

Az R adatelemzési nyelv alapjai II.

Egészségügyi informatika és biostatisztika

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Functions

Functions

- Functions do things with data
 - “Input”: function arguments (0,1,2,...)
 - “Output”: function result (exactly one)
- Why do we write functions?
 - To simplify code
 - To reduce code redundancy

- Example:

```
add <- function(a,b)
{ result <- a+b
  return(result) }
```

General Form of Functions

```
function (arguments)
{
    # do things...
    expression
}
```

```
function (arguments)
{
    # do things...
    return( expression )
}
```

Calling Conventions for Functions

- Arguments may be specified in the same order in which they occur in function definition, in which case the values are supplied in order.
- Arguments may be specified as name=value, when the order in which the arguments appear is irrelevant.
- Above two rules can be mixed.

```
> t.test(x1, y1, var.equal=F,  
conf.level=.99)
```

```
> t.test(var.equal=F, conf.level=.99,  
x1, y1)
```

Missing Arguments

- R function can handle missing arguments two ways:
 - either by providing a default expression in the argument list of definition, or
 - by testing explicitly for missing arguments.

Missing Arguments in Functions

```
> add <- function(x, y=0) {x + y}
```

```
> add(4)
```

```
> add <- function(x, y) {
```

```
  if(missing(y)) x
```

```
  else x+y
```

```
}
```

```
> add(4)
```

Variable Number of Arguments

- The special argument name “...” in the function definition will match any number of arguments in the call.

Variable Number of Arguments

```
> mean.of.all <- function(...) { mean(c(...)) }  
> mean.of.all(1:10, 20:100, 12:14)  
  
> mean.of.means <- function(...)  
{  
  means <- numeric()  
  for(x in list(...))  
    means <- c(means, mean(x))  
  mean(means)  
}  
> mean.of.means(1:10, 20:100, 12:14)
```

Formulas

Formulas

- Formulation of models
- Expression: $y \sim \text{model}$
 - the response y is modelled by a linear predictor specified symbolically by `model`
 - operators
 - $+$ \Rightarrow `term1+term2` (main effects)
 - $:$ \Rightarrow `term1:term2` (interaction)
 - $*$ \Rightarrow `term1*term2 =`
`term1+term2+term1:term2`
(main effects and interaction)
- ?formula

Graphics in R

plot()

- If `x` and `y` are vectors, `plot(x, y)` produces a scatterplot of `x` against `y`.
- `plot(x)` produces a time series plot if `x` is a numeric vector or time series object.
- `plot(mtcars)`
`plot(~ expr)`
`plot(y ~ expr)`, where `df` is a data frame, `y` is any object, `expr` is a list of object names separated by '+' (e.g. `a + b + c`).
- The first two forms produce distributional plots of the variables in a data frame (first form) or of a number of named objects (second form). The third form plots `y` against every object named in `expr`.

plot()

```
> attach( mtcars )
```

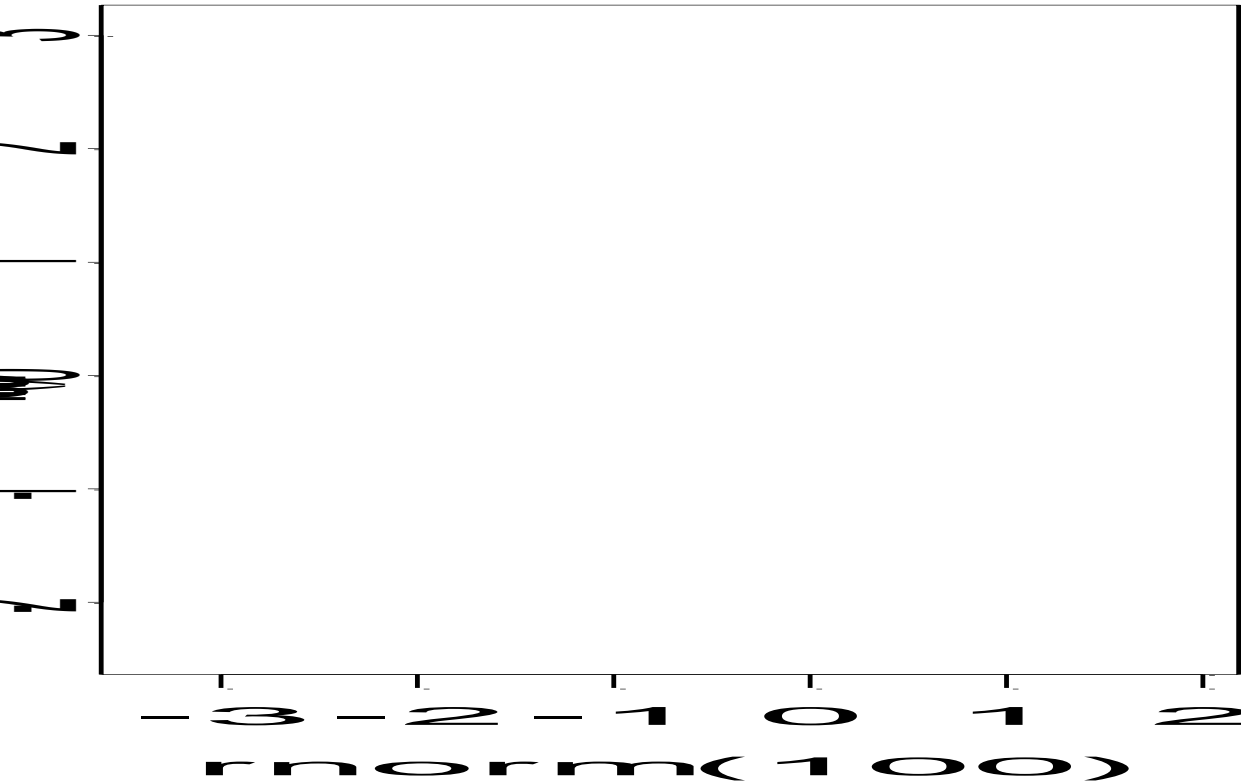
```
> plot( mtcars )
```

```
> plot( ~ cyl + hp + drat + wt )
```

```
> plot( mtcars, mpg ~ cyl + hp +  
drat + wt )
```

Graphics with `plot()`

```
> plot(rnorm(100), rnorm(100))
```



The function `rnorm()`
Simulates a random normal
distribution .

Help `?rnorm`,
and `?runif`,
`?rexp`,
`?binom`, ...

Graphics with `plot()`

```
> x <- seq(-2*pi, 2*pi, length=100)
```

```
> y <- sin(x)
```

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

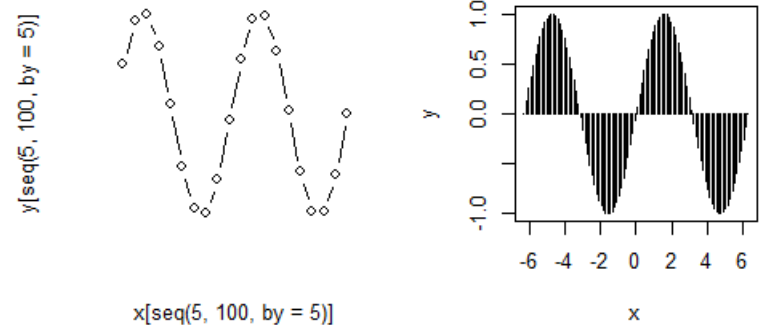
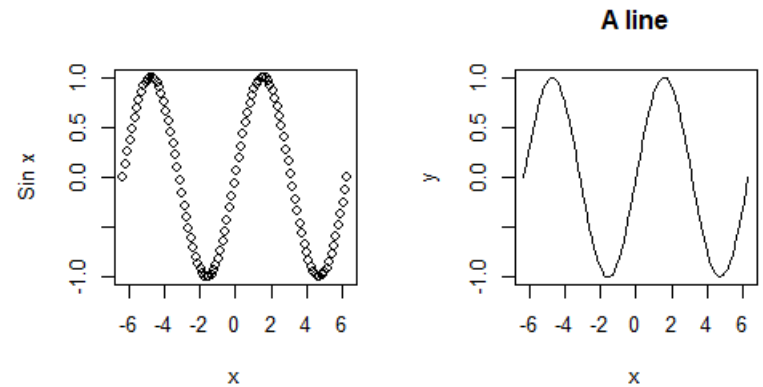
```
> plot(x, y, xlab="x",  
       ylab="Sin x")
```

```
> plot(x, y, type="l",  
       main="A Line")
```

```
> plot(x[seq(5, 100, by=5)],  
       y[seq(5, 100, by=5)],  
       type="b", axes=F)
```

```
> plot(x, y, type="h" )
```

```
> par(mfrow=c(1,1))
```



Graphical Parameters of `plot()`

```
type = "c": p (default), l,b,s,o,h,n  
pch= "+": character or numbers 1 - 18  
lty=1 : numbers #line type  
lwd=2 : numbers #line width  
xlab = "string", ylab = "string"  
sub = "string", main = "string"  
xlim = c(lo,hi), ylim = c(lo,hi)
```

And some more.

Graphical Parameters of `plot()`

```
x <- 1:10
y <- 2*x + rnorm(10,0,1)
plot(x,y,type="p") #Try l,b,s,o,h,n
# axes=T, F
# xlab="age", ylab="weight"
# sub="sub title",
# main="main title"
# xlim=c(0,12), ylim=c(-1,12)
```

Other graphical functions

See also:

`hist()`

`boxplot()`

`barplot()`

`image()`

`pairs()`

`persp()`

`piechart()`

`polygon()`

Other graphical functions

```
> axis(1, at=c(2, 4, 5),  
      legend("A", "B", "C"))
```

Axis details (“ticks”, legend, ...)
Use **xaxt="n"** ou **yaxt="n"** inside
plot()

```
> lines(x, y, ...)
```

Line plots

```
> abline(lsfite(x, y))
```

Add an adjustment

```
> abline(0, 1)
```

add a line of slope 1 and intercept 0

```
> legend(...)
```

Legends: very flexible

```
> text(x, y, "Hey")
```

Write text at coordinate x,y.

Histogram

- A *histogram* is a special kind of bar plot
- It allows you to visualize the *distribution* of values for a numerical variable
- When drawn with a *density scale*:
 - the *AREA* (NOT height) of each bar is the proportion of observations in the interval
 - the *TOTAL AREA* is 100% (or 1)

Making a histogram

- Type **?hist** to view the help file
 - Note some important arguments, esp **breaks**
- Simulate some data, make histograms varying the number of bars (also called 'bins' or 'cells'), e.g.

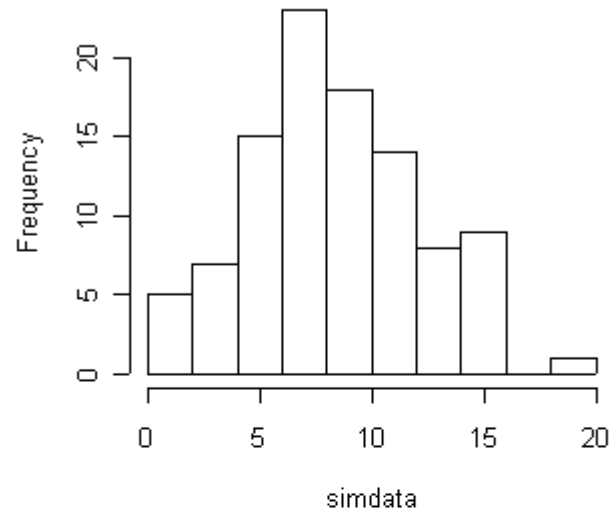
```
> par(mfrow=c(2,2)) # set up multiple  
plots
```

```
> simdata <- rchisq(100,8)
```

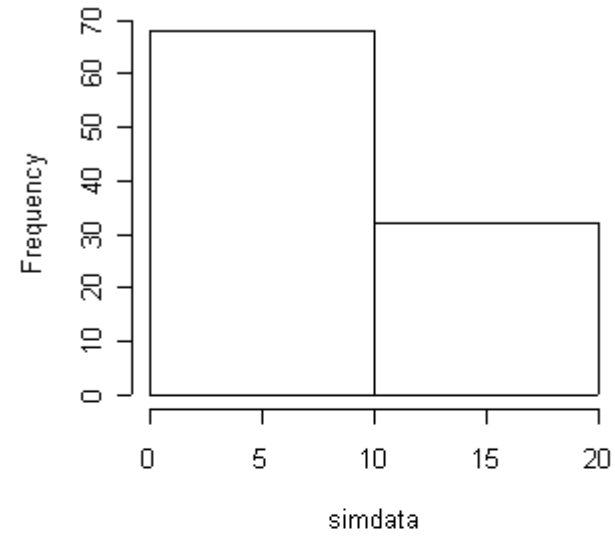
```
> hist(simdata) # default number of bins
```

```
> hist(simdata,breaks=2) # etc,4,20
```

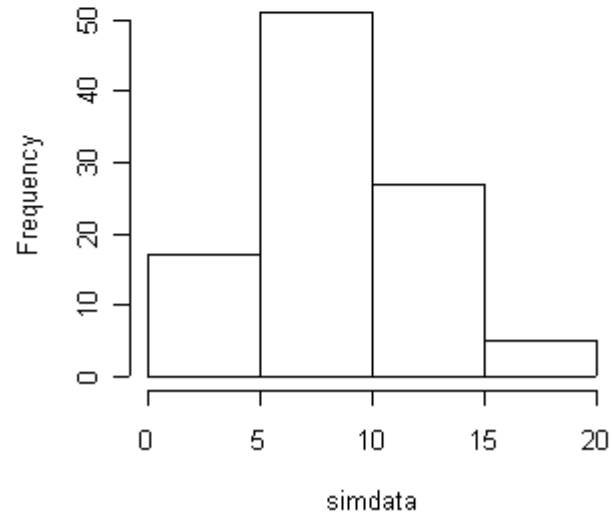
Histogram of simdata



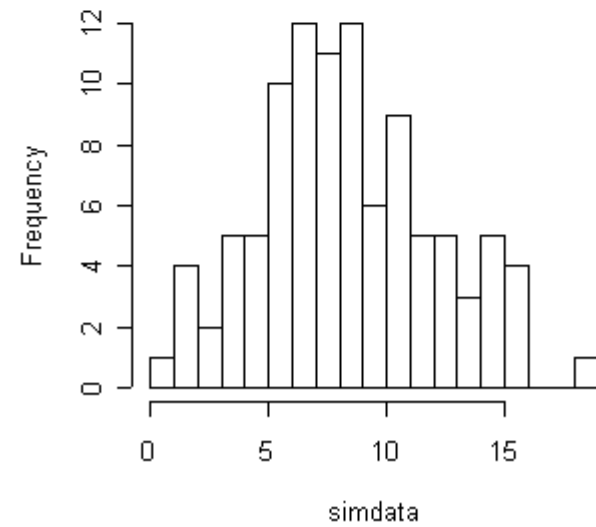
Histogram of simdata



Histogram of simdata



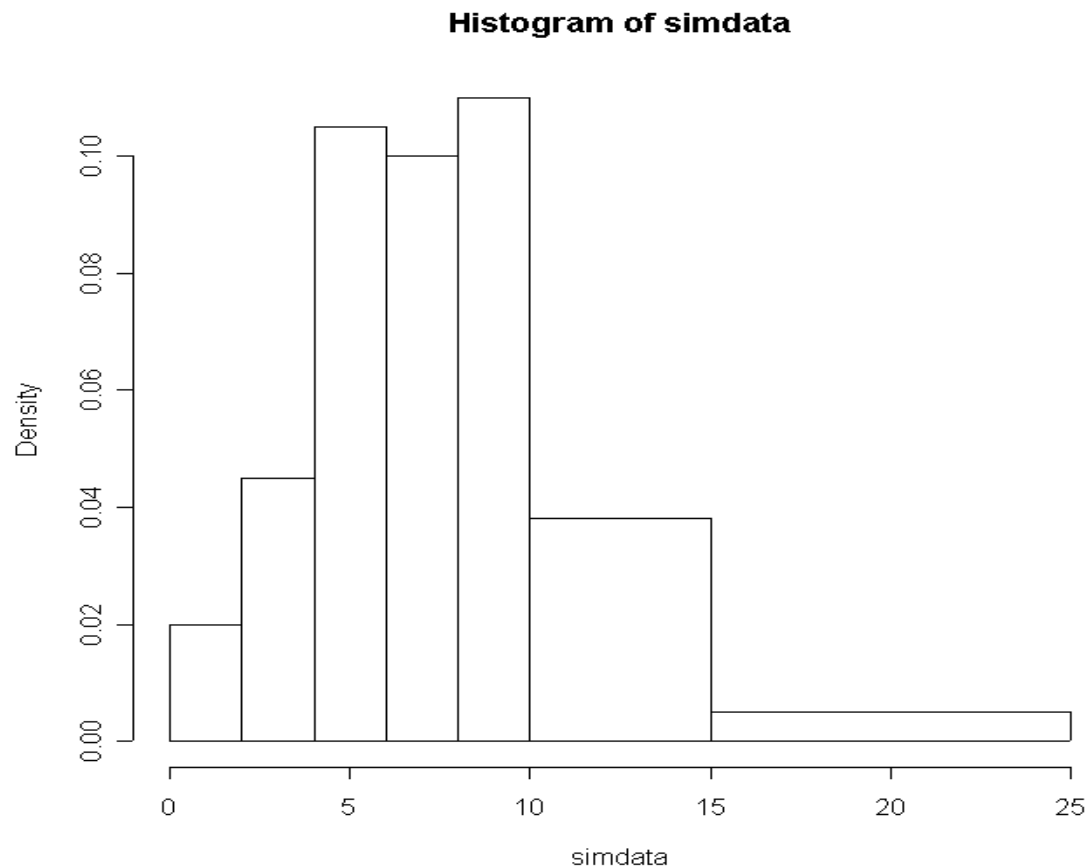
Histogram of simdata



Setting your own breakpoints

```
> bps <- c(0, 2, 4, 6, 8, 10, 15, 25)
```

```
> hist(simdata, breaks=bps)
```



Scatterplot

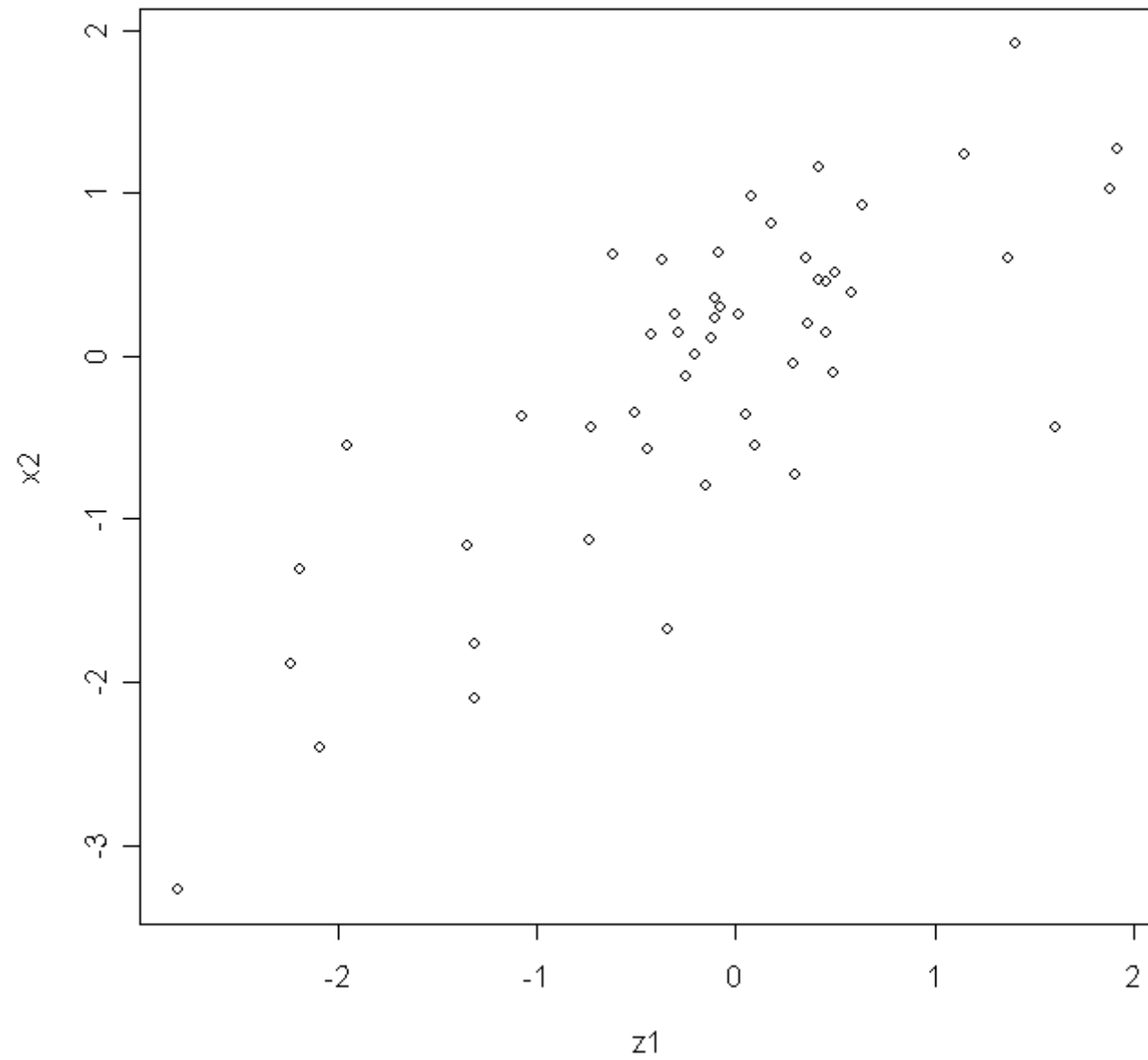
- A scatterplot is a standard two-dimensional (X,Y) plot
- Used to examine the relationship between two (continuous) variables
- It is often useful to plot values for a single variable against the order or time the values were obtained

Making a scatterplot

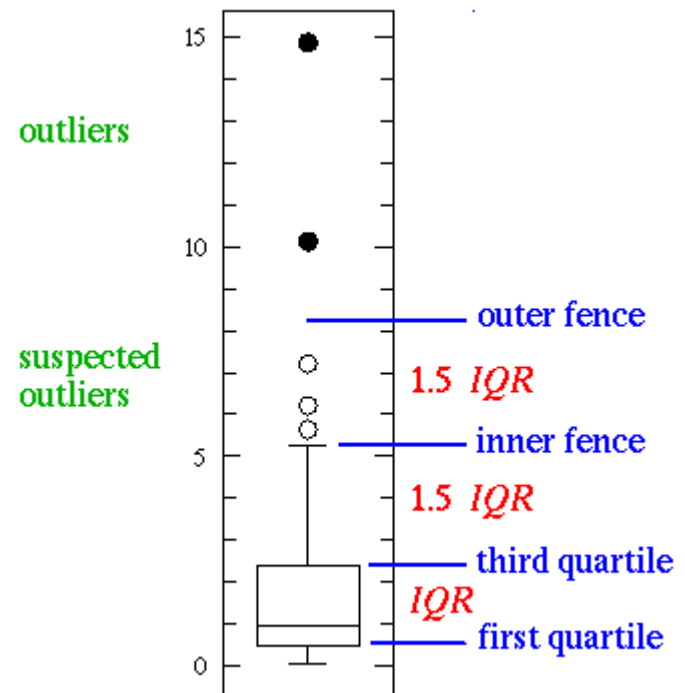
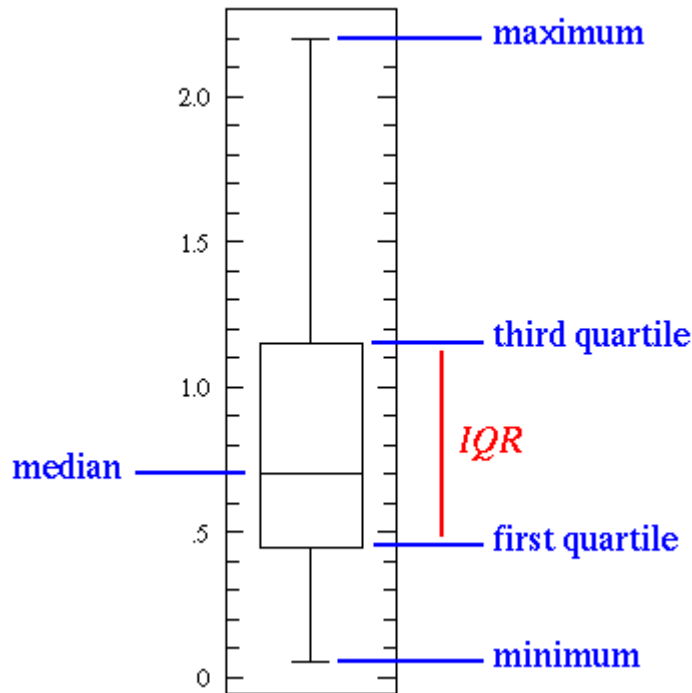
- Type `?plot` to view the help file
- Simulate a bivariate data set:

```
> z1 <- rnorm(50)
> z2 <- rnorm(50)
> rho <- .75          # (or any number between -1 and 1)
> x2 <- rho*z1+sqrt(1-rho^2)*z2
> plot(z1,x2)
```

Scatterplot of X2 vs. Z1



Boxplot (a.k.a. box and whisker diagram)



Making a boxplot

- Type `?boxplot` to view the help file
 - Split data by using formulas
- Simulate some data

```
> X <- c(rep(0,1000),rep(1,1000))
```

```
> Y <- X + rnorm(2000)
```

```
> simdata <- data.frame( X=X,Y=Y )
```

```
> boxplot( formula = Y ~ factor(X),  
           data = simdata )
```

