

# Reproducible Analyses with knitr and rmarkdown

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# Introduction

# NIH plans to enhance reproducibility

**Francis S. Collins** and **Lawrence A. Tabak** discuss initiatives that the US National Institutes of Health is exploring to restore the self-correcting nature of preclinical research.

A growing chorus of concern, from scientists and laypeople, contends that the complex system for ensuring the reproducibility of biomedical research is failing and is in need of restructuring<sup>1,2</sup>. As leaders of the US National Institutes of Health (NIH), we share this concern and here explore some of the significant interventions that we are planning.

Science has long been regarded as 'self-correcting', given that it is founded on the replication of earlier work. Over the long term, that principle remains true. In the

shorter term, however, the checks and balances that once ensured scientific fidelity have been hobbled. This has compromised the ability of today's researchers to reproduce others' findings.

Let's be clear: with rare exceptions, we have no evidence to suggest that irreproducibility is caused by scientific misconduct. In 2011, the Office of Research Integrity of the US Department of Health and Human Services pursued only 12 such cases<sup>3</sup>. Even if this represents only a fraction of the actual problem, fraudulent papers are vastly

At the lab?

## Trouble at the lab

Scientists like to think of science as self-correcting. To an alarming degree, it is not

Oct 19th 2013 | From the print edition

 *Timekeeper*

 Like

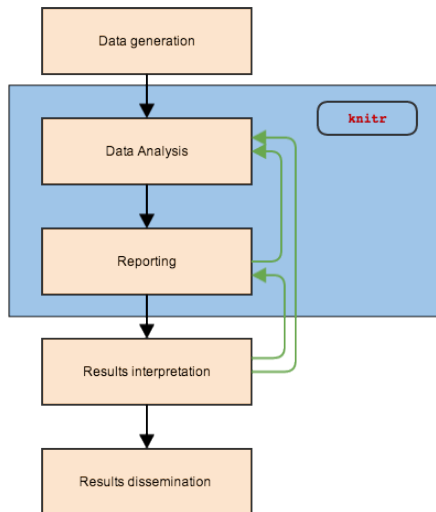
22k

 Tweet

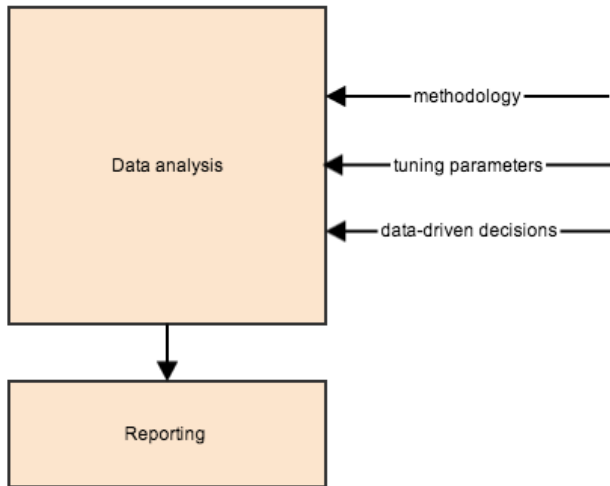
2,182



# Where do we fit in?



# Data analysis



Goal: code + prose = report

# Solution

- ▶ Cut and paste for report production is not a viable method
  - ▶ tedious
  - ▶ slow
  - ▶ error-prone
- ▶ Incorporate analysis code into text documents
  - ▶ `knit` to create results
  - ▶ results get incorporated into document
  - ▶ post-process to get any type of output format
- ▶ Literate documents
  - ▶ R Code creates results to inform report
  - ▶ Prose surrounding code informs analysis

# How? Markdown.

This is a first-level heading

#####

Second level

-----

### Third

Lists

- one
- two
- three

> Blockquotes are awesome

> - Me

Italics for *\*emphasis\** or bold for **\*\*more emphasis\*\***

```
```{r my-first-chunk, results='asis'}
```

```
## code goes in here
```

```
```
```

## This is a first-level heading

### Second level

#### Third

Lists

- one
- two
- three

Blockquotes are awesome

-Me

Italics for *emphasis* or bold for **more emphasis**

```
## code goes in here
```



# Contrast with html

```
<div id="this-is-a-first-level-heading" class="section level1">
<h1>This is a first-level heading</h1>
<div id="second-level" class="section level2">
<h2>Second level</h2>
<div id="third" class="section level3">
<h3>Third</h3>
<p>Lists</p>
<ul>
<li>one</li>
<li>two</li>
<li>three</li>
</ul>
<blockquote>
<p>Blockquotes are awesome</p>
<p>-Me</p>
</blockquote>
<p>Italics for <em>emphasis</em> or bold for <strong>more emphasis</strong></p>
<pre class="r"><code>## code goes in here</code></pre>
</div>
</div>
</div>
```

## ... and tex

```
\documentclass[]{article}
\begin{document}
\begin{center}
\normalsize
\end{center}
\section{This is a first-level heading}\label{this-is-a-first-level-heading}
```

```
\subsection{Second level}\label{second-level}
```

```
\subsubsection{Third}\label{third}
```

Lists

```
\begin{itemize}
\itemsep1pt\parskip0pt\parsep0pt
\item
  one
\item
  two
\item
  three
\end{itemize}
```

```
\begin{quote}
Blockquotes are awesome - Me
\end{quote}
```

Italics for `\emph{emphasis}` or bold for `\textbf{more emphasis}`

```
\begin{Shaded}
\begin{Highlighting}[]
\NormalTok{## code goes in here}
\end{Highlighting}
\end{Shaded}

\end{document}
```

## Incorporating code chunks

Three backticks, each chunk needs a unique name:

```
```{r my-first-chunk}
## code goes in here and gets evaluated
t.test(mpg ~ vs, data = mtcars)
```

##
## Welch Two Sample t-test
##
## data:  mpg by vs
## t = -4.6671, df = 22.716, p-value = 0.0001098
## alternative hypothesis: true difference in means is not
## 95 percent confidence interval:
##  -11.462508  -4.418445
## sample estimates:
## mean in group 0 mean in group 1
##          16.61667          24.55714
```

# Try it

- ▶ Open Rstudio
- ▶ Click New > Rmarkdown
- ▶ Select output format

Examine the markdown prose and the code.

What do you expect the output to look like?

- ▶ Click knit
- ▶ What do you get?

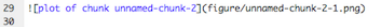
# How it works: knitr

Code chunks are evaluated sequentially the same, fresh R session:

## Rmd

```
1- ---
2- title: "Untitled"
3- author: "Michael Sachs"
4- date: "March 30, 2015"
5- output: html_document
6- ---
7-
8- This is an R Markdown document. Markdown is a simple formatting
9- syntax for authoring HTML, PDF, and MS Word documents. For more
10- details on using R Markdown see <http://rmarkdown.rstudio.com>.
11-
12- When you click the Knit button a document will be generated
13- that includes both content as well as the output of any embedded R
14- code chunks within the document. You can embed an R code chunk
15- like this:
16-
17- ```{r}
18- summary(cars)
19- ```
20-
21- You can also embed plots, for example:
22-
23- ```{r, echo=FALSE}
24- plot(cars)
25- ```
26-
27- Note that the `echo = FALSE` parameter was added to the code chunk
28- to prevent printing of the R code that generated the plot.
```

## md

```
1- ---
2- title: "Untitled"
3- author: "Michael Sachs"
4- date: "March 30, 2015"
5- output: html_document
6- ---
7-
8- This is an R Markdown document. Markdown is a simple formatting
9- syntax for authoring HTML, PDF, and MS Word documents. For more
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12- When you click the Knit button a document will be generated
13- that includes both content as well as the output of any embedded R
14- code chunks within the document. You can embed an R code chunk
15- like this:
16-
17- ```r
18- summary(cars)
19- ```
20-
21-
22- ## speed dist
23- ## Min. : 4.0 Min. : 2.00
24- ## 1st Qu.:12.0 1st Qu.: 26.00
25- ## Median :15.0 Median : 36.00
26- ## Mean :15.4 Mean : 42.98
27- ## 3rd Qu.:19.0 3rd Qu.: 56.00
28- ## Max. :25.0 Max. :120.00
29- ```
30-
31- You can also embed plots, for example:
32-
33- 
34-
35- Note that the `echo = FALSE` parameter was added to the code chunk
36- to prevent printing of the R code that generated the plot.
```

# How it works: pandoc

- ▶ Pandoc converts the markdown format to some other document type:
  - ▶ Word
  - ▶ Html
  - ▶ Pdf
  - ▶ ...
- ▶ Templates and output formats for specific uses:
  - ▶ Beamer slideshows
  - ▶ ioslides presentations
  - ▶ Tufte-style handouts

# Caveats

- ▶ Markdown is minimalistic
- ▶ Easy to write and read

rmarkdown + knitr is designed to *quickly* and *simply* generate analytic reports with minimal markup

- ▶ Not complex or precise enough for **complete** control over output
- ▶ How much precision do you need?

# Markdown specs

- ▶ Paragraphs, # headers, ## subheader, etc, > blockquotes
- ▶ Emphasis, *italics*, **bold**
- ▶ Images/links: ![name](pathtoimage), [text](link)
- ▶ Lists/ordered lists
- ▶ Code chunks
- ▶ Latex equations:  $\sum_{i=1}^n X_i/n = \sum_{i=1}^n X_i/n$
- ▶ Tables
- ▶ Citations: [@citekey], bibtex, endnote, others supported



# Front matter

- ▶ Metadata
- ▶ Document types
- ▶ Other options
  - ▶ Default figure size, table of contents, theme
  - ▶ See <http://rmarkdown.rstudio.com/> for complete documentation

---

title: "Reproducible Analyses with knitr and rmarkdown"

author: "Michael Sachs"

date: "March 31, 2015"

output:

ioslides\_presentation:

widescreen: true

---

## Controlling R output

# knitr chunk output

## Results

Default: results = 'markup'

```
```{r markup, results = 'markup'}  
head(mtcars, 4)  
```
```

| ## |                | mpg  | cyl | disp | hp  | drat | wt    | qsec  | vs | am |
|----|----------------|------|-----|------|-----|------|-------|-------|----|----|
| ## | Mazda RX4      | 21.0 | 6   | 160  | 110 | 3.90 | 2.620 | 16.46 | 0  | 1  |
| ## | Mazda RX4 Wag  | 21.0 | 6   | 160  | 110 | 3.90 | 2.875 | 17.02 | 0  | 1  |
| ## | Datsun 710     | 22.8 | 4   | 108  | 93  | 3.85 | 2.320 | 18.61 | 1  | 1  |
| ## | Hornet 4 Drive | 21.4 | 6   | 258  | 110 | 3.08 | 3.215 | 19.44 | 1  | 0  |

## knitr chunk output

```
results = 'asis'
```

```
```{r asis, results = 'asis'}  
head(mtcars, 4)  
```
```

|  | mpg | cyl | disp | hp | drat | wt | qsec | vs | am | gear | car |
|--|-----|-----|------|----|------|----|------|----|----|------|-----|
|--|-----|-----|------|----|------|----|------|----|----|------|-----|

|           |      |     |      |       |       |       |       |   |   |   |                |            |
|-----------|------|-----|------|-------|-------|-------|-------|---|---|---|----------------|------------|
| Mazda RX4 | 21.0 | 6   | 160  | 110   | 3.90  | 2.620 | 16.46 | 0 | 1 | 4 | 4              | Mazda RX4  |
| Wag       | 21.0 | 6   | 160  | 110   | 3.90  | 2.875 | 17.02 | 0 | 1 | 4 | 4              | Datsun 710 |
| 22.8      | 4    | 108 | 93   | 3.85  | 2.320 | 18.61 | 1     | 1 | 4 | 1 | Hornet 4 Drive | 21.4       |
| 6         | 258  | 110 | 3.08 | 3.215 | 19.44 | 1     | 0     | 3 | 1 |   |                |            |

## knitr chunk output

Make tables pretty with `knitr::kable` and `results = 'asis'`

```
kable(head(mtcars, 4), digits = 1)
```

|                | mpg  | cyl | disp | hp  | drat | wt  | qsec | vs | am |
|----------------|------|-----|------|-----|------|-----|------|----|----|
| Mazda RX4      | 21.0 | 6   | 160  | 110 | 3.9  | 2.6 | 16.5 | 0  | 1  |
| Mazda RX4 Wag  | 21.0 | 6   | 160  | 110 | 3.9  | 2.9 | 17.0 | 0  | 1  |
| Datsun 710     | 22.8 | 4   | 108  | 93  | 3.8  | 2.3 | 18.6 | 1  | 1  |
| Hornet 4 Drive | 21.4 | 6   | 258  | 110 | 3.1  | 3.2 | 19.4 | 1  | 0  |

## Tables and other output

Several other packages are available to customize table output:

- **pander**: Good for printing output from regression models:

```
pander::pander(lm(mpg ~ factor(cyl), data = mtcars))
```

Table 2:Fitting linear model: mpg ~ factor(cyl)

|                     | Estimate | Std. Error | t value | Pr(> t )  |
|---------------------|----------|------------|---------|-----------|
| <b>factor(cyl)6</b> | -6.921   | 1.558      | -4.441  | 0.0001195 |
| <b>factor(cyl)8</b> | -11.56   | 1.299      | -8.905  | 8.568e-10 |
| <b>(Intercept)</b>  | 26.66    | 0.9718     | 27.44   | 2.688e-22 |

## Other options

- ▶ `include = FALSE` evaluates code but doesn't include anything
- ▶ `echo = FALSE` don't display results
- ▶ `warning = FALSE` don't display warnings
- ▶ `cache = TRUE` cache results for long-running stuff
- ▶ `comment = NA` hide # from markup output

# Figure options

The important ones:

- ▶ `fig_width`, `fig_height`, in inches. Can also be set globally in the header.
- ▶ `fig_align`, left, right or center
- ▶ `fig_cap` = "Caption" add caption to figure



# Try it!

- ▶ Create a new rmarkdown document with an output format of your choice
- ▶ Use the BMI.CSV dataset to perform some basic analysis
- ▶ Display the data using `head`
- ▶ Summarize the data
- ▶ Do a t-test
- ▶ Make a figure using `ggplot`
- ▶ Make a table of regression coefficients
- ▶ *Write about what you are doing along the way*

# Summary

- ▶ Analysis + report writing should be easy and integrated
- ▶ knitr + rmarkdown + pandoc all in Rstudio
- ▶ You don't have to remember everything, use the menus and help documents
- ▶ Your future self will thank you for having a reproducible analysis

# Resources

| Topic                   | Link  |
|-------------------------|---|
| KBroman's UWisc Class   | <a href="https://kbroman.github.io/knitr_knu">https://kbroman.github.io/knitr_knu</a> |
| Knitr homepage          | <a href="http://yihui.name/knitr/">http://yihui.name/knitr/</a>                       |
| rmarkdown documentation | <a href="http://rmarkdown.rstudio.com/">http://rmarkdown.rstudio.com/</a>             |
| Another knitr tutorial  | <a href="http://sachsmc.github.io/knit-git-m">http://sachsmc.github.io/knit-git-m</a> |
| Pandoc reference        | <a href="http://johnmacfarlane.net/pandoc/">http://johnmacfarlane.net/pandoc/</a>     |