

Reproducilble Research

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1 What is reproducible research

In science replicability is of prime importance. There is a difference between “replication” and “reproduction”. Replication of an experiments is to repeat the same experiment to see whether the result holds up. There is a assumption that the results will differ next time. The results of experimental replication are of two types: biological and technical replicates. Unlike replication, reproduction of and experment is to repeat is exactly to check whether another groups’ results holds up under scrutiny. This is of vital importance.

In bioinformatics, reproducible research is a concept where data and code are packaged together into one live document. It provides the facility to reproduce the data analysis step exactly on another computer.

2 Text processing vs word processing

In word processing (like in Microsoft Word), the environment is WYSIWYG (What You See Is What You Get). You exactly see the formatting of the document live. In text processing, you think about the overall structure of the document and **markup** the document is special syntax then use a software to convert the document in more presentation format, like PDF. The prime software for text processing in scientific computing is L^AT_EX.

3 L^AT_EX

L^AT_EX has been a mainstay of scientific publishing. A very simple document in L^AT_EX is:

```
\documentclass{article}
\title{My first document}
\author{Malay}
\begin{document}
\maketitle
\section{This is the first section}
Hello World!
\end{document}
```

3.1 Exercise

Type the above code block in a text file. Name the file as `hello.tex`. You can compile the document in L^AT_EX using

```
xelatex hello.tex
```

Note: You might have to install xelatex and the associated packages.

4 Pandoc

L^AT_EX is a type-setting software and no one should use it directly. **Pandoc** (<http://johnmacfarlane.net/pandoc/>) is a highly useful text conversion software that can convert any form of text to another. Additionally, it supports a variant of a very popular and simple text markup language called **markdown**. When you convert a **markdown** formatted file to PDF using **pandoc**, it uses L^AT_EX behind the scene. This makes our job of creating L^AT_EX a lot easier.

Let's create our first **pandoc** file.

```
---
title: My first document
author: Malay
---
# This is the first section
Hello World!
```

Type the above code in a file called `hello1.md`. Then type the following:

```
pandoc --number-sections -o hello1.pdf hello1.md
```

Interesting thing is that the same source can be now used to create an HTML document:

```
pandoc -s --number-sections -o hello1.html hello1.md
```

Also a Microsoft Word file:

```
pandoc -o hello1.docx hello1.md
```

5 Literate programming

1. Originally created by Don Knuth.
2. Document that describes the data analysis and the code are in the same text file.
3. You **weave** a document to create the human readable text.
4. You **tangle** a document to create the machine readable code.

6 knitr

knitr is a package in R to mix **markdown** formatted document with R code called, **Rmarkdown**(<http://rmarkdown.rstudio.com/>). The R code is live and executed on the fly to generate the final document with the result of the R already embedded in. The overall steps are:

```
.Rmd -> (use knitr)-> .md -> (use pandoc)-> HTML or PDF
```

Within RStudio all these steps are put together under one button click.

6.1 Exercise

Type the following code into RStudio:

```
---
title: My first knitr document
author: Me
---

This is some text (text chunk).

Here is some code (code chunk).

```{r}
set.seed(1)
x<-rnorm(100)
mean(x)
```
```

6.2 Chunk options

For a list of chunk options see http://yihui.name/knitr/options#chunk_options. There are several options you can use to control what should be included in the final document. For example, **echo=FALSE** with not include the code in the final document.

```
```{r echo=FALSE}
set.seed(1)
x<-rnorm(100)
mean(x)
```
```

results="hide" will hide the output but the code will still be shown.

```
```{r results="hide"}
set.seed(1)
x<-rnorm(100)
mean(x)
```
```

include=FALSE will suppress both the output and the code, but the code will be executed.

```
```{r include=FALSE}
set.seed(1)
x<-rnorm(100)
mean(x)
```
```

```
head(x)
```

```
## [1] -0.6264538  0.1836433 -0.8356286  1.5952808  0.3295078 -0.8204684
```

To prevent the execution of the code, use `eval=FALSE`.

6.3 Global chunk option

You can set a chunk option globally for every chunk.

```
``{r include=FALSE}
knitr::opts_chunk$set(message=FALSE)
``
```

One important option for chunk is caching R calculations.

```
``{r include=FALSE}
knitr::opts_chunk$set(cache=TRUE)
``
```

6.4 In-line R

You can include a piece of data from the calculation inside a text chunk.

We have ``r length(x)`` rows in our data.

6.5 Nice tables

You can use R package `xtable` for creating nice looking table.

```
``{r fitmodel}
library(datasets)
data(airquality)
fit <- lm(Ozone ~ Wind, data=airquality)
``
```

We can now print a nice looking table of regression coefficients.

```
library(xtable)
options(xtable.comment=FALSE)
xt <- xtable( summary(fit) )
print.xtable(xt, type="latex")
```

| | Estimate | Std. Error | t value | Pr(> t) |
|-------------|----------|------------|---------|----------|
| (Intercept) | 96.8729 | 7.2387 | 13.38 | 0.0000 |
| Wind | -5.5509 | 0.6904 | -8.04 | 0.0000 |

You can also use the `knitr` function `kable` to create a table.

```
knitr::kable(summary(airquality))
```

| Ozone | Solar.R | Wind | Temp | Month | Day |
|----------------|---------------|----------------|---------------|---------------|--------------|
| Min. : 1.00 | Min. : 7.0 | Min. : 1.700 | Min. :56.00 | Min. :5.000 | Min. : 1.0 |
| 1st Qu.: 18.00 | 1st Qu.:115.8 | 1st Qu.: 7.400 | 1st Qu.:72.00 | 1st Qu.:6.000 | 1st Qu.: 8.0 |
| Median : 31.50 | Median :205.0 | Median : 9.700 | Median :79.00 | Median :7.000 | Median :16.0 |
| Mean : 42.13 | Mean :185.9 | Mean : 9.958 | Mean :77.88 | Mean :6.993 | Mean :15.8 |
| 3rd Qu.: 63.25 | 3rd Qu.:258.8 | 3rd Qu.:11.500 | 3rd Qu.:85.00 | 3rd Qu.:8.000 | 3rd Qu.:23.0 |
| Max. :168.00 | Max. :334.0 | Max. :20.700 | Max. :97.00 | Max. :9.000 | Max. :31.0 |
| NA's :37 | NA's :7 | NA | NA | NA | NA |

7 Git and github

Git is a version control system, created by Linus Tovalds. Github is website for storing and sharing `git` repositories.

7.1 Demo how to create github account and create a repository

Check the Jupyter notebook.
