# Introduction to Biostatistical Computing Literate Programming

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# Introduction

An article about computational science in a scientific publication is not the scholarship itself, it is merely advertising of the scholarship. The actual scholarship is the complete software development environment and the complete set of instructions which generated the figures.

- D. Donoho
  - The concept of reproducible research is based on the idea of literate programming such that the logic of the analysis is clearly represented in the final product by combining computer code/programs with ordinary human language
    - ⇒ Combine analysis code and report

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# Literate Programming with R

#### 2 main tools

- **Sweave** (2002): a tool that allows to embed the R code for complete data analyses in latex documents. The purpose is to create dynamic reports, which can be updated automatically if data or analysis change.
  - Sweave is part of every R installation
- knitr: "The knitr package was designed to be a transparent engine for dynamic report generation with R, solve some long-standing problems in Sweave, and combine features in other add-on packages into one package"
  - e.g., caching, html and markdown

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# Data Example

Televisions, Physicians, and Life Expectancy

The data set contains information on

- Life expectancies in 40 countries with population ≥ 20 millions (1990)
  - For female and male. Average used as the country's overall life expectancy
- Number of people per television set
- Number of people per physician

Data set taken from the The World Almanac and Book of Facts, 1993

# Data Example

Televisions, Physicians, and Life Expectancy

```
Country life tv phys fem male
   Argentina 70.5 4.0 370
                            74
                                  67
 Bangladesh 53.5 315.0 6166
                             53
                                  54
3
     Brazil 65.0
                   4.0
                        684
                             68
                                  62
4
     Canada 76.5
                   1.7
                        449
                             80
                                  73
5
                        643
                             72
                                  68
      China 70.0 8.0
6
   Colombia 71.0
                   5.6 1551
                             74
                                  68
```

# Data Example

Televisions, Physicians, and Life Expectancy

Is the number of people per TV set a good predictor of life expectancy?

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- A .tex file with .Rnw extension
- Include R code as 'chunks' or inline

- .Rnw file converted to .tex using Sweave
- The .tex file contains output from R; no raw code
- .tex file converted to .pdf (or .ps, .dvi)

```
.Rnw file
```

```
\documentclass{article}
\usepackage{Sweave}
\begin{document}
Some R code:
<<>>=
x <- rnorm(1000)
Let's display an histogram of {\tt x}
\begin{figure}[!h]
  \centering
<<myhist, fig = TRUE, eps = FALSE>>=
hist(x)
\caption{An histogram of {\tt x}}
   \label{fig:hist}
\end{figure}
\end{document}
```

#### .tex file

\end{document}

```
\documentclass{article}
\usepackage{Sweave}
\begin{document}
Some R code:
\begin{Schunk}
\begin{Sinput}
> x <- rnorm(1000)
\end{Sinput}
\end{Schunk}
Let's display an histogram of {\tt x}
\begin{figure}[!h]
 \centering
\begin{Schunk}
\begin{Sinput}
> hist(x)
\end{Sinput}
\end{Schunk}
\includegraphics{simple_example-myhist}
\caption{An histogram of {\tt x}}
  \label{fig:hist}
\end{figure}
```

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Some R code:  $> x \leftarrow rnorm(1000)$  Let's display an histogram of x

> hist(x)

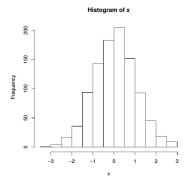


Figure 1: An histogram of x

#### Code chunks

R code is entered in the LTFX document using code chunks.

- Text within the code chunk is interpreted as R code
- Arguments for the code chunk are entered within <<here>>=

#### Code chunks options

- eval: Default is TRUE. If set to FALSE, code is not evaluated
- echo: Include R code in the output file. Default is TRUE
- keep.source: Default to FALSE. If TRUE, the original source code is copied to the file when echoing. Otherwise it gets through R's parsing engine
- results: Default to verbatim, the output of the R code is copied "verbatim" to the file. Other option is tex for latex output, e.g., table, or hide
- include: Default to TRUE. If you want to place the figure yourself, add include=FALSE

#### Code chunks options

- fig: Default to FALSE. Indicate whether the code chunk produces a graphical output
- eps: Indicates whether a .eps figure should be generated. Default is TRUE but ignored if fig is FALSE
- pdf: generate .pdf figure. Default to TRUE
- width and height: width and height of the figures in inches (default is
   6)

```
\SweaveOpts{option1 = value1, option2 = value2, ...}
```

in the preamble sets default for all the code chunks in the document

D 1 4 3 1 4

#### Named code chunks

Named code chunks can be reused in other code chunks later in the document

```
<<a>>>=
```

(

$$x + y$$

0

ര

#### **Figures**

Easy to include figures in your document

```
\begin{figure}
\begin{center}
<<my_fig, fig=TRUE, echo=FALSE, width=10, eps = FALSE>>=
plot(x, y)
@
\caption{A beautiful figure}
\end{center}
\end{figure}
```

The chunk name is used as figure name, i.e., myfile-myfig.pdf

#### **Figures**

#### Alternative:

```
<<my_fig, fig=TRUE, echo=FALSE, width=10, include = FALSE>>=
plot(x, y)
@
\begin{figure}
\begin{center}
\includegraphics{myfile-myfig.pdf}
\caption{A beautiful figure}
\end{center}
\end{figure}
```

#### **Tables**

results option should be set to tex

Necessitates R packages that convert R object into LTFX

- xtable: The most general
- stargazer, texreg, ..., provides LTEX representations for several models (e.g., linear models, mixed models)

See http://cran.r-project.org/web/views/ReproducibleResearch.html for a relatively exhaustive list

#### xtable

```
<<results=tex, echo=FALSE>>=
library(xtable)
y <- rnorm(100)
x <- rnorm(100)
fit.lm <- lm(y ~ x)
xtable(summary(fit.lm)$coefficients)
@</pre>
```

#### xtable

```
<<results=tex, echo=FALSE>>=
library(xtable)
y <- rnorm(100)
x <- rnorm(100)
fit.lm <- lm(y ~ x)
xtable(summary(fit.lm)$coefficients)
@</pre>
```

```
\begin{table}[ht]
\centering
\begin{tabular}{rrrrr}
\hline
& Estimate & Std. Error & t value & Pr($>$$|$t$|$) \\
\hline
(Intercept) & -0.037 & 0.090 & -0.418 & 0.677 \\
        x & -0.020 & 0.092 & -0.217 & 0.829 \\
\hline
\end{tabular}
\end{tabular}
```

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#### xtable

## Useful options in xtable()

- caption
- label: For cross-reference in LTEX
- align: alignment of the columns
- digits: number of digits to display

## Useful options in print.xtable()

- floating.environment: control the floating environment. Default is table
- table.placement: Control placement of the table in the pdf. Default is ht.
- include.rownames and include.colnames

See ?xtable and ?print.xtable for all the options



#### **Expressions**

All objects within a code chunk is in R's global environment each time a document is compiled

This allows the information saved in the global environment to be reproduced in the final document as inline text via *expressions* 

```
<<echo = FALSE>>=
x <- rnorm(100)
@
```

The mean of x is  $\operatorname{Sexpr}\{\operatorname{round}(\operatorname{mean}(x), 2)\}$ 

#### **Expressions**

All objects within a code chunk is in R's global environment each time a document is compiled

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```
<<echo = FALSE>>=
x <- rnorm(100)
@
```

```
The mean of x is \operatorname{Sexpr}\{\operatorname{round}(\operatorname{mean}(x), 2)\}
```

The mean of x is 0.08



#### Run Sweave

#### In Rstudio

- Set up an option that decides on using Sweave or knitr
- Click on "compile pdf"

#### From within R

- Sweave("foo.Rnw") produces the .tex file
- texi2dvi("foo.tex", pdf = TRUE) for compiling the .tex file within R

#### In the shell

- R CMD Sweave foo.Rnw produces the .tex file
- pdflatex foo.tex
- pdflatex foo.tex produces the pdf (done 2 times to actualise references, etc.)

## In emacs (+ESS)

• M-n s for Sweave, M-n P for pdflatex



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# knitr essentially reimplements and extends Sweave

- Support for additional document types: Markdown, html, reStructuredText
- Improved chunk options
- Code decoration
- Support multiple graphics devices
- Better plot manipulation capabilities
- Cacheing
- Different error handling

knitr not included in R, so you might have to install it

```
install.packages("knitr")
```

and load it in R

library("knitr")

To process a report document from within R

knit("my\_report.Rnw")

knitr is well integrated within Rstudio

## **Chunk options**

For .Rnw files, it works mostly as with Sweave

```
<<some_code, eval=TRUE, results="asis">>=
```

- As in Sweave, chunk options should be written in one line
- Option values must be valid R expressions (unlike in Sweave)

#### **Chunk Options**

- eval
- echo
- results character that takes 3 possible values
  - 'markup' mark up the results using the output hook, e.g. put results in a special LaTeX environment
  - 'asis' output as-is, i.e., raw output of R
  - 'hide'
- highlight whether to highlight the source code
- dev The graphical device to record plots on, e.g., 'pdf' for latex, 'png' for html
- cache caching?

Comprehensive list of options yihui.name/knitr/options



## **Chunk Options**

Unlike Sweave, global options for knitr are set in R

```
<<setup, include=FALSE>>=
opts_chunk$set(fig.width=5, fig.height=5)
@
```

This example will constrain the dimensions of all figures to  $5\times5$  inches

# Inline code

As in Sweave using \Sexpr{}

## **Figures**

- No need of an extra option in the chunk header to include a figure. In Sweave, fig=true was needed
- (at least for latex) no need to wrap the R chunk around \begin{figure} \end{figure} if fig.cap='caption' is specified
- It is possible to include multiple graphs within one chunk. Figures will be automatically put side-by-side with one caption as indicated by fig.cap
- Choice of the graphical device through the dev argument
- You can automatically put all automatically generated figures into a separate directory using the fig.path chunk option, e.g.,

<<fig.path="Graphics", caption="Some text">>=

## Figures

# Control of the plot size made easier

- fig.height and fig.width control the size of the plot made by the plotting device. Defaults are 7 inches
- out.width and out.height control the size of the plot in the output document. For Lag.,

```
out.width="10cm"
out.width=".7\\linewidth"
```

# knitr with HTML

html is the the standard markup language used to create web pages HTML is written in the form of HTML elements consisting of tags enclosed in angle brackets (like <html>). HTML tags most commonly come in pairs like <h1> and </h1>

# knitr with HTML

## Code chunks

```
<!--begin.rcode echo=FALSE, results='asis'
x <- rnorm(100)
end.rcode-->
```

# knitr with HTML

#### Code chunks

```
<!--begin.rcode echo=FALSE, results='asis'
x <- rnorm(100)
end.rcode-->
```

- In my experience, medical doctors don't like pdf
- An HTML file can be opened in Word

# knitr with Markdown

Markdown is a plain text formatting syntax[5] designed so that it can optionally be converted to HTML using a tool by the same name

```
Heading
======
Sub-heading
Paragraphs are separated
by a blank line.
Text attributes *italic*.
**bold**, 'monospace'.
A [link] (http://example.com).
Shopping list:
  * apples
  * oranges
```

\* pears

# knitr with Markdown

# Heading

## Sub-heading

Paragraphs are separated by a blank line.

Text attributes italic, bold, monospace.

#### A link .

Shopping list:

- apples
- oranges
- pears

Numbered list:

- 1. apples
- 2. oranges
- pears

The rain—not the reign—in Spain.

# knitr with Markdown

## Code chunk

```
'''{r, echo=FALSE}
x <- rnorm(100)
'''</pre>
```