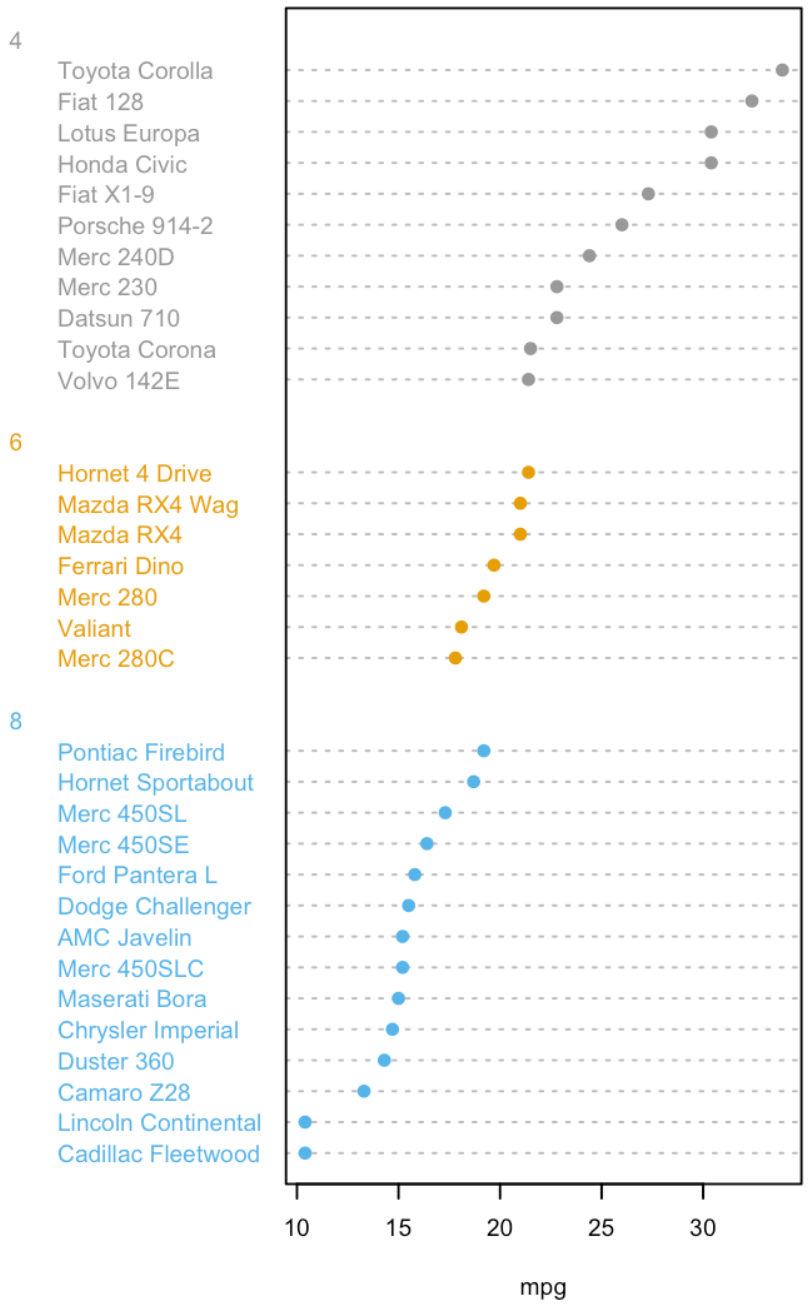
```
grps <- as.factor(mtcars$cyl)
```

#Colours for each group

```
my_cols <- c("#999999", "#E69F00", "#56B4E9")
```

#Plot

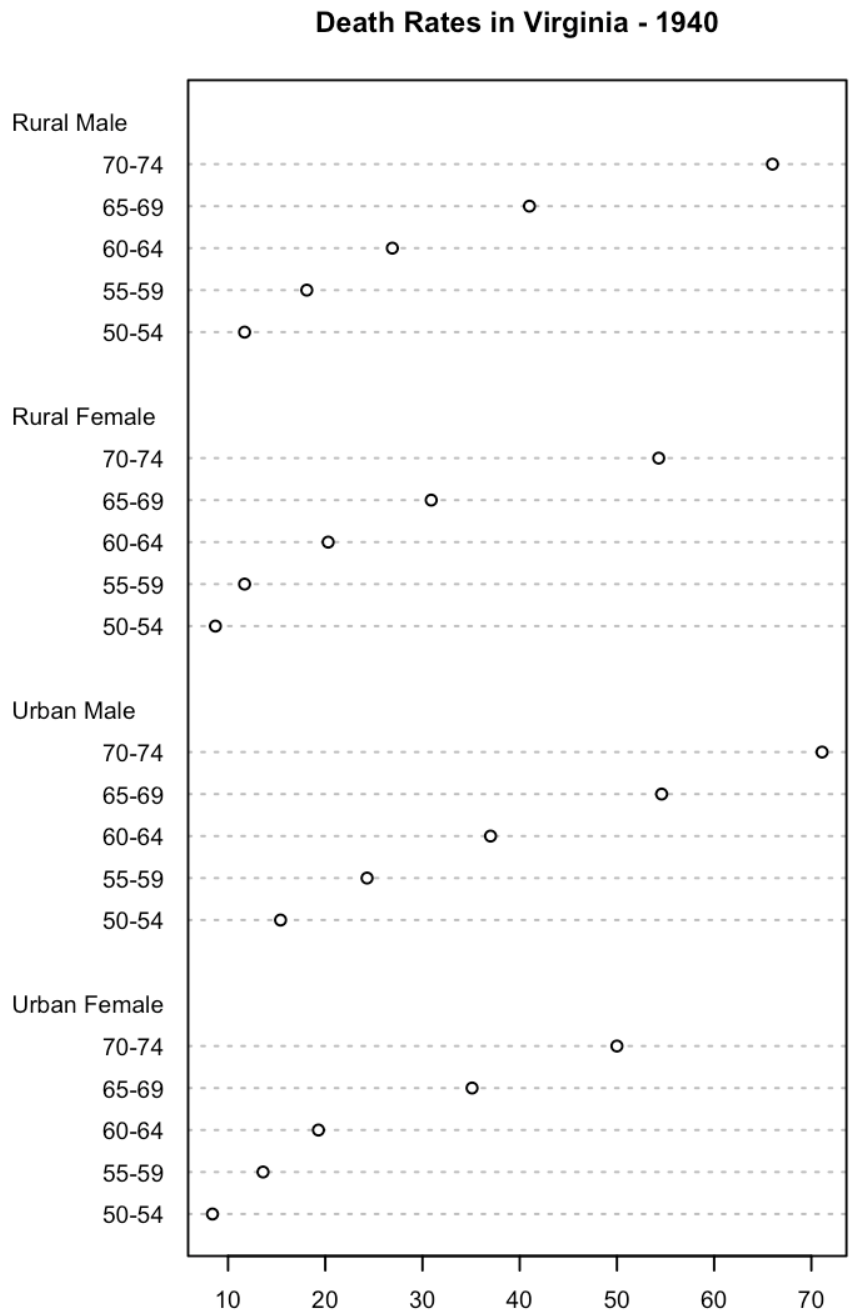
```
dotchart(mtcars$mpg, labels = row.names(mtcars), groups = grps, gcolor = my_cols, color = my_cols[grps], cex = 0.6, pch = 19, xlab = "mpg")
```



# Dot Plots: dotchart()

**Dot chart of a matrix**

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



# Bar Plots

Create barplots with the **barplot(*height*)** function

***height* is a vector or matrix**

If **height is a vector:** the values determine the heights of the bars in the plot.

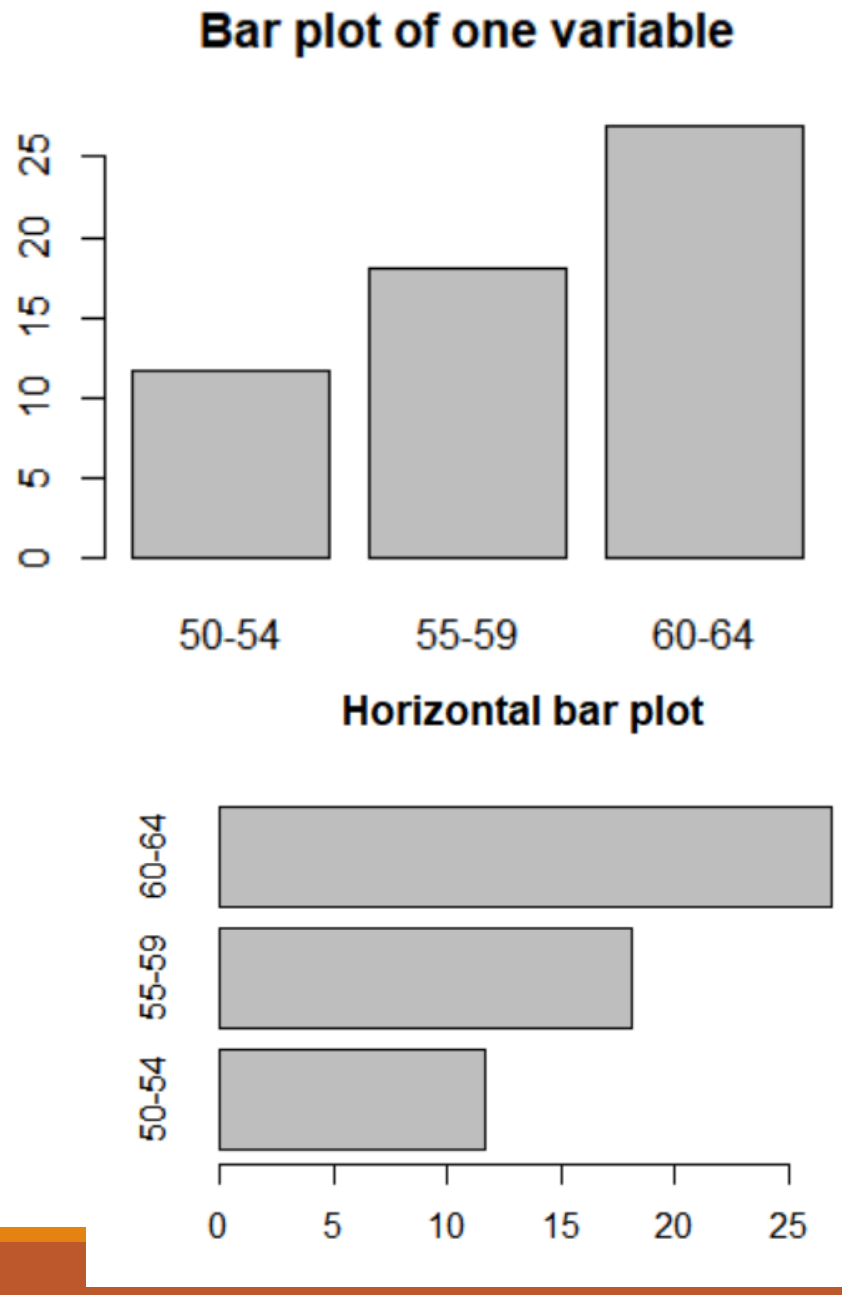
If **height is a matrix** and the option **beside=FALSE** then each bar of the plot corresponds to a column of height, with the values in the column giving the heights of stacked "sub-bars".
If **height is a matrix** and **beside=TRUE**, the values in each column are juxtaposed rather than stacked

**names.arg=(*character vector*)** labels the bars

**horiz=TRUE** creates a horizontal barplot

# Bar Plots:basic

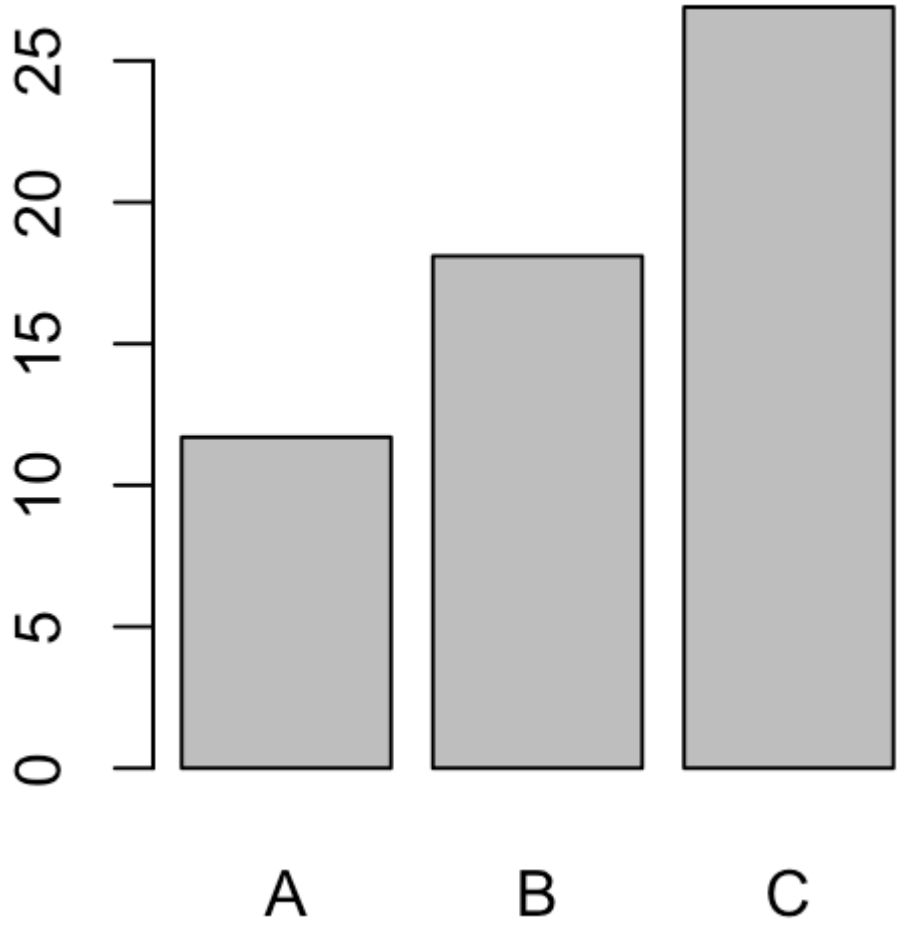
```
# Subset
x <- VADeaths[1:3, "Rural Male"]
x
## 50-54 55-59 60-64
##  11.7  18.1  26.9
# Bar plot of one variable
barplot(x)
# Horizontal bar plot
barplot(x, horiz = TRUE)
```



# Bar Plots

**Change group names**

```
barplot(x, names.arg = c("A", "B", "C"))
```



# Bar Plots

**Change color**

**# Change border and fill color using one single color**

```
barplot(x, col = "white", border = "steelblue")
```

**# Use different colors for each group**

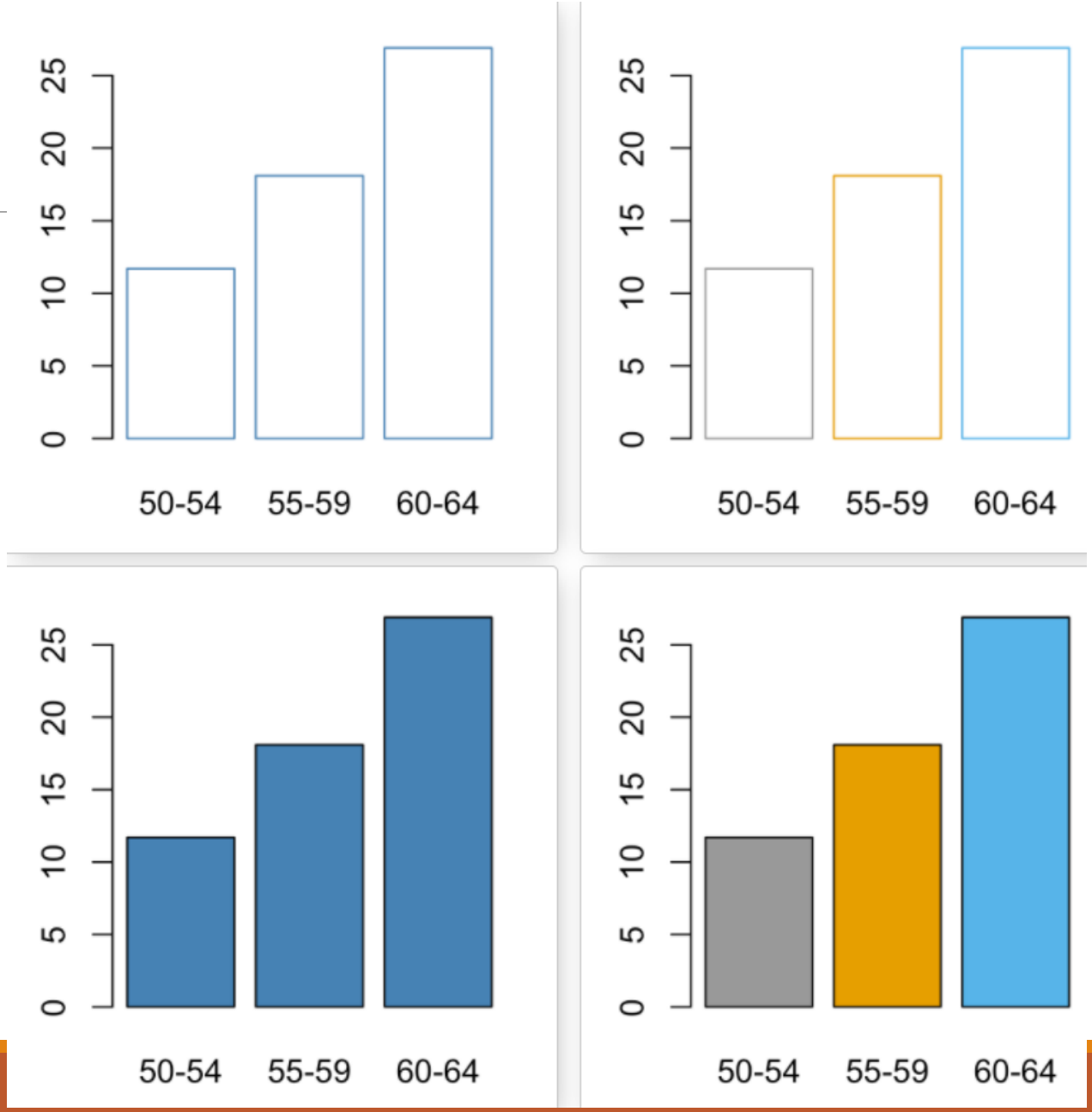
```
barplot(x, col = "white",
        border = c("#999999", "#E69F00", "#56B4E9"))
```

**# Change fill color : single color**

```
barplot(x, col = "steelblue")
```

**# Change fill color: multiple colors**

```
barplot(x, col = c("#999999", "#E69F00", "#56B4E9"))
```



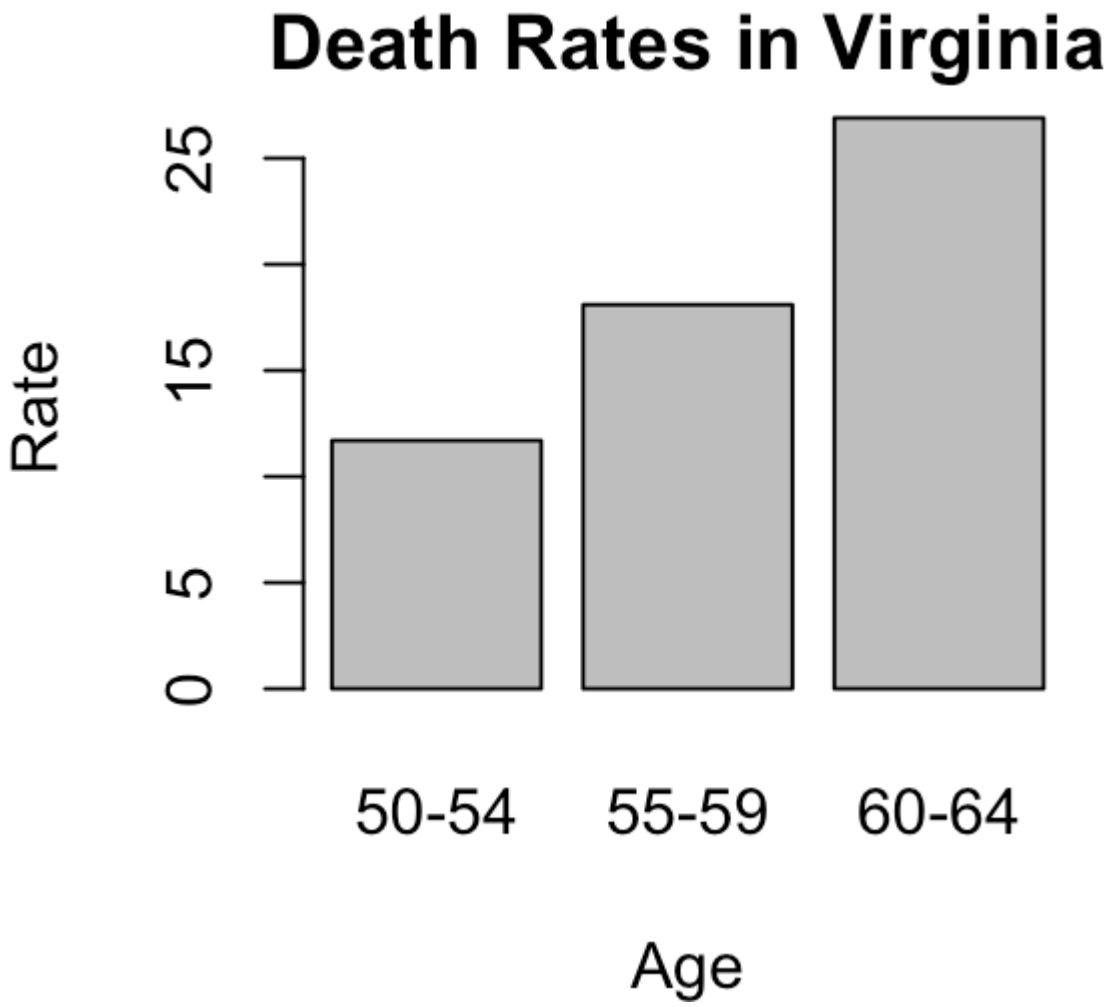
# Bar Plots

**Change main title and axis labels**

# Change axis titles

# Change color (col = "gray") and remove frame

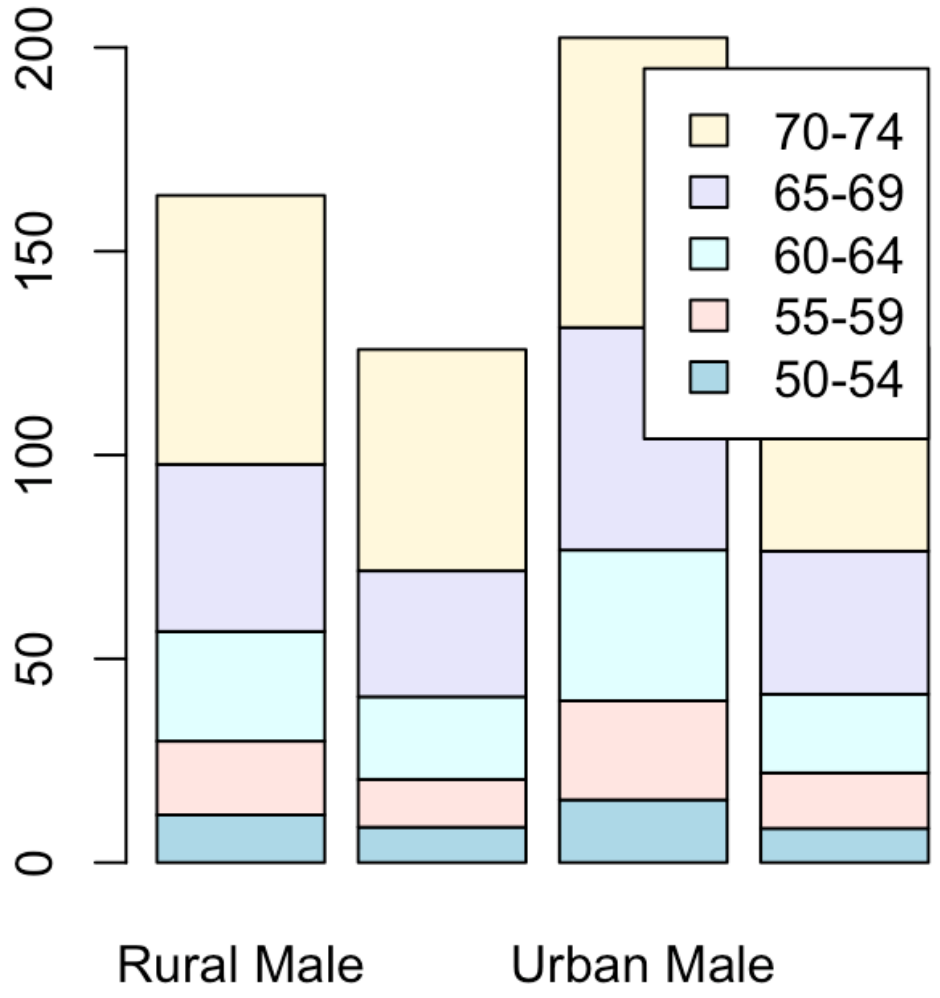
barplot(x, main = "Death Rates in Virginia",  xlab = "Age", ylab = "Rate")



# Bar Plots

**Stacked bar plots**

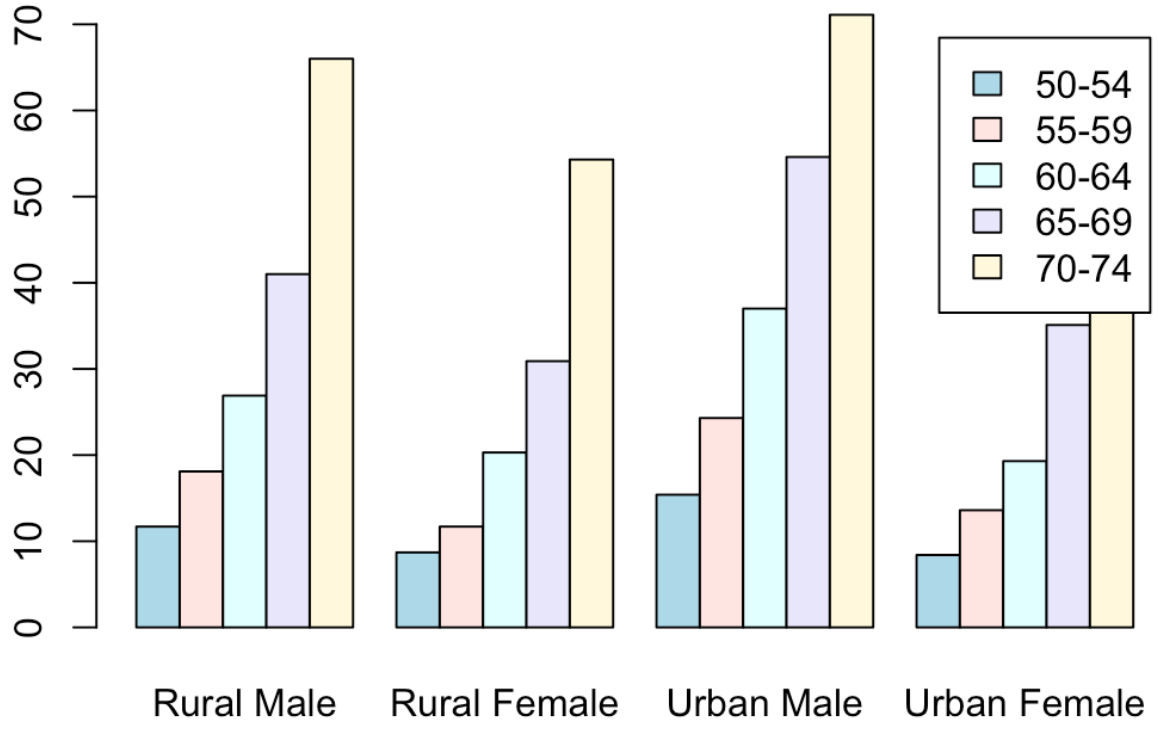
```
barplot(VADeaths,   col = c("lightblue", "mistyrose", "lightcyan",  "lavender", "cornsilk"),       legend = rownames(VADeaths))
```



# Bar Plots

**Grouped bar plots**

```
barplot(VADeaths,      col =
c("lightblue", "mistyrose", "lightcyan",

"lavender", "cornsilk"),   legend =
rownames(VADeaths), beside = TRUE)
```



# Bar Plots

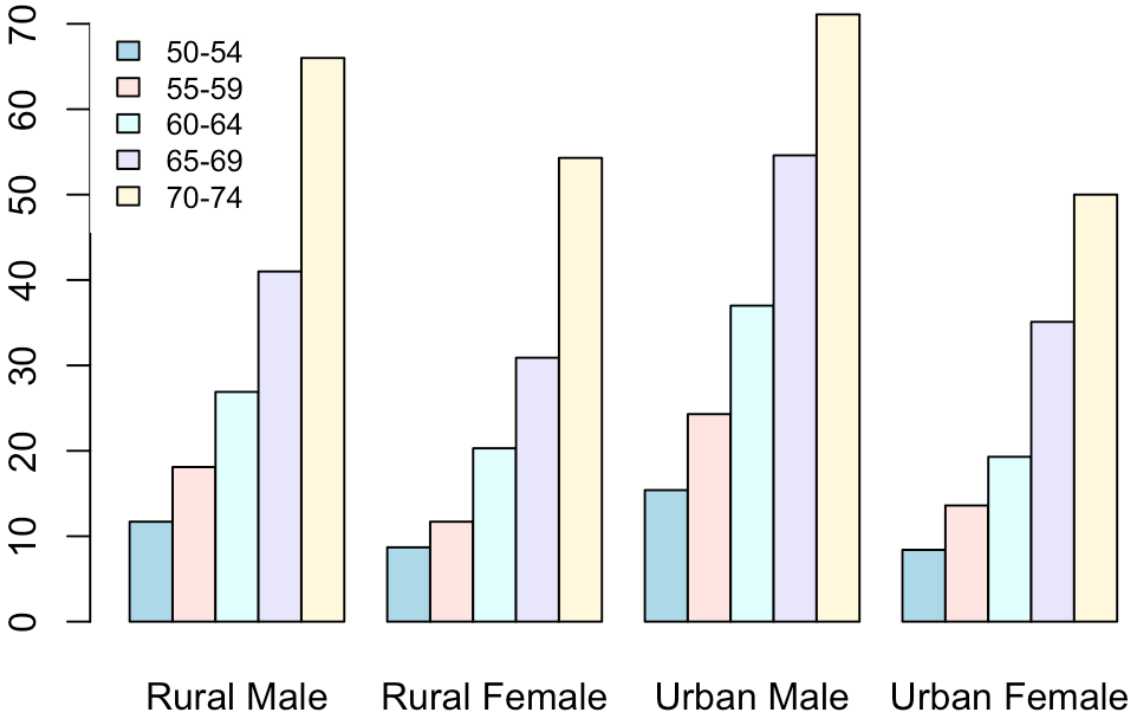
It’s also possible to add legends to a plot using the function legend() as follow.

```
# 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)
```

Notes:

**box.lty = 0**: Remove the box around the legend

**cex = 0.8**: legend text size



# Line Plots: plot() and lines()

The simplified format of plot() and lines() is as follow.

plot(x, y, type = "l", lty = 1)

lines(x, y, type = "l", lty = 1)

**x, y**: coordinate vectors of points to join

**type**: character indicating the type of plotting. Allowed values are:

- “p” for points
- “l” for lines
- “b” for both points and lines
- “c” for empty points joined by lines
- “o” for overplotted points and lines
- “s” and “S” for stair steps
- “n” does not produce any points or lines

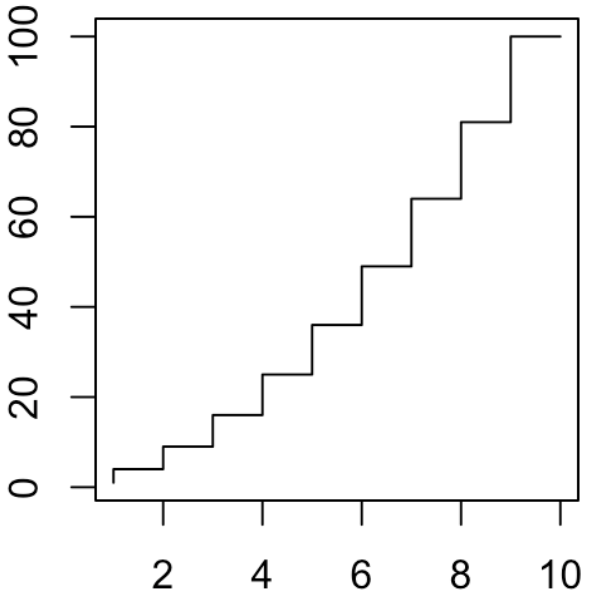
**lty**: line types.

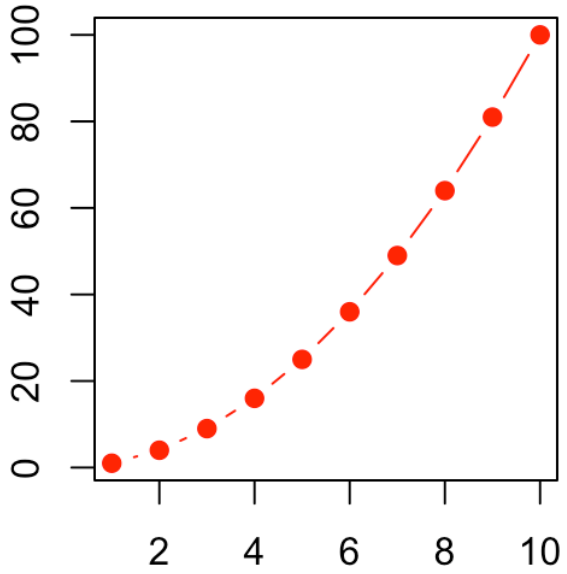
Line types can **either** be specified as **an integer** (0=blank, 1=solid (default), 2=dashed, 3=dotted, 4=dotdash, 5=longdash, 6=twodash)

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).

# Line Plots: plot() and lines()

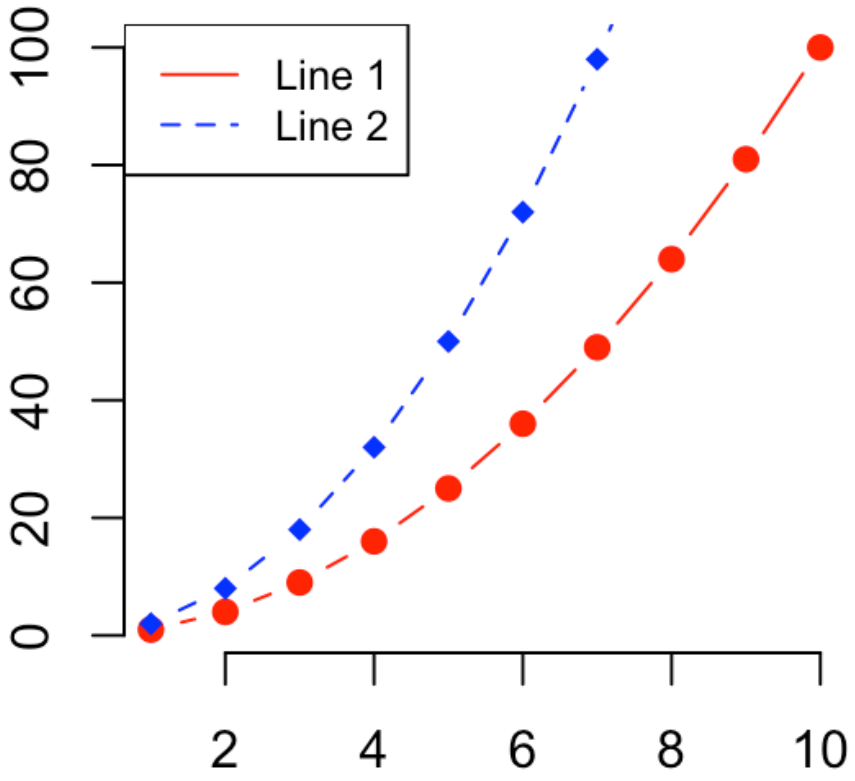
```
# Create some variables

x <- 1:10

y1 <- x*x

y2  <- 2*y1

# Create a basic stair steps plot

plot(x, y1, type = "S")

# Show both points and line

plot(x, y1, type = "b", pch = 19, col = "red",
xlab = "x", ylab = "y")
```



# Line Plots: plot() and lines()

```
# Create a first line

plot(x, y1, type = "b", frame = FALSE, pch = 19,
col = "red", xlab = "x", ylab = "y")

# Add 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)
```



# Pie Charts

**Pie charts are not recommended:** their features are somewhat limited.

Bar or dot plots recommended over pie charts because people are able to judge **length** more accurately than **volume**.

Pie charts are created with the function **pie(*x*, labels=)**

***x*** is a non-negative numeric vector indicating the area of each slice

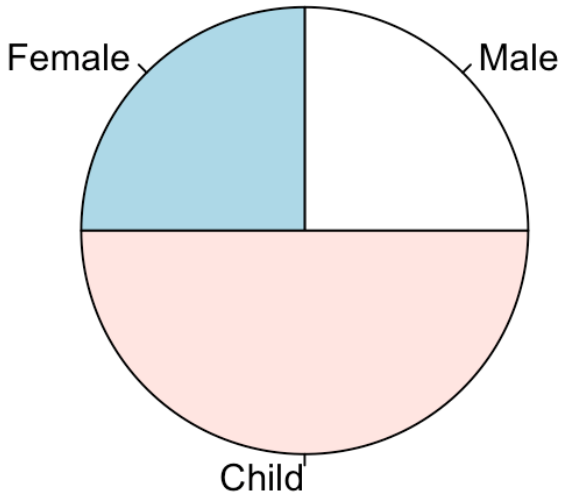
**labels**= notes a character vector of names for the slices.

**radius**: radius of the pie circle. If the character strings labeling the slices are long it may be necessary to use a smaller radius.

# Pie Charts: basic pie chart

Create some data

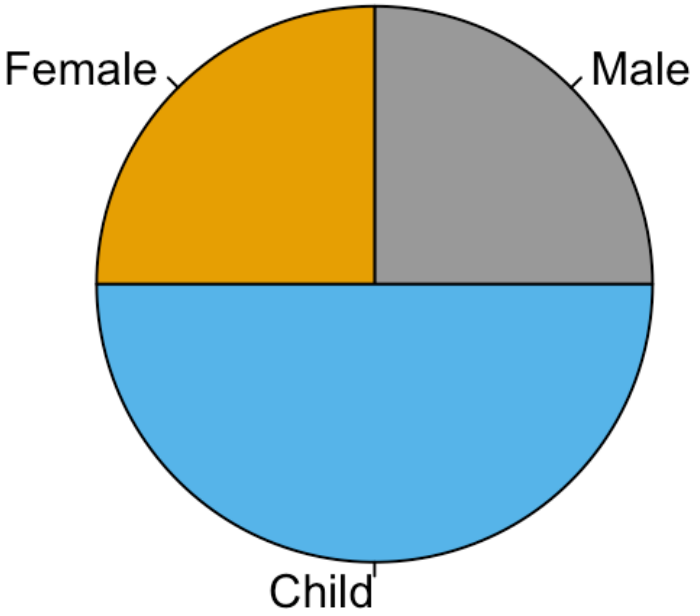
```
df <- data.frame(
  group = c("Male", "Female", "Child"),
  value = c(25, 25, 50)
  )
df


pie(df$value, labels = df$group, radius = 1)
```



# Pie Charts

# Change colors

```
pie(df$value, labels = df$group, radius = 1, col
= c("#999999", "#E69F00", "#56B4E9"))
```



# Pie Charts: Create 3D pie charts: plotix::pie3D()

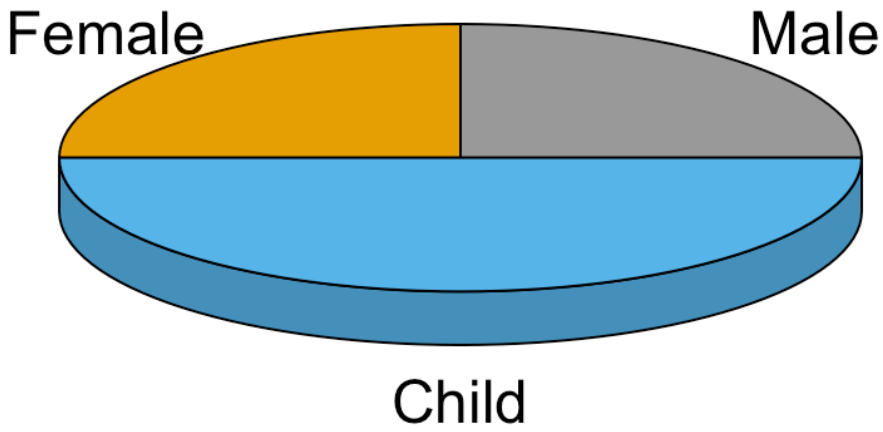
The function **pie3D**()[in **plotrix** package] can be used to draw a 3D pie chart.

**Install** plotrix package:

```
install.packages("plotrix")
```

Use **pie3D():**

```
# 3D pie chart

library("plotrix")

pie3D(df$value, labels = df$group, radius = 1.5,  col = c("#999999", "#E69F00", "#56B4E9"))
```

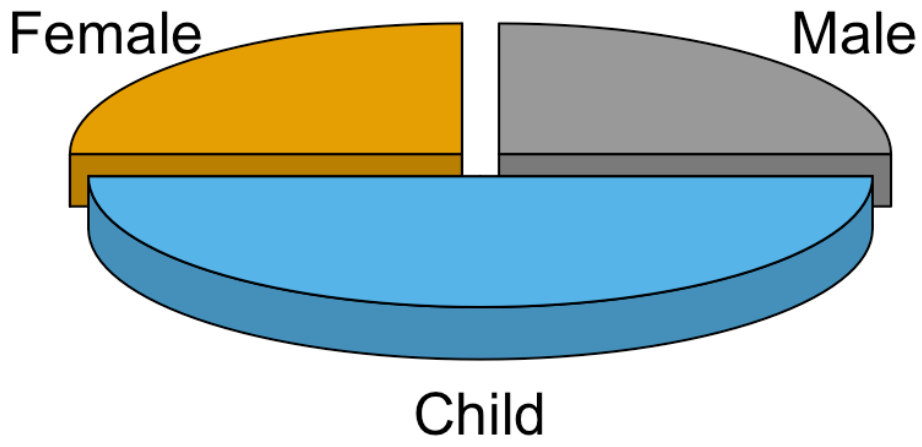


# Pie Charts: Create 3D pie charts: plotix::pie3D()

---

# 3D Exploded Pie Chart

```
pie3D(df$value, labels = df$group, radius = 1.5,
      col = c("#999999", "#E69F00", "#56B4E9"),
      explode = 0.1)
```



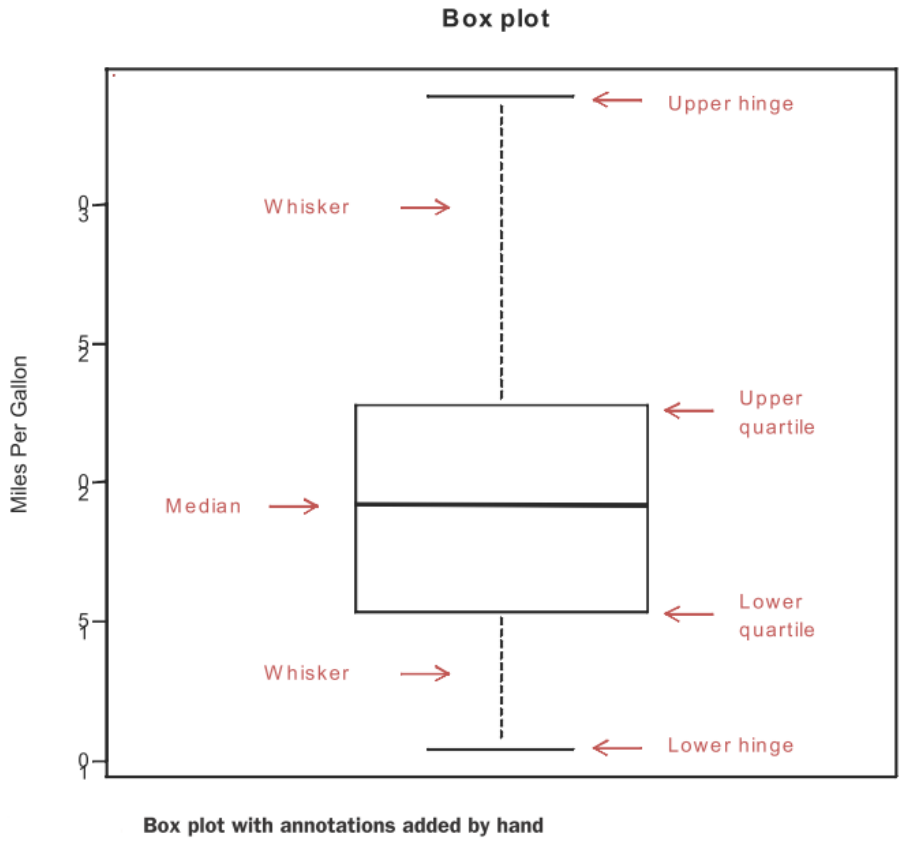
# Boxplots

A "box-and-whiskers" plot describes the distribution of a continuous variable by plotting its five-number summary: the **minimum**, **lower quartile** (25th percentile), **median** (50th percentile), **upper quartile** (75th percentile), and **maximum**.

It can also display observations that may be **outliers** (values outside the range of ± 1.5***IQR**,

where

IQR is the **interquartile range** defined as the **upper quartile minus the lower quartile**).



# Boxplots

Boxplots can be created for **individual variables** or for **variables by group**.

The format is:

boxplot(x, data=),

where **x** is a formula

**data=** denotes the data frame providing the data.

Example formula is y~group - a separate boxplot for numeric variable **y** is generated for each value of **group**.

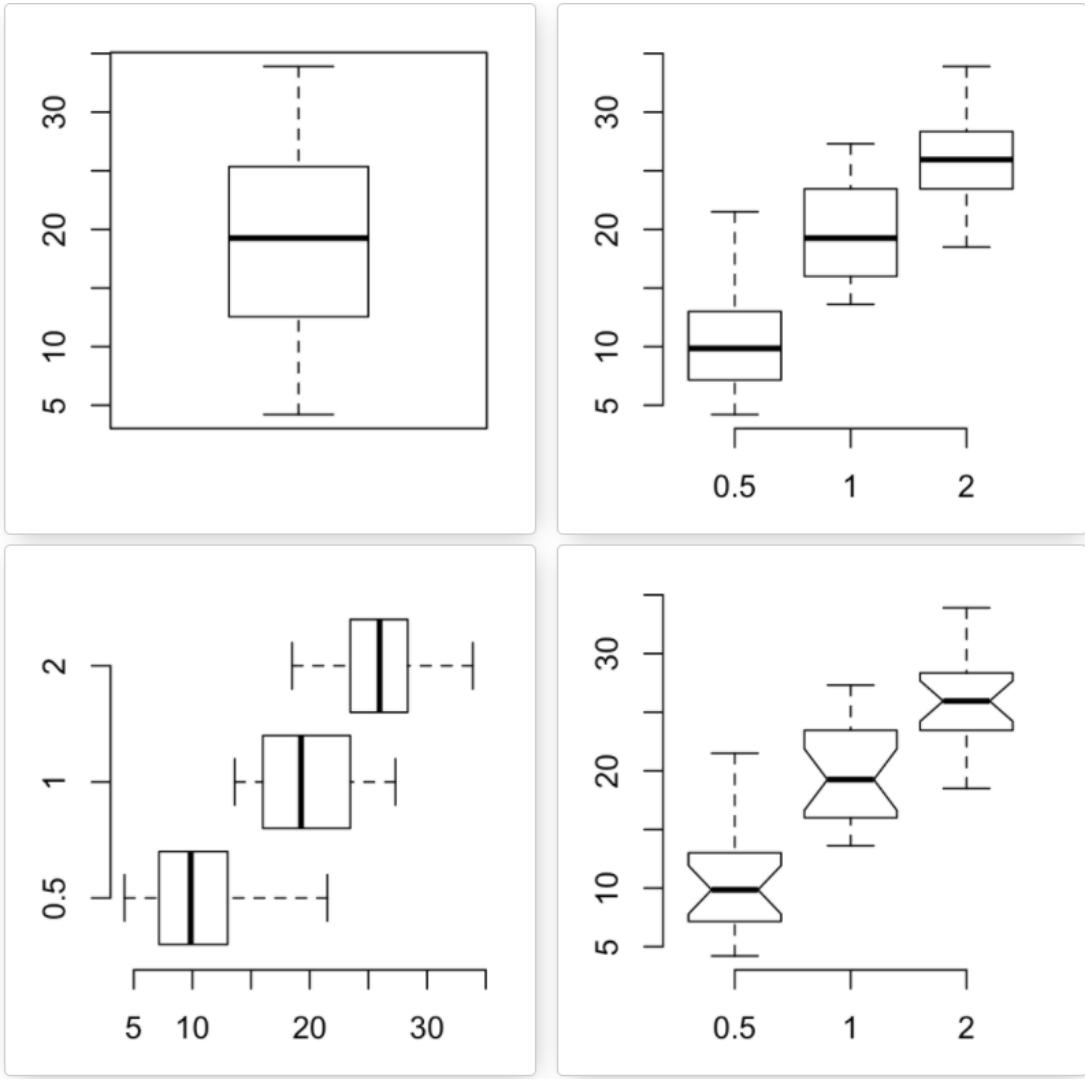
**varwidth=TRUE:** boxplot widths are made proportional to the square root of the samples sizes.

**horizontal=TRUE** reverses the axis orientation.

# Boxplots

Here, we'll use the R built-in ToothGrowth data set.

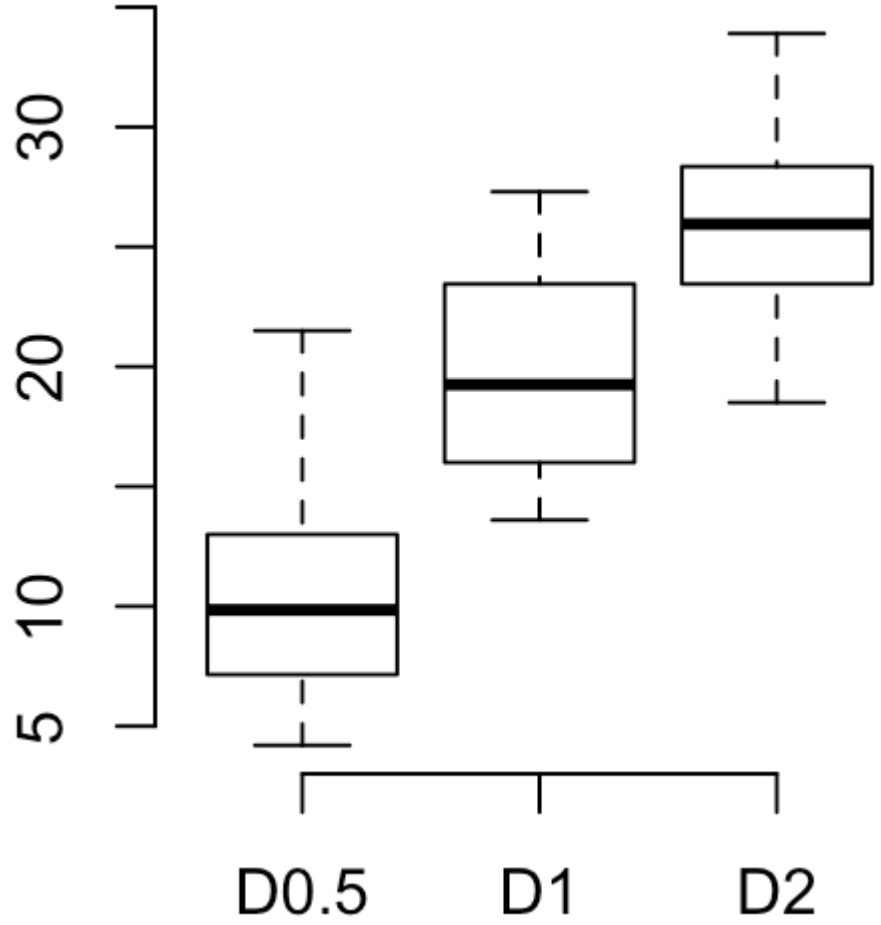
```
# Box plot of one variable
boxplot(ToothGrowth$len)
# Box plots by groups (dose) removing frame
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE)
# Horizontal box plots
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        horizontal = TRUE)
# Notched box plots
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        notch = TRUE)
```



# Boxplots

**Change group names**

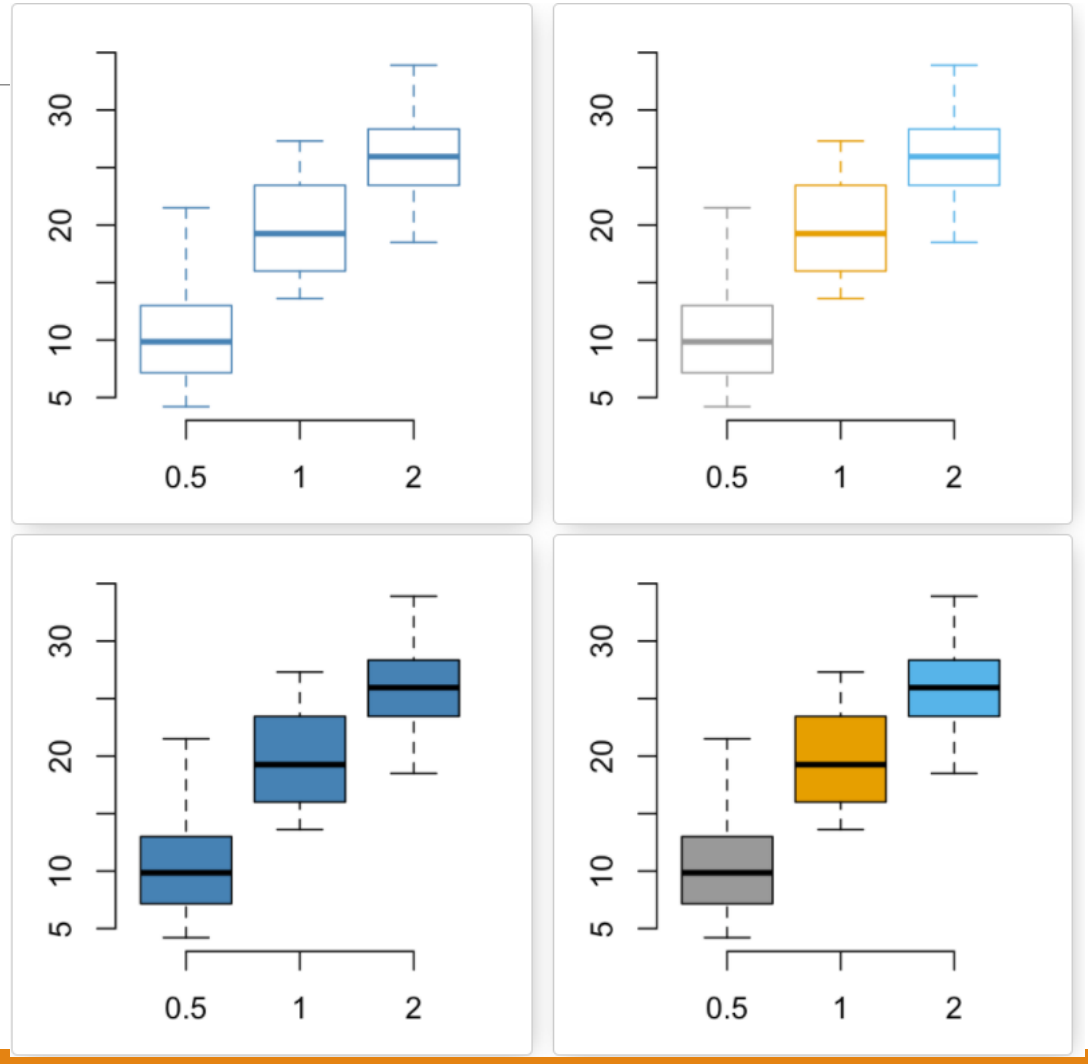
```
boxplot(len ~ dose, data = ToothGrowth, frame
= FALSE, names = c("D0.5", "D1", "D2"))
```



# Boxplots

**Change color**

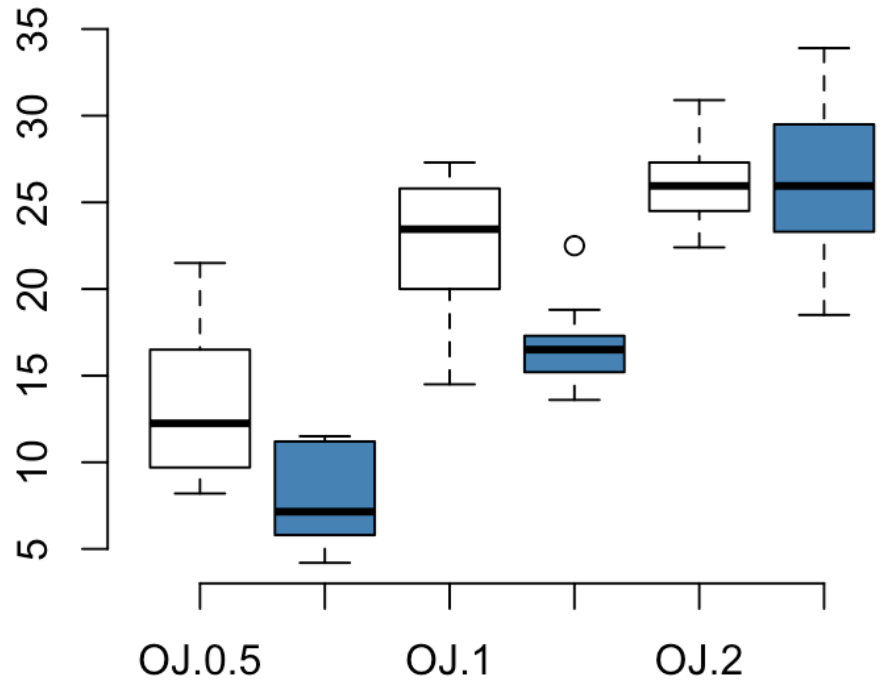
```
# Change the color of border using one single color
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        border = "steelblue")
#  Use different border colors for each group
boxplot(len ~ dose, data = ToothGrowth, frame = FALSE,
        border = c("#999999", "#E69F00", "#56B4E9"))
# Change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



# Boxplots

Box plot with multiple groups

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



# Boxplots

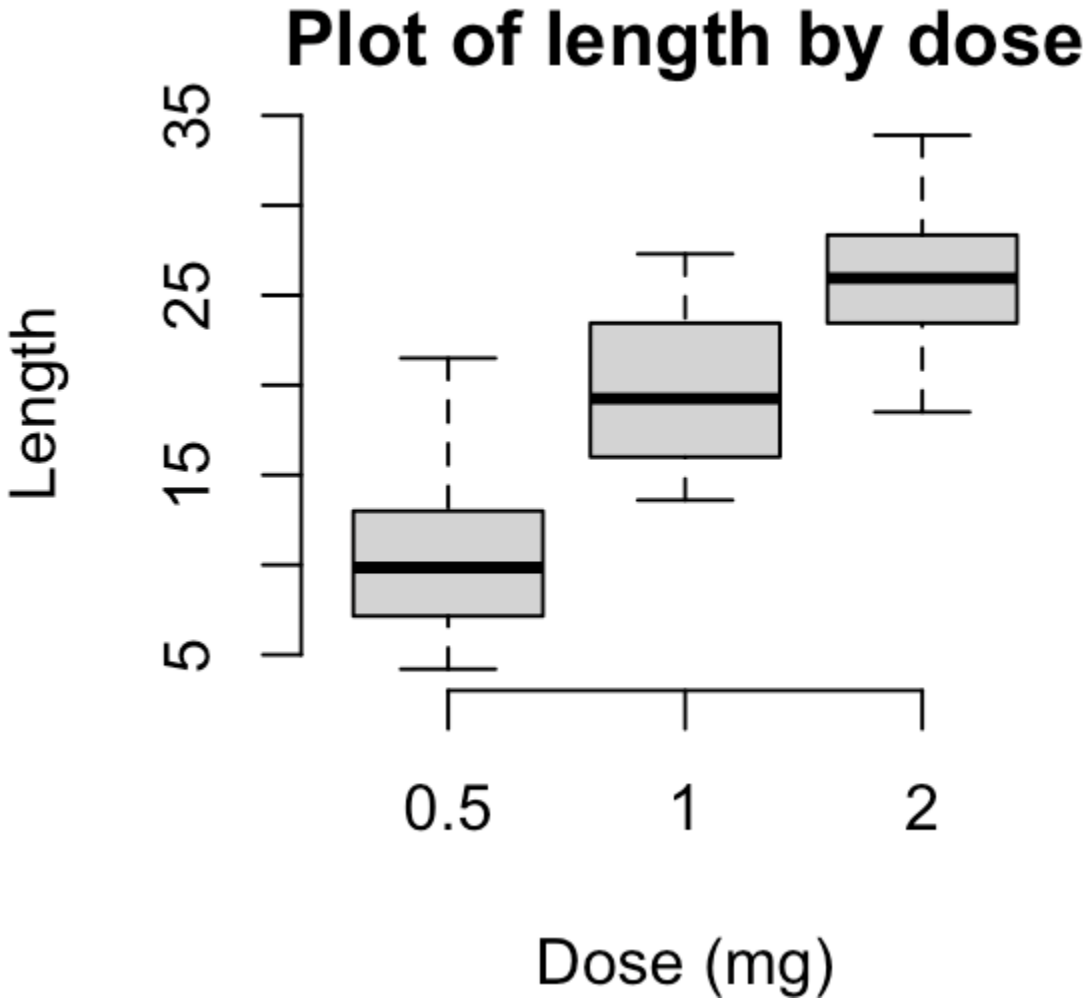
Change main title and axis labels

# 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)
```



# Scatterplots

A convenient method of plotting a bivariate relationship- relationships between two variables (more statistical details to be covered in a later session)

Scatter plots can be created using the function plot(x, y).

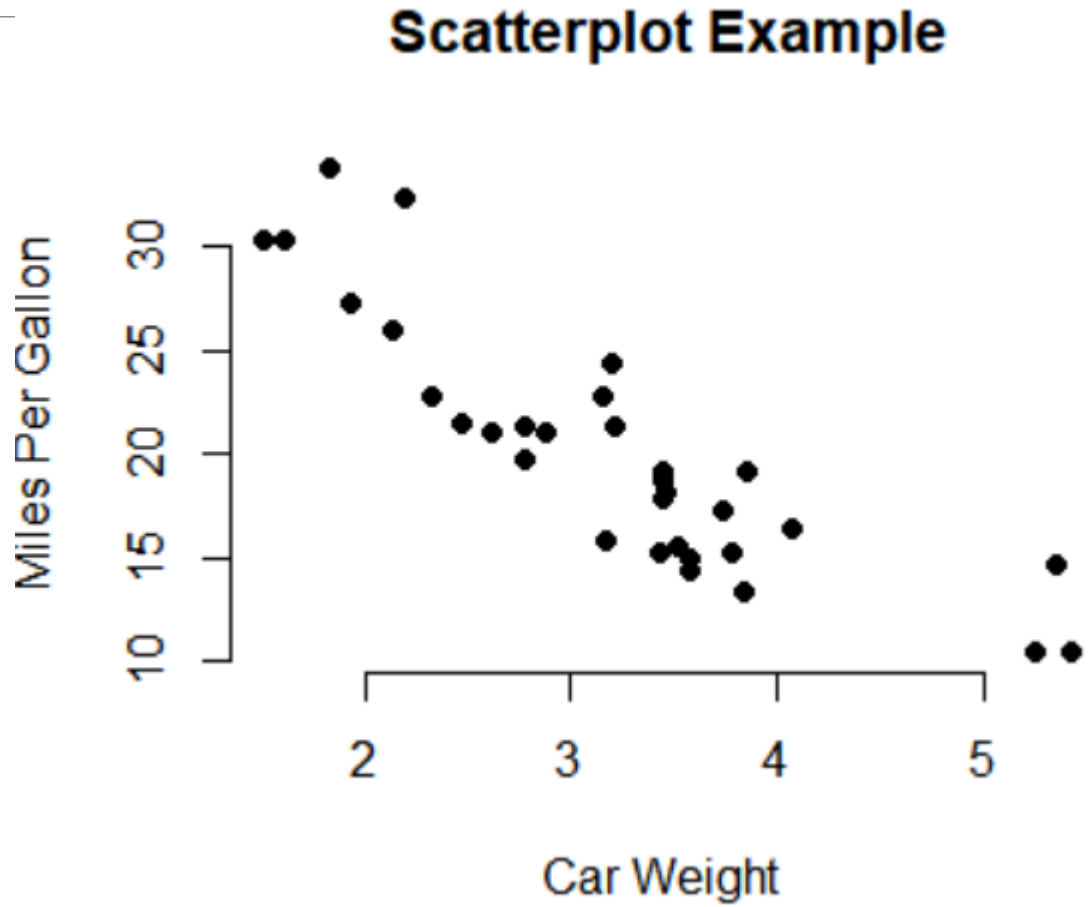
The function lm() will be used to fit linear models between y and x.

A regression line will be added on the plot using the function abline(), which takes the output of lm() as an argument.

Smoothing lines can be added using the function loess().

# Scatterplots: basic

```
attach(mtcars)

# Plot with main and axis titles

# Change point shape (pch = 19) and remove
frame.


plot(wt, mpg, main="Scatterplot Example",
   xlab="Car Weight ", ylab="Miles Per Gallon ",
pch=19, frame=FALSE)
```



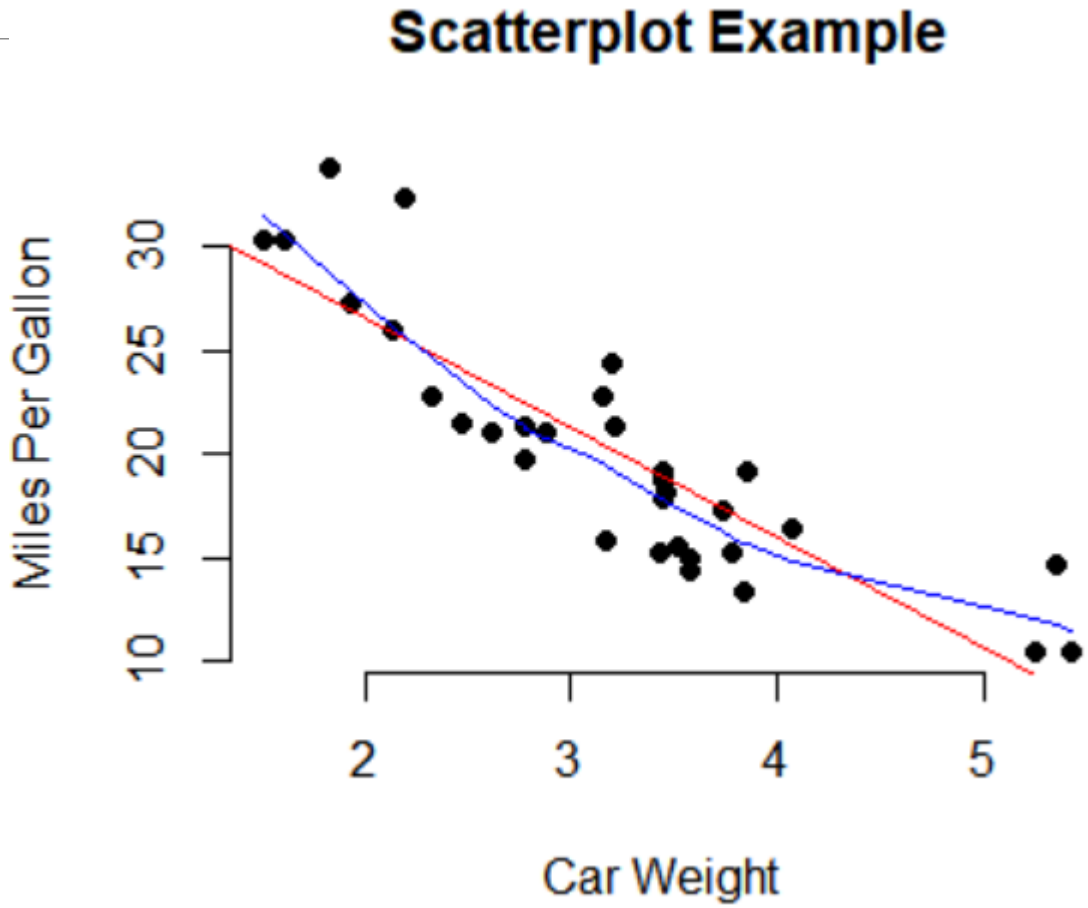
# Scatterplots

# Add regression line

```
plot(wt, mpg, main="Scatterplot Example",
   xlab="Car Weight ", ylab="Miles Per Gallon ",
pch=19, frame=FALSE)
```

# Add fit lines

```
abline(lm(mpg~wt), col="red") # regression line (y~x)
lines(lowess(wt,mpg), col="blue") # lowess line (x,y)
```



# Enhanced scatter plots: car::scatterplot()

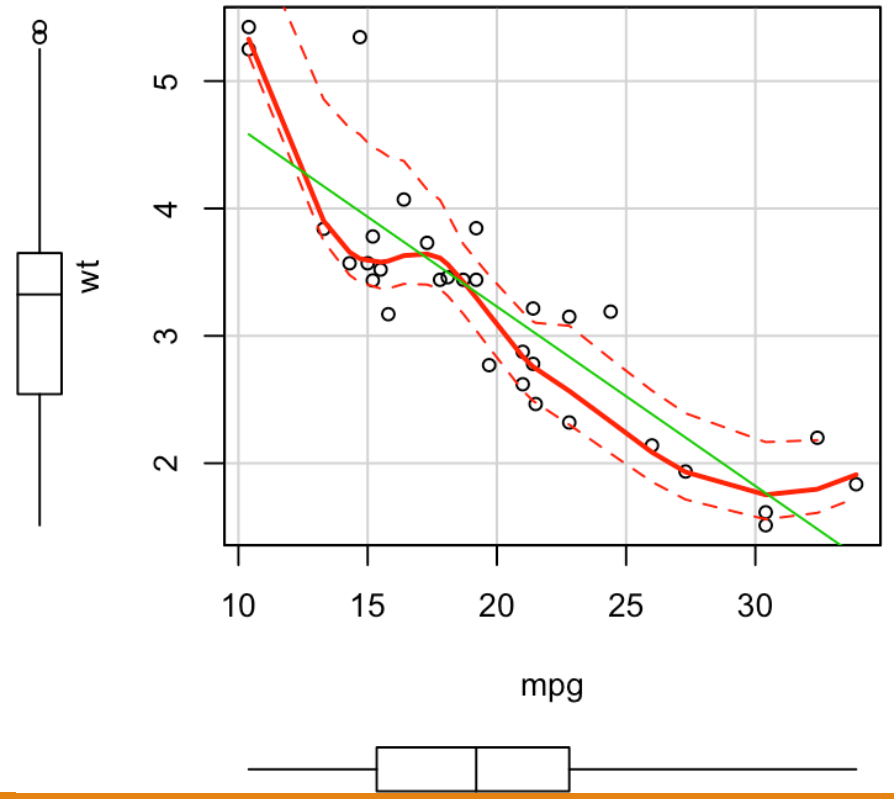
The function scatterplot() [in car package] makes enhanced scatter plots, with box plots in the margins, a non-parametric regression smooth, smoothed conditional spread, outlier identification, and a regression line, …

Install car package:

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

Use scatterplot() function:

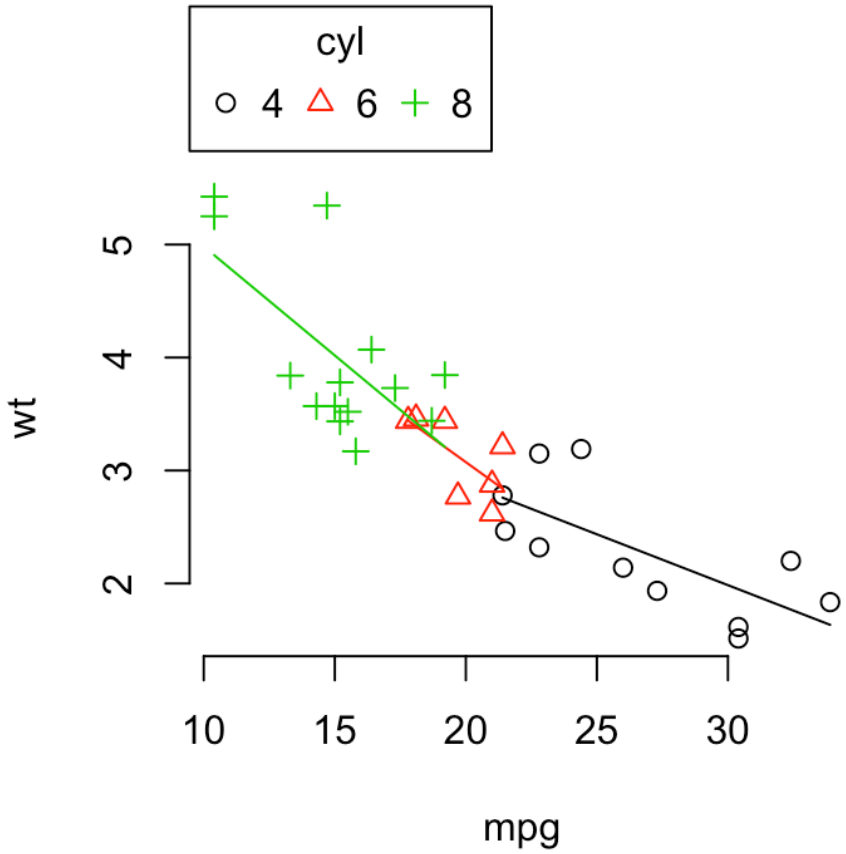
```
library("car")

scatterplot(wt ~ mpg, data = mtcars)
```



# Scatterplots: grouped

# Scatter plot by groups ("cyl")

```
scatterplot(wt ~ mpg | cyl, data = mtcars,
smoother = FALSE, grid = FALSE, frame =
FALSE)
```

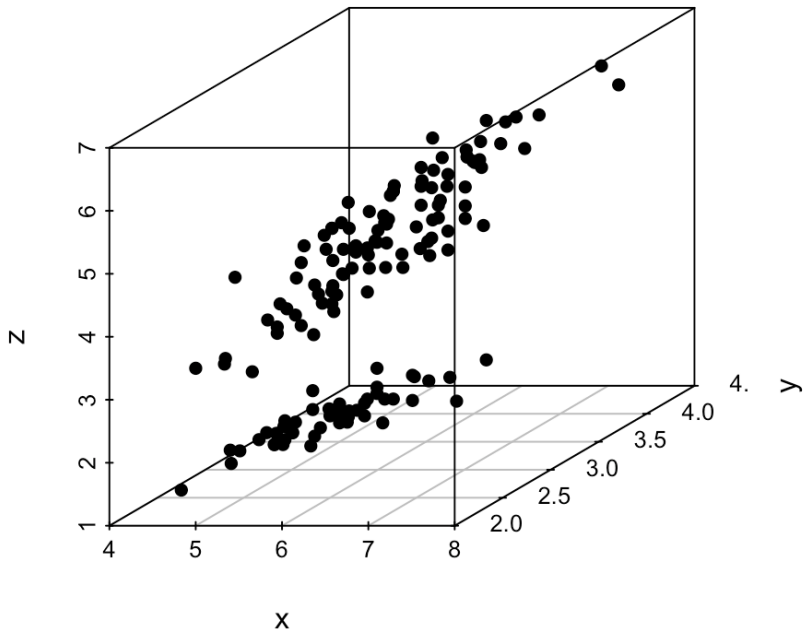


# Scatterplots: 3D

Function scatterplot3D [in scatterplot3D package can be used].

The following R code plots a 3D scatter plot using iris data set.

```
# Prepare the data set
x <- iris$Sepal.Length
y <- iris$Sepal.Width
z <- iris$Petal.Length
grps <- as.factor(iris$Species)
# Plot
library(scatterplot3d)
scatterplot3d(x, y, z, pch = 16)
```



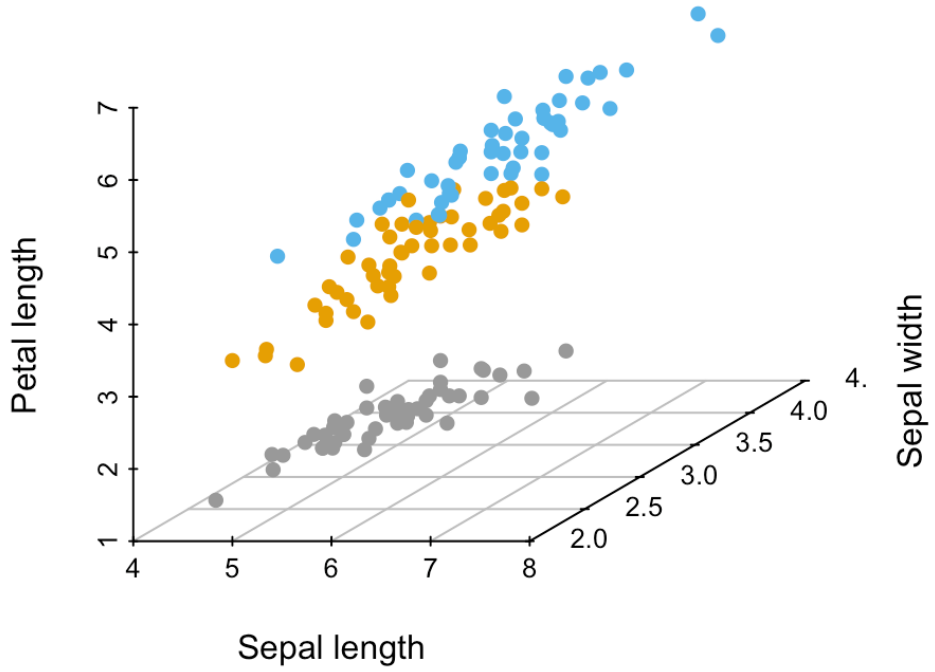
# Scatterplots: 3D

Change color by groups

Add grids and remove the box around the plot

Change axis labels: xlab, ylab and zlab

```
colors <- c("#999999", "#E69F00", "#56B4E9")

scatterplot3d(x, y, z, pch = 16, color = colors[grps],    grid = TRUE, box = FALSE, xlab = "Sepal length", ylab = "Sepal width", zlab = "Petal length")
```



# Advanced Graphics
- Graphical parameters
- Axes and text
- Combining plots

# Graphical Parameters

- You can customize many features of your graphs (fonts, colors, axes, titles) through graphic options.
- The par( ) function: parameter values set here are in effect for the rest of the session or until you change them again.
- The format is par(*optionname*=*value*, *optionname*=*value*, ...)
- # Set a graphical parameter using par()

```
par()             # view current settings
opar <- par()     # make a copy of current settings
par(col.lab="red") # red x and y labels
hist(mtcars$mpg)  # create a plot with these new settings
par(opar)         # restore original settings
```

# Graphical Parameters

- A second way to specify graphical parameters is by providing the *optionname=value* pairs directly to a high-level plotting function. In this case, the options are only in effect for that specific graph.
- # Set a graphical parameter within the plotting function hist(mtcars$mpg, col.lab="red")
- See the help for a specific high level plotting function (e.g. plot, hist, boxplot) to determine which graphical parameters can be set this way.
- The remainder of this section describes some of the more important graphical parameters that you can set.

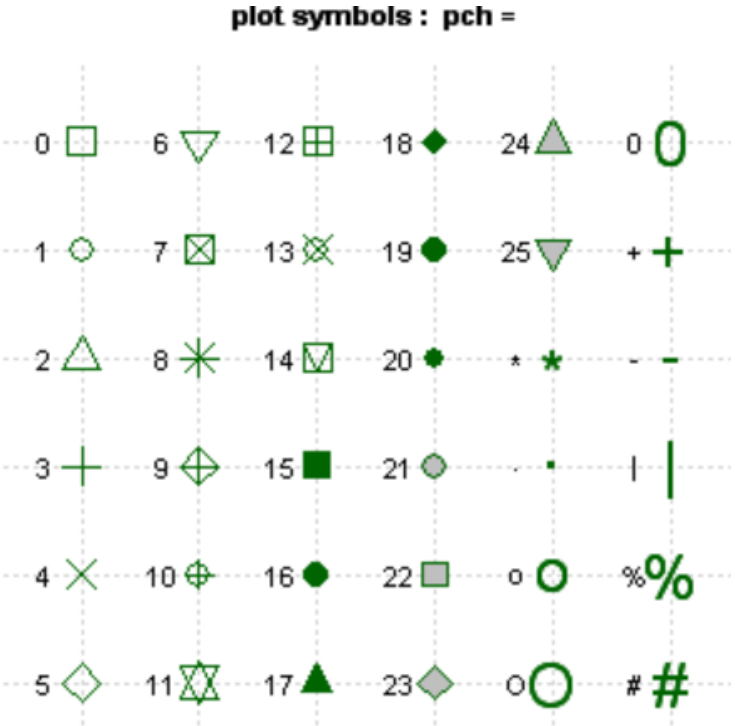
# Graphical Parameters

- **Text and Symbol Size**
- The following options can be used to control text and symbol size in graphs.

| option | description |
|---|---|
| cex | number indicating the amount by which plotting text and symbols should be scaled relative to the default. 1=default, 1.5 is 50% larger, 0.5 is 50% smaller, etc. |
| cex.axis | magnification of axis annotation relative to cex |
| cex.lab | magnification of x and y labels relative to cex |
| cex.main | magnification of titles relative to cex |
| cex.sub | magnification of subtitles relative to cex |

# Graphical Parameters

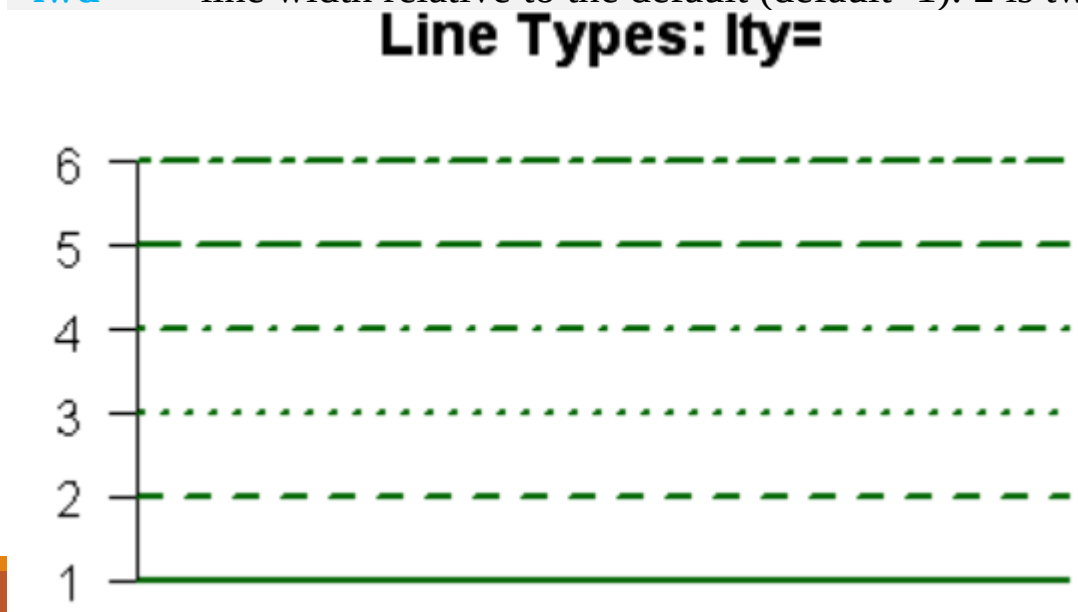
- **PLOTTING SYMBOLS**
- Use the **pch=** option to specify symbols to use when plotting points. For symbols 21 through 25, specify border color (col=) and fill color (bg=).



# Graphical Parameters

- **LINES**
- **You can change lines using the following options. This is particularly useful for reference lines, axes, and fit lines.**

| option | description |
|---|---|
| lty | line type. see the chart below. |
| lwd | line width relative to the default (default=1). 2 is twice as wide. |



# Graphical Parameters

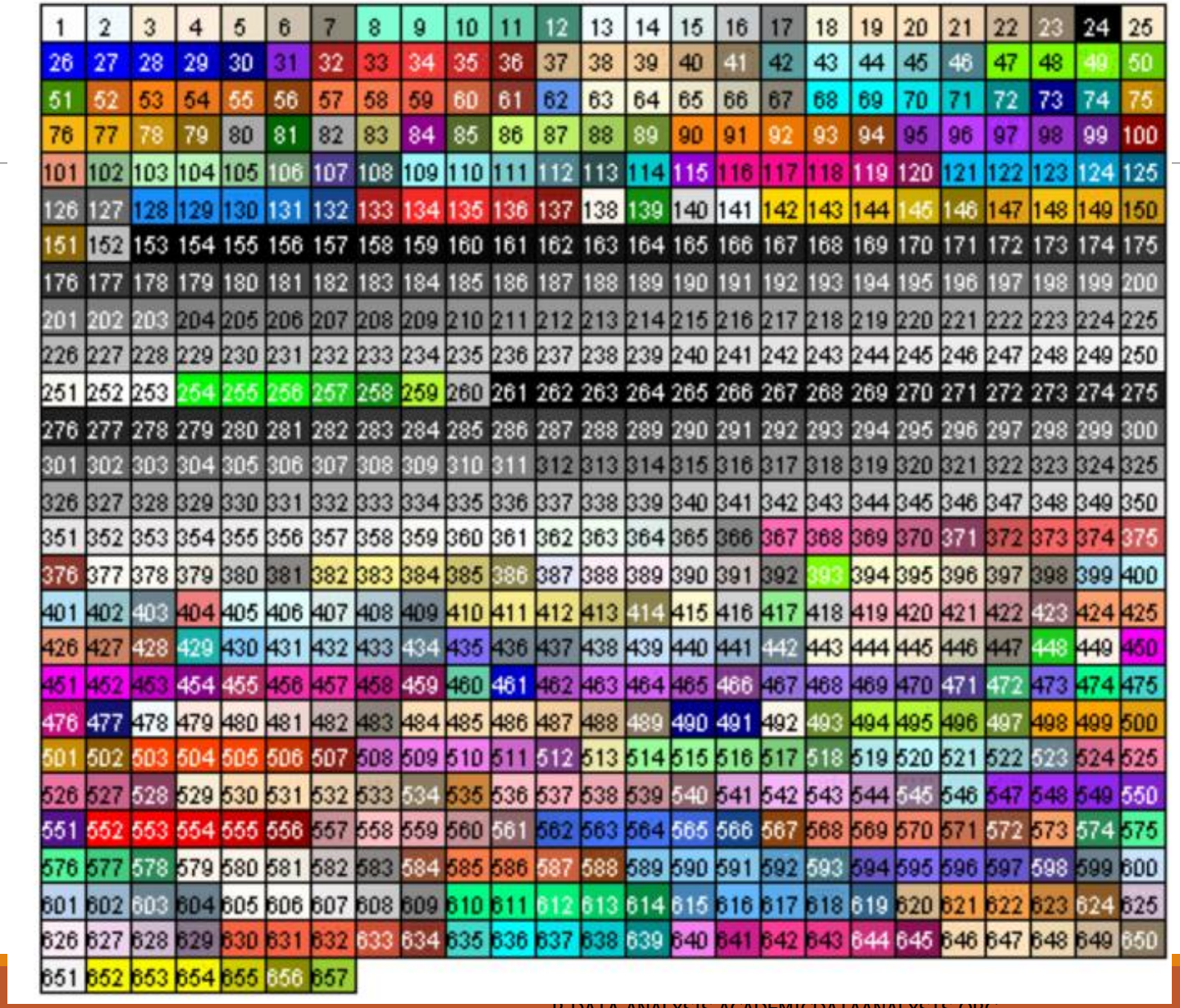
- **COLORS**
- Options that specify colors include the following.

| option | description |
|---|---|
| col | Default plotting color. Some functions (e.g. lines) accept a vector of values that are recycled. |
| col.axis | color for axis annotation |
| col.lab | color for x and y labels |
| col.main | color for titles |
| col.sub | color for subtitles |
| fg | plot foreground color (axes, boxes - also sets col= to same) |
| bg | plot background color |

# Graphical Parameters

- You can specify colors in R by index, name, hexadecimal, or RGB.
  For example col=1, col="white", and col="#FFFFFF" are equivalent.
- The following slide prersents a chart was produced with code developed by Earl F. Glynn.
- You can also create a vector of *n* contiguous colors using the functions **rainbow(*n*)**, **heat.colors(*n*)**, **terrain.colors(*n*)**, **topo.colors(*n*)**, and **cm.colors(*n*)**.
- **colors()** returns all available color names.

# Graphical Parameters: color code chart



# Graphical Parameters

- fonts
- You can easily set font size and style, but font family is a bit more complicated.

| option | description |
|---|---|
| font | Integer specifying font to use for text.<br>1=plain, 2=bold, 3=italic, 4=bold italic, 5=symbol |
| font.axis | font for axis annotation |
| font.lab | font for x and y labels |
| font.main | font for titles |
| font.sub | font for subtitles |
| ps | font point size (roughly 1/72 inch)<br>text size=ps*cex |
| family | font family for drawing text. Standard values are "serif", "sans", "mono", "symbol". Mapping is device dependent. |

# Graphical Parameters

- Axes and Text
- Many high-level plotting functions (plot, hist, boxplot, etc.) allow you to include axis and text options (as well as other graphical paramters). For example
- # Specify axis options within plot()
  plot(*x*, *y*, main="*title*", sub="*subtitle*",
  xlab="*X-axis label*", ylab="*y-axix label*",
  xlim=c(*xmin*, *xmax*), ylim=c(*ymin*, *ymax*))
- For finer control or for modularization, you can use the functions described below.

# Graphical Parameters

- Titles
- Use the title( ) function to add labels to a plot.
- title(main="*main title*", sub="*sub-title*",
  xlab="*x-axis label*", ylab="*y-axis label*")
- Many other ***graphical parameters*** (such as text size, font, rotation, and color) can also be specified in the title( ) function.
- # Add a red title and a blue subtitle. Make x and y
  # labels 25% smaller than the default and green.
  title(main="My Title", col.main="red",
  sub="My Sub-title", col.sub="blue",
  xlab="My X label", ylab="My Y label",
  col.lab="green", cex.lab=0.75)

# Graphical Parameters

If you are going to create a custom axis, you should suppress the axis automatically generated by your high level plotting function.

The option axes=FALSE suppresses both x and y axes. xaxt="n" and yaxt="n" suppress the x and y axis respectively.

# Graphical Parameters

**Axes**

You can create custom axes using the **axis( )** function.

axis(*side*, at=, labels=, pos=, lty=, col=, las=, tck=, ...)

where

| option | description |
|---|---|
| **side** | an integer indicating the side of the graph to draw the axis (1=bottom, 2=left, 3=top, 4=right) |
| **at** | a numeric vector indicating where tic marks should be drawn |
| **labels** | a character vector of labels to be placed at the tickmarks<br>(if NULL, the *at* values will be used) |
| **pos** | the coordinate at which the axis line is to be drawn.<br>(i.e., the value on the other axis where it crosses) |
| **lty** | line type |
| **col** | the line and tick mark color |
| **las** | labels are parallel (=0) or perpendicular(=2) to axis |
| **tck** | length of tick mark as fraction of plotting region (negative number is outside graph, positive number is inside, 0 suppresses ticks, 1 creates gridlines) default is -0.01 |
| (...) | other graphical parameters |

# Graphical Parameters

```
# An Axis Example
# specify the data
x <- c(1:10); y <- x; z <- 10/x
# create extra margin room on the right for an axis
par(mar=c(5, 4, 4, 8) + 0.1)
# plot x vs. y
plot(x, y,type="b", pch=21, col="red", yaxt="n", lty=3, xlab="", ylab="")
# add x vs. 1/x
lines(x, z, type="b", pch=22, col="blue", lty=2)
# draw an axis on the left
axis(2, at=x,labels=x, col.axis="red", las=2)
# draw an axis on the right, with smaller text and ticks
axis(4, at=z,labels=round(z,digits=2),col.axis="blue", las=2, cex.axis=0.7, tck=-.01)
# add a title for the right axis
mtext("y=1/x", side=4, line=3, cex.lab=1,las=2, col="blue")
# add a main title and bottom and left axis labels
title("An Example of Creative Axes", xlab="X values", ylab="Y=X")
```

# Graphical Parameters

Reference Lines

Add reference lines to a graph using the abline( ) function.

abline(h=*yvalues*, v=*xvalues*)

Other ***graphical parameters*** (such as line type, color, and width) can also be specified in the abline( ) function.

```
# add solid horizontal lines at y=1,5,7
  abline(h=c(1,5,7))
  # add dashed blue verical lines at x = 1,3,5,7,9
  abline(v=seq(1,10,2),lty=2,col="blue")
```

Note: You can also use the grid( ) function to add reference lines.

# Graphical Parameters

- **Legend**
- Add a legend with the **legend()** function.
- legend(location, title, legend, ...)
- Common options are described below.

| option | description |
|---|---|
| location | There are several ways to indicate the location of the legend. You can give an **x,y coordinate** for the upper left hand corner of the legend. You can use **locator(1)**, in which case you use the mouse to indicate the location of the legend. You can also use the **keywords** "bottom", "bottomleft", "left", "topleft", "top", "topright", "right", "bottomright", or "center". If you use a keyword, you may want to use **inset=** to specify an amount to move the legend into the graph (as fraction of plot region). |
| title | A character string for the legend title (optional) |
| legend | A character vector with the labels |
| ... | Other options. If the legend labels colored lines, specify **col=** and a vector of colors. If the legend labels point symbols, specify **pch=** and a vector of point symbols. If the legend labels line width or line style, use **lwd=** or **lty=** and a vector of widths or styles. To create colored boxes for the legend (common in bar, box, or pie charts), use **fill=** and a vector of colors. |

# Graphical Parameters

- # Legend Example

```
attach(mtcars)
boxplot(mpg~cyl, main="Milage by Car Weight",
   yaxt="n", xlab="Milage", horizontal=TRUE,
   col=terrain.colors(3))
legend("topright", inset=.05, title="Number of
Cylinders",
   c("4","6","8"), fill=terrain.colors(3), horiz=TRUE)
```

# Graphical Parameters

- **Combining Plots**
- **R** makes it easy to combine multiple plots into one overall graph, using either the **par( )** or **layout( )** function.
- With the **par( )** function, you can include the option **mfrow=c(***nrows, ncols***)** to create a matrix of *nrows x ncols* plots that are filled in by row. **mfcol=c(***nrows, ncols***)** fills in the matrix by columns.
- # 4 figures arranged in 2 rows and 2 columns

```
attach(mtcars)
par(mfrow=c(2,2))
plot(wt,mpg, main="Scatterplot of wt vs. mpg")
plot(wt,disp, main="Scatterplot of wt vs disp")
 hist(wt, main="Histogram of wt")
boxplot(wt, main="Boxplot of wt")
```

# Graphical Parameters

- Three figures arranged in 3 rows and 1 column

```
attach(mtcars)
par(mfrow=c(3,1))
hist(wt)
hist(mpg)
hist(disp)
```

# Graphical Parameters

- The **layout( )** function has the form **layout(*mat*)** where *mat* is a matrix object specifying the location of the N figures to plot.
- # One figure in row 1 and two figures in row 2

```
attach(mtcars)
layout(matrix(c(1,1,2,3), 2, 2, byrow = TRUE))
hist(wt)
hist(mpg)
hist(disp)
```

# Graphical Parameters

- Optionally, you can include widths= and heights= options in the **layout( )** function to control the size of each figure more precisely. These options have the form
  **widths=** a vector of values for the widths of columns
  **heights=** a vector of values for the heights of rows.
- Relative widths are specified with numeric values. Absolute widths (in centimetres) are specified with the **lcm()** function.
- # One figure in row 1 and two figures in row 2
  # row 1 is 1/3 the height of row 2
  # column 2 is 1/4 the width of the column 1

```
attach(mtcars)
layout(matrix(c(1,1,2,3), 2, 2, byrow = TRUE),
  widths=c(3,1), heights=c(1,2))
hist(wt)
hist(mpg)
hist(disp)
```

# Thank You