

R workshop

Action!

(Operators and Functions)

Acting on variables

- In the previous section, we talked about objects and data sets
- Now let's do something with them
- Verbs
 - Operators
 - Functions

Operator

- Simple calculation

Operator

+	addition
-	subtraction
*	multiplication
/	division
^	taking powers

Order of Operations

- Important note: Order of operations matters
 - PEMDAS is your friend

> $(8-4)/2$

[1] 2

> $8-(4/2)$

[1] 6

Example

- Take 14, add 4 and multiply the whole thing by 18.
- Take 14 and add the product of 4 and 18.

Example

- $> (14 + 4) * 18$
- [1] 324
- $> 14 + 4 * 18$
- [1] 86

Logical Operator

==	equality
!=	inequality
>	greater than
>=	greater than or equal to
<	less than
<=	less than or equal to

Logical Operator

- Returns a value of TRUE or FALSE

Example

- Test whether height is greater than 5.5

Logical Operators on Characters

- Height == 5.5
- Height = 5.5
- **WARNING:** a SINGLE equals sign will change your data!

```
install.packages("babynames")  
install.packages("ggplot2")
```

```
library(babynames)  
library(ggplot2)
```

```
MyName <- "Sara"  
birthday <- 1990  
MySex <- "F"
```

```
data("babynames")  
colnames(babynames)  
myName.df <- subset(babynames, name == MyName)
```

```
ggplot(myName.df, aes(x = year, y = prop, color=sex)) +  
  geom_line() +  
  geom_point(aes(x = birthday,  
                 y = myName.df[myName.df$name == MyName &  
                               myName.df$year == birthday &  
                               myName.df$sex == MySex, "prop"]),  
             color="black") +  
  ggtitle(paste("Popularity of", MyName))
```

Functions

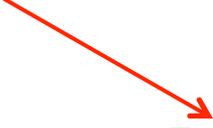
- Sometimes, you want to do more than add or multiply variables.
- To perform more complicated actions, use functions.
 - Functions are commands that describe, manipulate or analyze objects.

Functions

```
> log(10)  
[1] 2.302585
```

Functions

The **function** is called "log".



```
> log(10)
[1] 2.302585
```

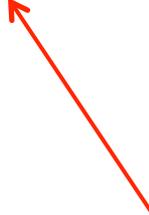
Functions

10 is an **object**

```
> log(10)
[1] 2.302585
```

Functions

```
> log(10)  
[1] 2.302585
```



2.302.... is the
output
returned by the
function

Mathematical functions

<code>sqrt()</code>	square root
<code>round()</code>	round a number
<code>log()</code>	logarithm
<code>exp()</code>	exponentiation
<code>abs()</code>	absolute value

Example

- Find the square root of 85.
- Take the log of 100.

Example

```
> sqrt(85)
[1] 9.219544
> log(100)
[1] 4.60517
```

```
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install.packages("ggplot2")
```

```
library(babynames)  
library(ggplot2)
```

```
MyName <- "Sara"  
birthday <- 1990  
MySex <- "F"
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```
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```
ggplot(myName.df, aes(x = year, y = prop, color=sex)) +  
  geom_line() +  
  geom_point(aes(x = birthday,  
                 y = myName.df[myName.df$name == MyName &  
                               myName.df$year == birthday &  
                               myName.df$sex == MySex, "prop"]),  
             color="black") +  
  ggtitle(paste("Popularity of", MyName))
```

Functions have three parts

- Function name
 - *Ex: log*
- Arguments
 - *Ex: 10*
- Output
 - *Ex: 2.302*

Functions have three parts

- Function name

- *Ex: log*

- Arguments

- *Ex: 10*

- Output

- *Ex: 2.302*

Each function has one and only one name.

Functions have three parts

- Function name

 - *Ex: log*

One argument is always specified: the input. This is the object that the function acts on.

- Arguments

 - *Ex: 10*

Other arguments control **how** the function acts. For example, do you want the natural log? Or log base 10?

- Output

 - *Ex: 2.302*

Each function has defaults for its arguments. You should know what those are and how to change them.

Functions have three parts

- Function name

 - *Ex: log*

- Arguments

 - *Ex: 10*

- Output

 - *Ex: 2.302*

Output can be a:

- number/integer

- a TRUE/FALSE statement

- a character value

- all of the above

Output can be a:

- single value

- vector

- data frame

- matrix

- list

You can store the output by assigning it to another object.

An Analogy

- Catch the ball quickly
- `Catch(item = ball, speed = "quickly")`

Exercise

- Use the `seq()` function to list numbers 0 to 100
- Arguments:
 - `from`: starting value of sequence
 - `to`: end value of sequence

Exercise

- Use the `seq()` function to list numbers 0 to 100, by intervals of 10
- Arguments:
 - `from`: starting value of sequence
 - `to`: end value of sequence
 - `by`: increment of the sequence

Exercise

- Use the `seq()` function to list numbers 0 to 100
- `>seq(from=0, to=100)`
- Now try to list numbers by intervals of 10
- `>seq(from=0, to=100, by=10)`

REMOVE POST-ITS!!



(and save!)

Multiple arguments

- Most functions take more than one argument.
- Separate arguments with commas.

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```
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[1] 2.305
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Number that
needs to be
rounded.

Multiple arguments

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- Separate arguments with commas.

```
> round (x = 2.30467, digits = 3)  
[1] 2.305
```



Number of
digits to round
to.

Arguments have Names

- Most arguments in functions have names.
- It is recommended that you use those names when using a function.

```
> round (x = 2.30467, digits = 3)  
[1] 2.305
```

Order of arguments

- Technically, you don't have to name the argument.
 - However, if you do not use the names, arguments MUST go in the right order.
 - Also you cannot skip arguments.
- If you name the arguments, you can put them in any order that you want and you can skip some.

Order of arguments

This is a
bad idea!



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 - However, if you do not use the names, arguments MUST go in the right order.
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- If you name the arguments, you can put them in any order that you want and you can skip some.

Order matters!

```
> round (2.30467, 3)
```

```
[1] 2.305
```

```
> round (3, 2.30467)
```

```
[1] 3
```

```
> round (x = 2.30467, digits = 3)
```

```
[1] 2.305
```

```
> round (digits = 3, x = 2.30467)
```

```
[1] 2.305
```

```
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            color="black") +  
  ggtitle(paste("Popularity of", MyName))
```

Great, but how do I know what the arguments are?

- Look in the R documentation
 - > `?round`
- In RStudio, hit tab to see names of arguments and descriptions.

The image shows a screenshot of the R Documentation viewer interface. At the top, there is a menu bar with 'Files', 'Plots', 'Packages', 'Help', and 'Viewer'. Below the menu bar is a toolbar with navigation icons (back, forward, home, print, refresh) and a search box. The main content area displays the title 'R: Rounding of Numbers' and a 'Find in Topic' search box. The page content includes the text 'Round {base}' and 'R Documentation' in the top right corner. The main heading is 'Rounding of Numbers'. Below this is a 'Description' section with paragraphs describing the functions `ceiling`, `floor`, `trunc`, `round`, and `signif`. Finally, there is a 'Usage' section listing the function signatures: `ceiling(x)`, `floor(x)`, and `trunc(x, ...)`.

Files Plots Packages Help Viewer

← → 🏠 🖨️ ↻ 🔍

R: Rounding of Numbers ▾ Find in Topic

Round {base} R Documentation

Rounding of Numbers

Description

`ceiling` takes a single numeric argument `x` and returns a numeric vector containing the smallest integers not less than the corresponding elements of `x`.

`floor` takes a single numeric argument `x` and returns a numeric vector containing the largest integers not greater than the corresponding elements of `x`.

`trunc` takes a single numeric argument `x` and returns a numeric vector containing the integers formed by truncating the values in `x` toward 0.

`round` rounds the values in its first argument to the specified number of decimal places (default 0).

`signif` rounds the values in its first argument to the specified number of significant digits.

Usage

```
ceiling(x)
floor(x)
trunc(x, ...)
```

You try!

- Look up documentation for the correlation function, `cor()`

You try!

> ?cor

Correlation, Variance and Covariance (Matrices)

Description

`var`, `cov` and `cor` compute the variance of `x` and the covariance or correlation of `x` and `y` if these are vectors. If `x` and `y` are matrices then the covariances (or correlations) between the columns of `x` and the columns of `y` are computed.

`cov2cor` scales a covariance matrix into the corresponding correlation matrix *efficiently*.

Usage

```
var(x, y = NULL, na.rm = FALSE, use)
```

```
cov(x, y = NULL, use = "everything",  
    method = c("pearson", "kendall", "spearman"))
```

```
cor(x, y = NULL, use = "everything",  
    method = c("pearson", "kendall", "spearman"))
```

```
cov2cor(V)
```

Arguments

- `x` a numeric vector, matrix or data frame.
- `y` `NULL` (default) or a vector, matrix or data frame with compatible dimensions to `x`. The default is equivalent to `y = x` (but more efficient).
- `na.rm` logical. Should missing values be removed?
- `use` an optional character string giving a method for computing covariances in the presence of missing values. This must be (an abbreviation of) one of the strings "everything", "all.obs", "complete.obs", "na.or.complete", or

cor {stats}

Correlation, Variance and Covariance (Matrices)

`cor {stats}`

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```
cov2cor(V)
```

```
cor {stats}
```

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cov2cor(V)
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`na.rm` logical. Should missing values be removed?

`use` an optional character string giving a method for computing covariances in the presence of missing values. This must be (an abbreviation of) one of the strings "everything", "all.obs", "complete.obs", "na.or.complete", or "pairwise.complete.obs".

`method` a character string indicating which correlation coefficient (or covariance) is to be computed. One of "pearson" (default), "kendall", or "spearman", can be abbreviated.

`V` symmetric numeric matrix, usually positive definite such as a covariance matrix.

Details

For `cov` and `cor` one must *either* give a matrix or data frame for `x` or give both `x` and `y`.

The inputs must be numeric (as determined by [is.numeric](#); logical values are also allowed for historical compatibility): the "kendall" and "spearman" methods make sense for ordered inputs but [xtfrm](#) can be used to find a suitable prior transformation to numbers.

Details

For `cov` and `cor` one must *either* give a matrix or data frame for `x` or give both `x` and `y`.

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Value

For `r <- cor(*, use = "all.obs")`, it is now guaranteed that `all(r <= 1)`.

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Value

For `r <- cor(*, use = "all.obs")`, it is now guaranteed that `all(r <= 1)`.

Examples

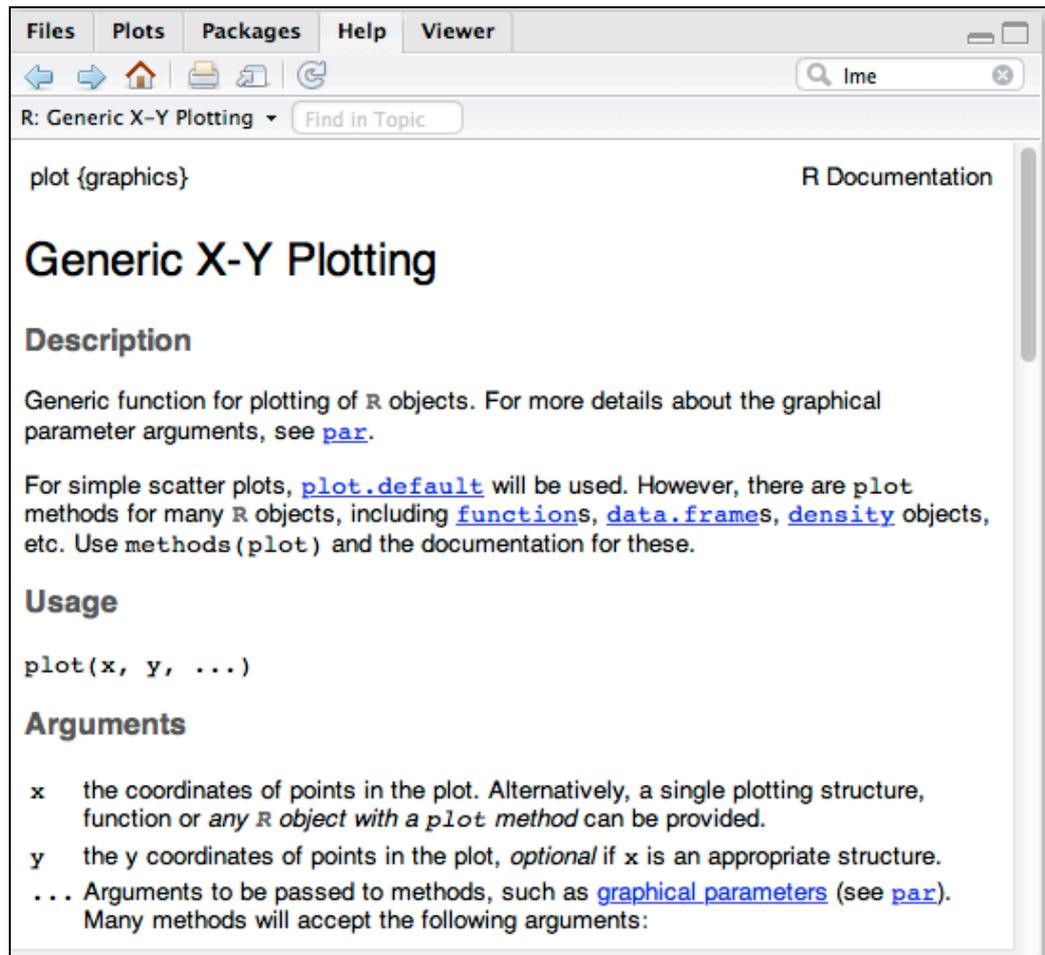
```
var(1:10) # 9.166667
var(1:5, 1:5) # 2.5
## Two simple vectors
cor(1:10, 2:11) # == 1
## Correlation Matrix of Multivariate sample:
(C1 <- cor(longley))
## Graphical Correlation Matrix:
symnum(C1) # highly correlated
```

Example

- 5 volunteers, how many cups of coffee did you drink today?
- Create a vector, add to data frame

Example

- Look up documentation for plot
- Make a scatter plot of coffee by height
- Add a title to your graph
- Add labels to x and y axes
- What happens if you add `type='l'`
 - What is the default for type?

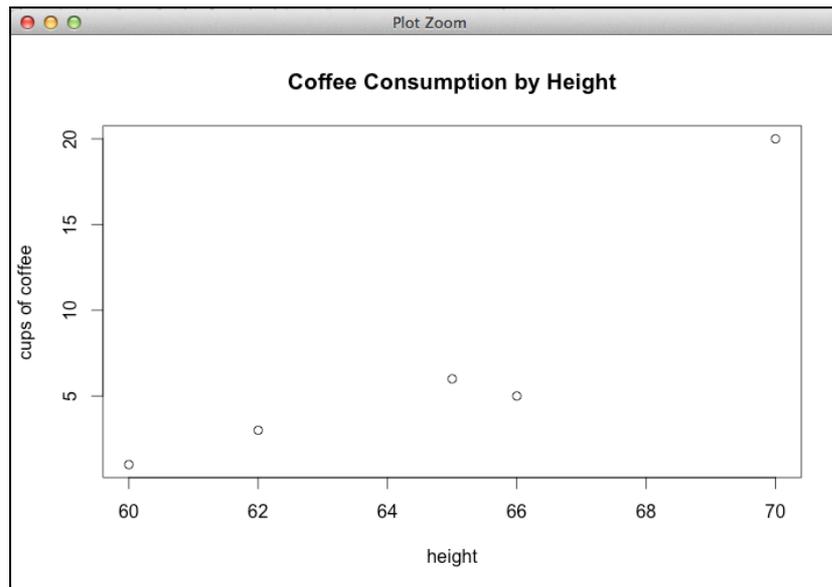


The image shows a screenshot of the R Documentation viewer. The window title is "R: Generic X-Y Plotting". The top menu bar includes "Files", "Plots", "Packages", "Help", and "Viewer". Below the menu bar is a search bar with the text "ime" and a search icon. The main content area displays the title "plot {graphics}" and "R Documentation". The main heading is "Generic X-Y Plotting". Below this is the "Description" section, which states: "Generic function for plotting of R objects. For more details about the graphical parameter arguments, see [par](#)." It also mentions: "For simple scatter plots, [plot.default](#) will be used. However, there are `plot` methods for many R objects, including [functions](#), [data.frames](#), [density](#) objects, etc. Use `methods(plot)` and the documentation for these." The "Usage" section shows the function signature: `plot(x, y, ...)`. The "Arguments" section lists:

- `x` the coordinates of points in the plot. Alternatively, a single plotting structure, function or *any R object with a plot method* can be provided.
- `y` the y coordinates of points in the plot, *optional* if `x` is an appropriate structure.
- `...` Arguments to be passed to methods, such as [graphical parameters](#) (see [par](#)). Many methods will accept the following arguments:

R Code

```
>plot(x=classdata$height,  
y=classdata$coffee,  
main="Coffee Consumption by Height",  
xlab="height (inches)",  
ylab="cups of coffee")
```



Back to the Documentation!

Arguments

- x** the coordinates of points in the plot. Alternatively, a single plotting structure, function or *any R object with a `plot` method* can be provided.
- y** the y coordinates of points in the plot, *optional* if **x** is an appropriate structure.
- ...** Arguments to be passed to methods, such as [graphical parameters](#) (see [par](#)). Many methods will accept the following arguments:

`type`

what type of plot should be drawn. Possible types are

- "p" for points,
- "l" for lines,
- "b" for both,
- "c" for the lines part alone of "b",
- "o" for both 'overplotted',
- "h" for 'histogram' like (or 'high-density') vertical lines,
- "s" for stair steps,
- "S" for other steps, see 'Details' below,
- "n" for no plotting.

All other types give a warning or an error; using, e.g., `type = "punkte"` being equivalent to `type = "p"` for S compatibility. Note that some methods, e.g. [plot.factor](#), do not accept this.