

# Bayesian Methods for Data Science (DATS 6450 - 11)

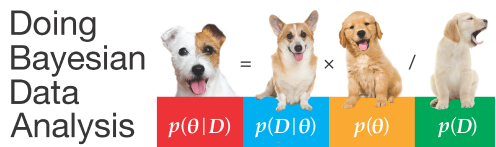
## The R Programming Language

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# Reference



Picture courtesy of the book website

- This set of slides is an excerpt of the book by Professor John K. Kruschke, with some trivial changes by the creator of the slides
- Please find the reference to and website of the book below:
  - Kruschke, J. K. (2014). *Doing Bayesian Data Analysis: A Tutorial with R, JAGS, and Stan. 2nd Edition.* Academic Press / Elsevier
  - <https://sites.google.com/site/doingbayesiandataanalysis/>

# Overview

- 1 Get the software
- 2 Basic commands and operators in R
- 3 Variable types
- 4 Loading and saving data
- 5 Some utility functions
- 6 Programming in R
- 7 Graphical plots: opening and saving

# Get R and RStudio

- Go to: <http://cran.r-project.org/> (Google “R language” if broken link)
- Download R for your operating system
- Go to: <http://www.rstudio.com/> (Google “RStudio” if broken link)
- Download RStudio for your operating system

## Example: plot $y = x^2$

- R:
  - see SimpleGraph.R (available on the book website)
  - explain:
    - Seq
    - from, to, by
    - type = "l"
- RStudio:
  - Run: executes the line of code where the cursor currently resides, or the lines of code currently selected<sup>1</sup>
  - Source: executes the entire document<sup>1</sup>

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<sup>1</sup><https://support.rstudio.com/hc/en-us/articles/200484448-Editing-and-Executing-Code>

# Learn from examples

- Reading from (good) examples can get you start quickly
- One such example is ExampleOfR.R (available on the book website)
- Read the code and comment to see how things are done in R
- Tweak the code and see if the result is the same as what you expected

## Other resources

- *An Introduction to R*, available at this address:  
<http://cran.r-project.org/doc/manuals/r-release/R-intro.pdf>
- *R Reference Card*, available at this address:  
<https://cran.r-project.org/doc/contrib/Short-refcard.pdf>

# Getting help in R

- `help.start()`: returns a list of online documentation, including *An Introduction to R*
- `help(name)` or `?name`: returns a single help page explaining the topic with name *name*
- `??name`: returns a list of help pages containing the word *name*
- Google it!



# Arithmetic operators

- Arithmetic operators:

- $+$
- $-$
- $*$
- $/$
- $^$ : power

- Precedence

- from left to right
- $^$  higher than  $*$ ,  $/$  higher than  $+$ ,  $-$
- what is  $1 + 2 * 3^2$  ?

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- what is  $1 + 2 * 3^2$  ?
- use parentheses to explicitly express the order that you intended
- $1 + 2 * (3^2)$

# Logic operators

- Logic values:
  - TRUE
  - FALSE
- Logic operators:
  - `!`: negation
  - `&&`: logic and
  - `||`: logic or
- Precedence
  - from left to right
  - `!` higher than `&&` higher than `||`
  - again, use parentheses to explicitly express the order that you intended

# Relational operators and tests of equality

- Relational operators:
  - $<$
  - $>$
  - $!=$
  - $==$
  - `all.equal()`
- Tests of equality:
  - $==$ 
    - what is  $0.5 - 0.3 == 0.3 - 0.1$ ?

# Relational operators and tests of equality

- Relational operators:

- $<$
- $>$
- $!=$
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- `all.equal()`

- Tests of equality:

- $==$

- what is  $0.5 - 0.3 == 0.3 - 0.1$ ?

- FALSE! (due to the limited precision of representing numbers in the computer's memory)

- `all.equal()`

- equal up to precision of computer

- `all.equal(0.5 - 0.3, 0.3 - 0.1)`: TRUE

- use `isTRUE(all.equal())` in *if* expressions<sup>1</sup>

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<sup>1</sup>R Documentation

# Assignment operators

- `<-`
  - if there is an empty space between `<` and `-`, “`< -`”
  - then this is a logic operator followed by a negative sign
  - `x < - 1`: TRUE when `x` is smaller than -1, FALSE otherwise
- `=`
  - preferred
  - differentiate `=` with `==`

# Vector

- A vector is an ordered (not sorted) list of elements of the same type
- Some useful functions
  - the combine function
  - component-by-component vector operations
  - the colon operator and sequence function
  - the replicate function
  - getting at elements of a vector
- See `ExampleOfR.R` for details (available on the book website)

# Factor

- A factor is a vector of indices (transformed from a vector of categorical values), along with a legend that decodes each index into a level name
- Some useful functions
  - the factor function
  - reorder the levels
  - labels
- See `ExampleOfR.R` for details (available on the book website)



# Matrix and array

- A matrix is simply a two-dimensional array of values of the same type
- An array is a generalization of a matrix to multiple dimensions
- See `ExampleOfR.R` for details (available on the book website)

# List and data frame

- The list structure is a generic vector in which components can be of different types, and named
- A data frame is a type of list in which each component is thought of as a named column of a matrix, with different columns possibly of different types
- See `ExampleOfR.R` for details (available on the book website)

# The read.csv function

- The read.csv function loads comma separated values (a.k.a. CSV) files into R's memory
- The data in the CSV files are stored as a data frame
- The categorial columns in the CSV files are transformed into factors in the data frame
- Some useful functions
  - re-order the levels
  - as.vector()
  - factor()
- See ExampleOfR.R for details (available on the book website)

## Saving data from R

- The `write.csv()` saves a data frame to a CSV file
- The CSV file loses all information about levels in factors in the data frame
- The `save()` function saves a data frame with all the factor information to a Rdata file
- The `load()` function loads a Rdata file into R's working memory
- The `objects()` function shows the objects in R's working memory
- See `ExampleOfR.R` for details (available on the book website)

# Some utility functions

- `summary()`: returns a summary appropriate for the argument
- `aggregate()`: summarizes data according to factor characteristics
- `apply()`: collapses arrays across specified dimensions, and applies a function to the data within the collapsed dimensions
- `melt()`: rearranges data so that there is one datum per row; needs the `reshape2` package
- See `ExampleOfR.R` for details (available on the book website)

# Variable names in R

- Variable names should be meaningful
- Variable names are case sensitive
- No space in the name of a variable
- Use *camelBack* notation for naming convention
  - the first letter of each word (except for the first one) is capitalized
  - e.g., `bayesianMethodsForDataScience`

# Running a program

- Set the working directory
- The `source()` function
  - accept all the commands from the argument
  - different from the `load()` function, which reads a Rdata format file
- The Run button: runs the line on which the cursor is presently positioned, or multiple lines if they are presently selected, and echoes the lines in the command window
- The Source button: runs the entire program without echoing the lines in the command window

# Programming a function

- Arguments with explicit labels may go in any order
- Arguments without explicit labels must be in the order used in the definition of the function
- Arguments with default value do not have to be specified in the function call
- Arguments without default value must be specified
- Unlabeled argument provided in the function call is assigned to the first argument in the definition, regardless of whether the argument was defined with a default value
- See `ExampleOfR.R` for details (available on the book website)



# Conditions and loops

- The *if-else* expression: the line containing “else” begins with a closing curly brace; a line that begins with “else” causes an error
- The *for* loop
- See ExampleOfR.R for details (available on the book website)

# Measuring processing time

- The `proc.time()` function: returns the current computer-system time
- To measure the duration of a process, use `proc.time()` at the beginning and end of the process, and compute the difference
- The ‘user time’ is the CPU time charged for the execution of user instructions of the calling process<sup>1</sup>
- The ‘system time’ is the CPU time charged for execution by the system on behalf of the calling process<sup>1</sup>
- See `ExampleOfR.R` for details (available on the book website)

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<sup>1</sup>R Documentation

# Debugging

- Hints for avoiding producing errors
  - when creating a new program, always start with a new file name
  - use explicit parentheses to be sure that operators are applied in the intended order
  - use visual white space to make your code easily readable
  - use indenting to meaningfully group sections of the program
    - select a section of program or the entire program
    - press ctrl-i to have the program properly indented
- Hints for debugging the errors
  - read the error messages displayed by R
  - locate the first point in the code where the error occurs by printing the result of the program section by section
  - see details of debugging in RStudio at:  
<http://www.rstudio.com/ide/docs/debugging/overview>

# The openGraph and saveGraph function

- The openGraph and saveGraph function are used for opening and saving graphs
- Need to install other packages
- See ExampleOfR.R and DBDA2E-utilities.R for details (available on the book website)