

# Creating basic R workflows and literate programming

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26 November 2024

# Outline

- Project-oriented workflows in R
  - Why organise workflows into projects
  - How to organise workflows into projects
- Literate programming
  - Why use literate programming
  - RMarkdown for literate programming
- Practical session

# Project-oriented workflows in R

# Why organise workflows into projects

- key word is "*organise*";
- organisation ensures *coherence* and *order* which is helpful to the user whenever he/she goes back to the same work again later;
- organisation supports good documentation; and,
- organisation allows for collaboration.

# How to organise workflows into projects

- a unique piece of "*work*" should be its own *project*;
- all resources and tools needed for a *project* should be within the same directory; and,
- group resources and tools within appropriately into their own directories with the project directory.

OxfordIHTM / Awareness-Driven-Behaviour (Private)

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main 1 branch 0 tags Go to file Add file <> Code

|                      |                       |                        |           |
|----------------------|-----------------------|------------------------|-----------|
| grecomalijan         | Delete Model Code.Rmd | 2e57851 on Aug 8, 2021 | 2 commits |
| App                  | Add files via upload  |                        | last year |
| Model_Comparison.csv | Add files via upload  |                        | last year |
| PHSI.csv             | Add files via upload  |                        | last year |
| PH_NPI+STA+LTA.R     | Add files via upload  |                        | last year |
| PHcases.csv          | Add files via upload  |                        | last year |
| PHdeaths.csv         | Add files via upload  |                        | last year |
| README.md            | Add files via upload  |                        | last year |
| Sensi.csv            | Add files via upload  |                        | last year |

OxfordIHTM / sc-vaccine-study-draft (Public)

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|                                 |                |                        |            |
|---------------------------------|----------------|------------------------|------------|
| gsrajahram                      | initial commit | 3affb15 on Dec 9, 2022 | 1 commit   |
| data                            | initial commit |                        | last month |
| output                          | initial commit |                        | last month |
| .gitignore                      | initial commit |                        | last month |
| sc-vaccine-study-draft.Rproj    | initial commit |                        | last month |
| sc_vaccines_study_non_targets.R | initial commit |                        | last month |

Help people interested in this repository understand your project by adding a README. Add a README

# Literate programming

# Why use literate programming

- write code that satisfies both the machine/computer and the human reader/user;
- highlight and give as much importance to the documentation of the processes and the outputs that your code represents;
- focus on communication and understanding of the process and the output; and,
- single document that integrates both code and textual documentation.

# R Markdown for literate programming

- R Markdown provides an authoring framework for literate programming;
- A single R Markdown file is used to both save and execute code, and generate high quality reports that can be shared with an audience; and,
- R Markdown documents are fully reproducible and support dozens of static and dynamic output formats.

# R Markdown example - PDF report

```
2189 lines (1748 sloc) | 142 KB
1  ---
2  #title: Report on Coverage Assessment of Direct Nutrition Interventions in Liberia
3  #author: "Valid International"
4  #date: `r format(Sys.Date(), "%d %B %Y")`'
5  fontsize: 12pt
6  geometry: margin=2cm
7  documentclass: article
8  classoption: a4paper
9  bibliography: bibliography.bib
10 lot: TRUE
11 lof: TRUE
12 link-citations: TRUE
13 links-as-notes: FALSE
14 colorlinks: TRUE
15 linkcolor: blue
16 citecolor: blue
17 urlcolor: blue
18 ...
19
20 ````{r setup, include = FALSE}
21 knitr::opts_chunk$set(echo = TRUE,
22                       warning = FALSE,
23                       message = FALSE)
24
25 if(!require(stringr)) install.packages("stringr")
26 if(!require(raster)) install.packages("raster")
27 if(!require(bbw)) install.packages("bbw")
28 if(!require(tidyR)) install.packages("tidyR")
29 if(!require(ggplot2)) install.packages("ggplot2")
30 if(!require(knitr)) install.packages("knitr")
31 if(!require(kableExtra)) install.packages("kableExtra")
32 if(!require(RColorBrewer)) install.packages("RColorBrewer")
33 if(!require(rgdal)) install.packages("rgdal")
34 if(!require(cowplot)) install.packages("cowplot")
35
36 if(!require(remote)) install.packages("remote")
37 if(!require(liberia)) install_github("validmeasures/liberia")
38 if(!require(liberiaData)) install_github("validmeasures/liberiaData")
39
40 options(stringsAsFactors = FALSE)
41 options(kableExtra.latex.load_packages = FALSE)
42
43 themeSettings <- theme_bw() +
44 theme(
```



THE POWER OF  
nutritio<sup>n</sup> **VALID**

Report on Coverage Assessment of Direct Nutrition Interventions in Liberia

# R Markdown example - HTML report

505 lines (440 sloc) | 9.3 KB

```
1  ---
2  title: Lifetables Seychelles 2021
3  author: Ministry of Health
4  date: 2022-06-16
5  output: html_document
6  ---
7
8  ``{r setup, include=FALSE}
9  knitr::opts_chunk$set(echo = FALSE)
10
11 ## Load packages and functions
12 suppressPackageStartupMessages(source("packages.R"))
13 for (f in list.files(here::here("R"), full.names = TRUE)) source (f)
14
15 ## Load required targets
16 tar_load(
17   c(
18     sc_lifetable, smooth_sc_lifetable,
19     demogdat, smooth_demogdat,
20     #lca_male, predict_lca_male,
21     #lca_female, predict_lca_female,
22     lca_total, predict_lca_total,
23     lca_smooth_total, predict_lca_smooth_total
24   )
25 )
26 ...
27
28 ## Lifetable
29
30 2021 life table samples, with forecast of mortality rates for next 30 years (based on 2011-2021 lifetable data, using lee-carter approach)
31
32 ### Lifetable by sex - 2021
33
34 ``{r lifetable-2021, fig.align = "center", fig.width = 12, fig.height = 8}
35 par(mfrow = c(1, 2))
36
37 plot(
38   sc_lifetable,
39   series = "male",
40   datatype = "rate",
41   transform = FALSE,
42   main = "Male death rates by age - 2021",
43   lty = 0,
44   ylim = c(0, max(sc_lifetable$rate$male, sc_lifetable$rate$female)),
```

## Lifetables Seychelles 2021

Ministry of Health

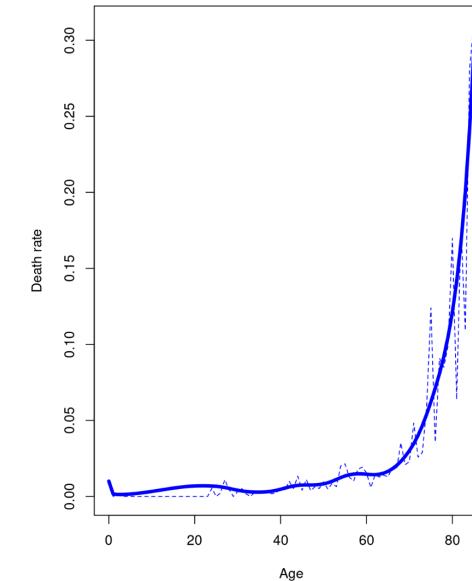
2022-06-16

### Lifetable

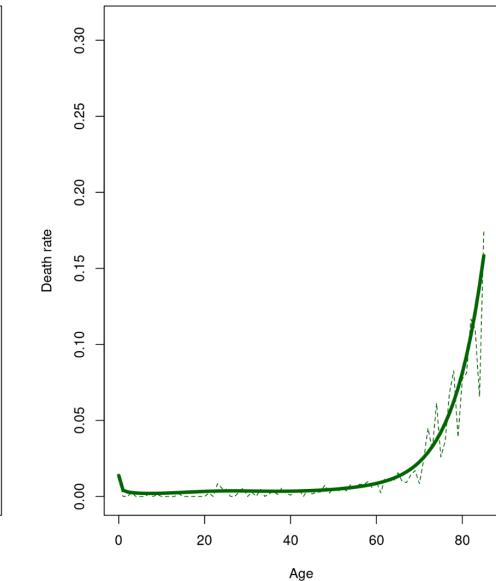
2021 life table samples, with forecast of mortality rates for next 30 years (based on 2011-2021 lifetable data, using lee-carter approach)

### Lifetable by sex - 2021

Male death rates by age - 2021



Female death rates by age - 2021



Male death rates by age - 2021

Female death rates by age - 2021

# R Markdown example - Website

295 lines (156 sloc) | 18.5 KB

```
1 ---  
2 output: github_document  
3 ---  
4  
5 <!-- README.md is generated from README.Rmd. Please edit that file -->  
6  
7 ```{r, include = FALSE}  
8 knitr::opts_chunk$set(  
9   collapse = TRUE,  
10   comment = "#>"  
11 )  
12 ...  
13  
14 # Open Science and Reproducible Research in R: *An Oxford International Health and Tropical Medicine Lecture Series*  
15  
16 <!-- badges: start -->  
17 [![license for data](https://img.shields.io/badge/(for%20data)-CC0-blue.svg)](https://creativecommons.org/publicdomain/zero/1.0/)  
18 [![license for code](https://img.shields.io/badge/(for%20code)-GPL3.0-blue.svg)](https://opensource.org/licenses/gpl-3.0.html)  
19 [![license for slide deck](https://img.shields.io/badge/(for%20slide%20decks)-CC%20BY%204.0-blue.svg)](https://creativecommons.org/licenses/by/4.0/)  
20 [![pages-build-deployment](https://github.com/OxfordIHTM/open-reproducible-science/actions/workflows/pages/pages-build-deployment/badge.svg)]  
  (https://github.com/OxfordIHTM/open-reproducible-science/actions/workflows/pages/pages-build-deployment)  
21 [![DOI](https://zenodo.org/badge/451333046.svg)](https://zenodo.org/badge/latestdoi/451333046)  
22 <!-- badges: end -->  
23  
24 ## Part 1: All about R  
25  
26 [R](https://r-project.org) is a language and environment for statistical computing and graphics. It is a [GNU](https://en.wikipedia.org/wiki/GNU) project  
  which is similar to the [S language and environment](https://en.wikipedia.org/wiki/S_(programming_language)) which was developed at Bell Laboratories  
  (formerly AT&T, now Lucent Technologies) by John Chambers and colleagues. R can be considered as a different implementation of S. There are some important  
  differences, but much code written for S runs unaltered under R.  
27  
28 [R](https://r-project.org) provides a wide variety of statistical (linear and nonlinear modelling, classical statistical tests, time-series analysis,  
  classification, clustering, etc.) and graphical techniques, and is highly extensible. The S language is often the vehicle of choice for research in  
  statistical methodology, and R provides an Open Source route to participation in that activity.  
29  
30 One of [R](https://r-project.org)'s strengths is the ease with which well-designed publication-quality plots can be produced, including mathematical symbols  
  and formulae where needed. Great care has been taken over the defaults for the minor design choices in graphics, but the user retains full control.  
31  
32 R is available as Free Software under the terms of the Free Software Foundation's GNU General Public License in source code form. It compiles and runs on a  
  wide variety of UNIX platforms and similar systems (including FreeBSD and Linux), Windows and MacOS.  
33  
34 R is unique in that it is not general-purpose. It does not compromise by trying to do a lot of things. It does a few things very well, mainly statistical  
  analysis and data visualization. While you can find data analysis and machine learning libraries for languages like [Python](https://www.python.org/), R has  
  many statistical functionalities built into its core. No third-party libraries are needed for much of the core data analysis you can do with the language.
```

## Open Reproducible Science

Open Science and Reproducible Research in R Lecture Series

[View the Project on GitHub](#)  
OxfordIHTM/open-reproducible-science

license (for data) CC0 license (for code) [GPL3.0](#)  
license (for slide decks) CC BY 4.0  pages-build-deployment passing  
DOI [10.5281/zenodo.7515077](#)

### Part 1: All about R

R is a language and environment for statistical computing and graphics. It is a [GNU](#) project which is similar to the [S language and environment](#) which was developed at Bell Laboratories (formerly AT&T, now Lucent Technologies) by John Chambers and colleagues. R can be considered as a different implementation of S. There are some important differences, but much code written for S runs unaltered under R.

R provides a wide variety of statistical (linear and nonlinear modelling, classical statistical tests, time-series analysis, classification, clustering, etc.) and graphical techniques, and is highly extensible. The S language is often the vehicle of choice for research in statistical methodology, and R provides an Open Source route to participation in that activity.

One of R's strengths is the ease with which well-designed publication-quality plots can be produced, including mathematical symbols and formulae where needed. Great care has been taken over the defaults for the minor design choices in graphics, but the user retains full control.

R is available as Free Software under the terms of the Free Software Foundation's GNU General Public License in source code form. It compiles and runs on a wide variety of UNIX platforms and similar systems (including FreeBSD and Linux), Windows and MacOS.

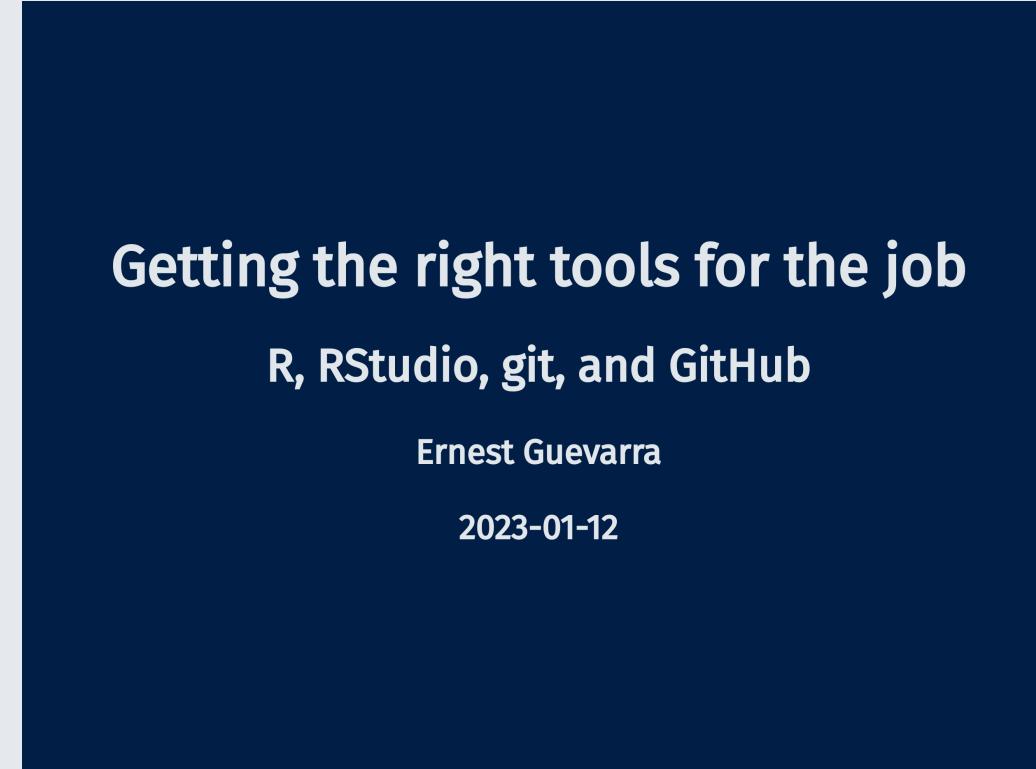
R is unique in that it is not general-purpose. It does not compromise by trying to do a lot of things. It does a few things very well, mainly statistical analysis and data visualization. While you can find data analysis and machine learning libraries for languages like Python, R has many statistical functionalities built into its core. No third-party libraries are needed for much of the core data analysis you can do with the language.

This project is maintained by [OxfordIHTM](#)

Hosted on GitHub Pages — Theme by [orderedlist](#)

# R Markdown example - slide deck

```
273 lines (173 sloc) | 8.7 KB
1  ---
2  title: "Getting the right tools for the job"
3  subtitle: "R, RStudio, git, and GitHub"
4  author:
5    - "Ernest Guevarra"
6  date: '2023-01-12'
7  output:
8    xaringan::moon_reader:
9      css: xaringan-themer.css
10     nature:
11       slideNumberFormat: "%current%"
12       highlightStyle: github
13       highlightLines: true
14       ratio: 16:9
15       countIncrementalSlides: true
16   ...
17
18 ``{r setup, include=FALSE}
19 options(htmltools.dir.version = FALSE)
20 knitr::opts_chunk$set(
21   fig.width=9, fig.height=3.5, fig.retina=3,
22   out.width = "100%",
23   cache = FALSE,
24   echo = TRUE,
25   message = FALSE,
26   warning = FALSE,
27   hline = TRUE
28 )
29
30 if (!require(remotes)) install.packages("remotes")
31 if (!require(fontawesome)) remotes::install_github("rstudio/fontawesome")
32 ...
33
34 ``{r xaringan-themer, include=FALSE, warning=FALSE}
35 library(xaringanthemr)
36 style_mono_light(
37   base_color = "#002147",
38   title_slide_background_image = "",
39   title_slide_background_size = "cover",
40   header_font_google = google_font("Fira Sans"),
41   text_font_google = google_font("Fira Sans Condensed"),
42   text_font_size = "1.2em",
43   link_color = "#214700",
44   header_h1_font_size = "50px",
```



# Basics of R Markdown

- An R Markdown document is written in markdown (an easy-to-write plain text format) and contains chunks of embedded R code, like the document below.

```
---
```

```
output: html_document
```

```
--
```

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see .

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
```{r}
```

```
summary(cars)
```

```
```
```

You can also embed plots, for example:

```
```{r, echo=FALSE}
```

```
plot(cars)
```

```
```
```

Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.

# Basic Markdown syntax

## Markdown syntax

- \*italics\* or italics
- \*\*bold\*\* or bold
- > block quote
- # Header 1
- ## Header 2
- ### Header 3
- ##### Header 4

## Resulting output

- *italics*
- **bold**
- | block quote
- # Header 1
- ## Header 2
- ### Header 3
- #### Header 4

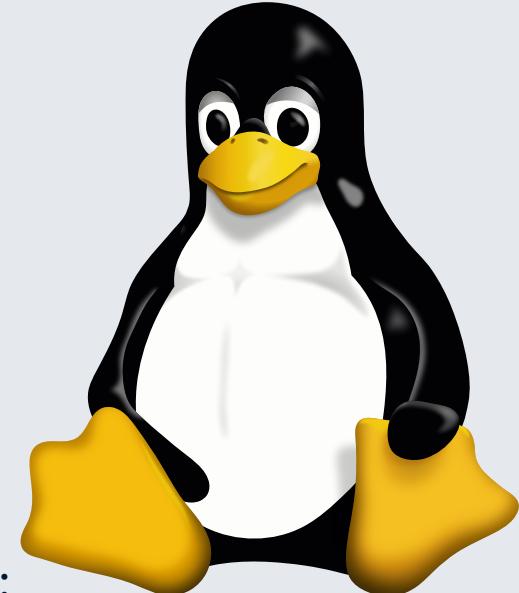
# Basic Markdown syntax

## Markdown syntax

- [link to Open Reproducible Science website](<https://oxford-ihtm.io/open-reproducible-science/>)
- adding an image: 

## Resulting output

- link to Open Reproducible Science website



- adding an image:

# Questions?

# Practical session

We'll work through *Exercise 5 - Extending R with packages* in Practical R for Epidemiologists (<https://practical-r.org/exercise5.html>) as a GitHub Classroom assignment

# Thank you!

Slides can be viewed at <https://oxford-ihtm.io/open-reproducible-science/session5.html>

PDF version of slides can be downloaded at <https://oxford-ihtm.io/open-reproducible-science/pdf/session5-literate-programming.pdf>

R scripts for slides available [here](#)