

# Reproducible Document Stack

Naomi Penfold, Innovation Officer  
Email: [innovation@elifesciences.org](mailto:innovation@elifesciences.org)  
Twitter: @eLifeInnovation

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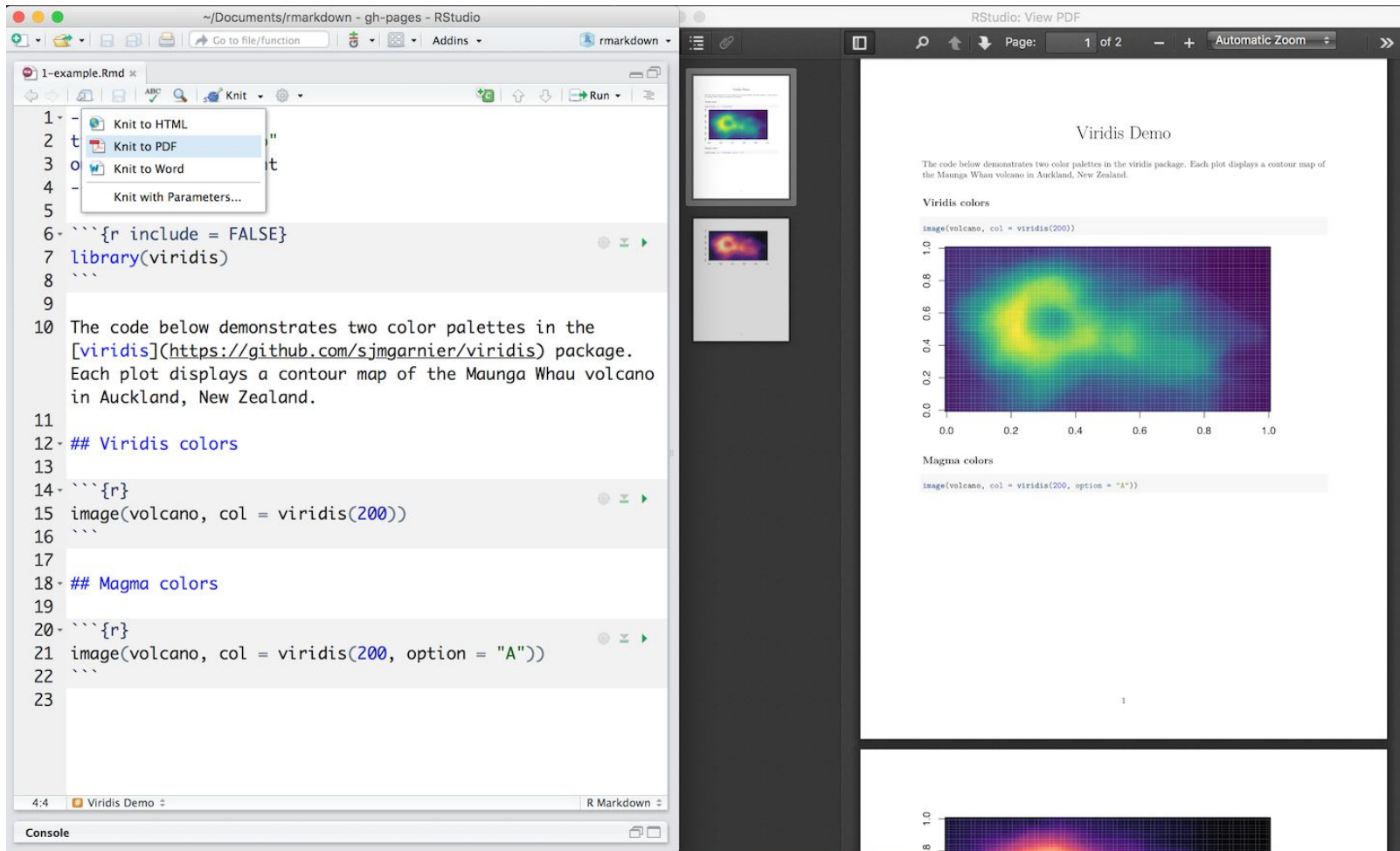


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# Is this you?

# Is this you?



The screenshot displays the RStudio interface with a file named '1-example.Rmd' open. A context menu is visible over the code editor, offering options: 'Knit to HTML', 'Knit to PDF' (highlighted), 'Knit to Word', and 'Knit with Parameters...'. The code in the editor includes R Markdown syntax for including an R chunk, loading the 'viridis' library, and two code blocks that generate contour maps of the Maunga Whau volcano using the 'viridis' color palette. The first block uses 'viridis(200)' and the second uses 'viridis(200, option = "A")'. The right pane shows the rendered PDF, titled 'Viridis Demo', which contains the same text and two contour plots. The top plot is labeled 'Viridis colors' and the bottom plot is labeled 'Magma colors'. The PDF viewer shows 'Page: 1 of 2' and 'Automatic Zoom'.

```
1- example.Rmd
1- 
2- t
3- o
4- 
5- 
6- ```{r include = FALSE}
7- library(viridis)
8- ```
9- 
10- The code below demonstrates two color palettes in the
11- [viridis](https://github.com/sjmgarnier/viridis) package.
12- Each plot displays a contour map of the Maunga Whau volcano
13- in Auckland, New Zealand.
14- 
15- ## Viridis colors
16- 
17- ```{r}
18- image(volcano, col = viridis(200))
19- ```
20- 
21- ## Magma colors
22- 
23- ```{r}
24- image(volcano, col = viridis(200, option = "A"))
25- ```
26- 
```

Viridis Demo

The code below demonstrates two color palettes in the [viridis](https://github.com/sjmgarnier/viridis) package. Each plot displays a contour map of the Maunga Whau volcano in Auckland, New Zealand.

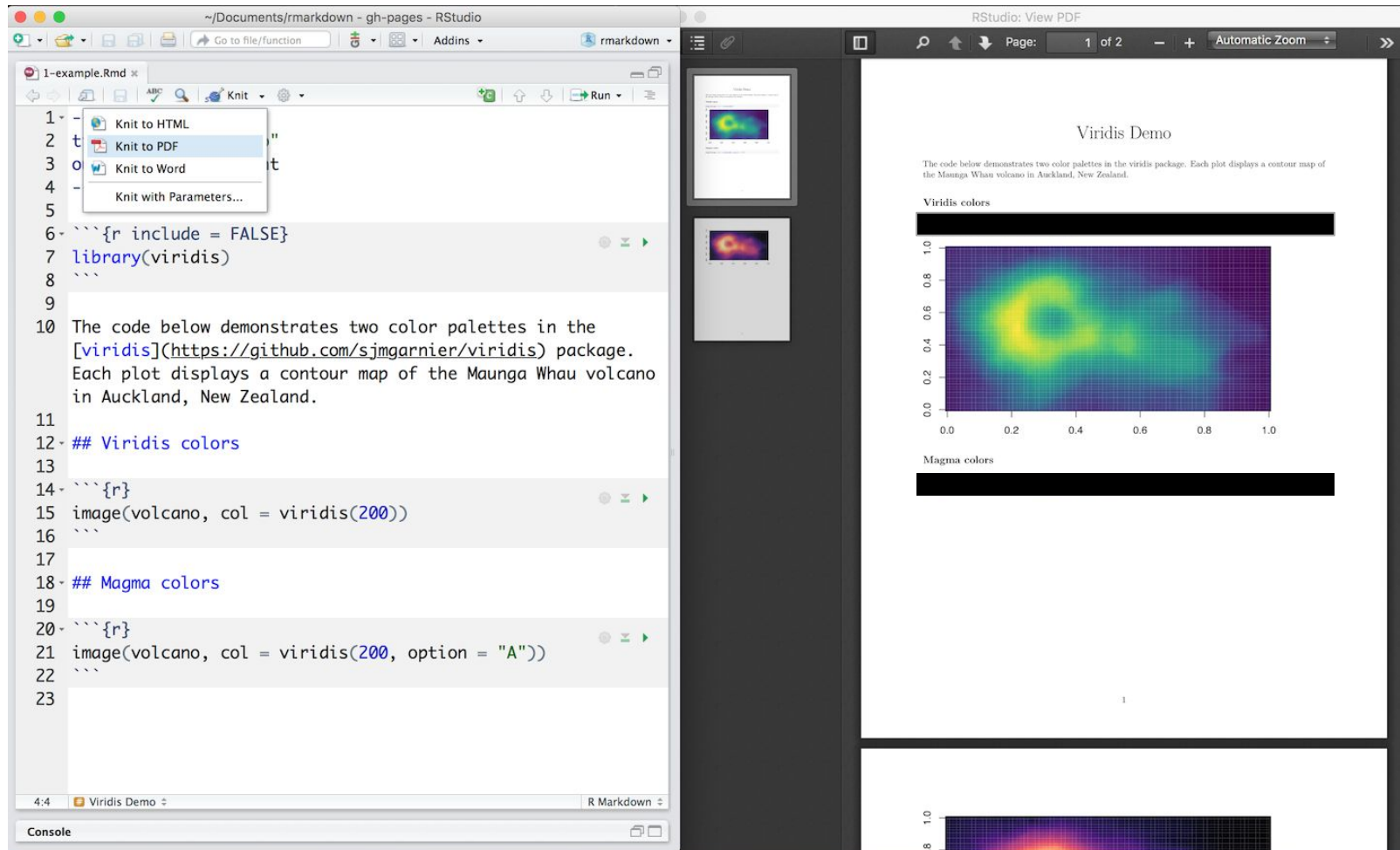
Viridis colors

```
image(volcano, col = viridis(200))
```

Magma colors

```
image(volcano, col = viridis(200, option = "A"))
```

# Is this you?



The screenshot displays the RStudio interface. On the left, the R Markdown source file '1-example.Rmd' is open. A context menu is visible over the code editor, showing options: 'Knit to HTML', 'Knit to PDF' (highlighted), 'Knit to Word', and 'Knit with Parameters...'. The code in the editor includes a YAML header, a title 'Viridis Demo', and two code blocks using the 'viridis' package to generate contour maps of the Maunga Whau volcano. The right pane shows the rendered PDF output, which includes the title, a description of the code, and two contour plots: 'Viridis colors' and 'Magma colors'. The 'Viridis colors' plot is a square contour map, while the 'Magma colors' plot is a horizontal rectangular contour map. Both plots show a color gradient from purple to yellow, representing the elevation of the volcano.

```

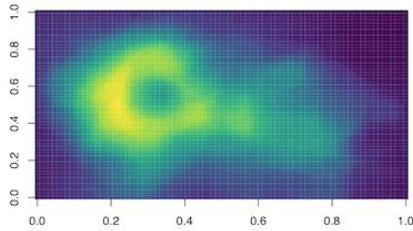
1- example.Rmd
2- title: "Viridis Demo"
3- output: pdf_document
4-
5-
6- {r include = FALSE}
7- library(viridis)
8-
9-
10- The code below demonstrates two color palettes in the
11- [viridis](https://github.com/sjmgarnier/viridis) package.
12- Each plot displays a contour map of the Maunga Whau volcano
13- in Auckland, New Zealand.
14-
15- ## Viridis colors
16- {r}
17- image(volcano, col = viridis(200))
18-
19- ## Magma colors
20- {r}
21- image(volcano, col = viridis(200, option = "A"))
22-
23-

```


Viridis Demo

The code below demonstrates two color palettes in the [\[viridis\]\(https://github.com/sjmgarnier/viridis\)](https://github.com/sjmgarnier/viridis) package. Each plot displays a contour map of the Maunga Whau volcano in Auckland, New Zealand.

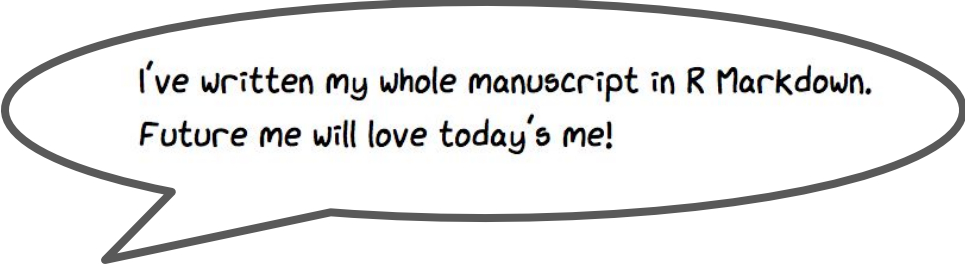
Viridis colors



Magma colors

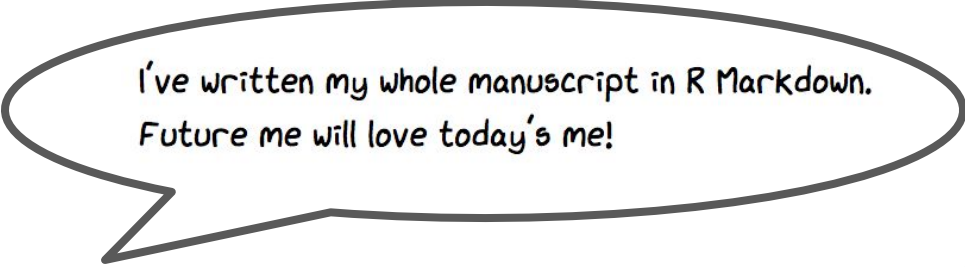


# Is this you?




I've written my whole manuscript in R Markdown.  
Future me will love today's me!

# Is this you?



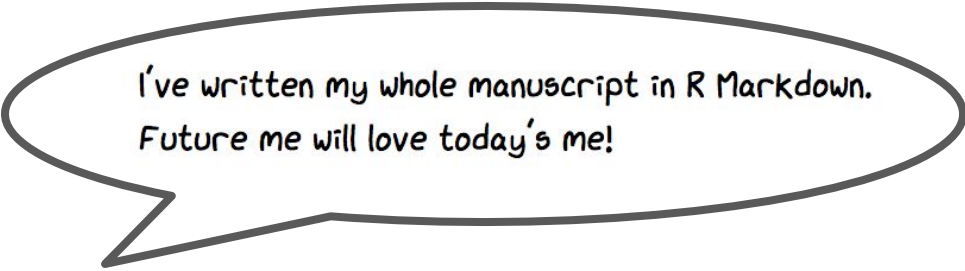
I've written my whole manuscript in R Markdown.  
Future me will love today's me!




Please could you send it to me in Word  
so I can track changes?



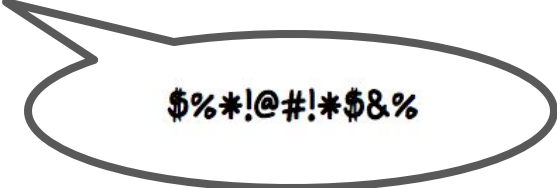
# Is this you?



I've written my whole manuscript in R Markdown.  
Future me will love today's me!

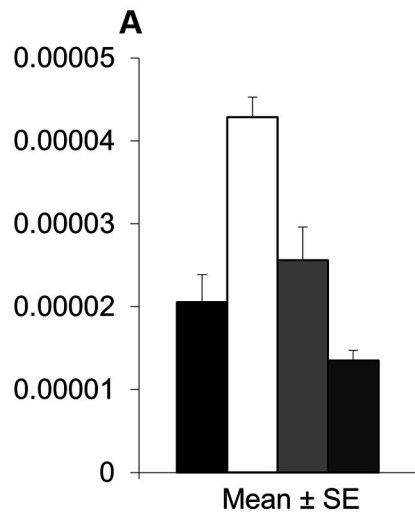


Please could you send it to me in Word  
so I can track changes?

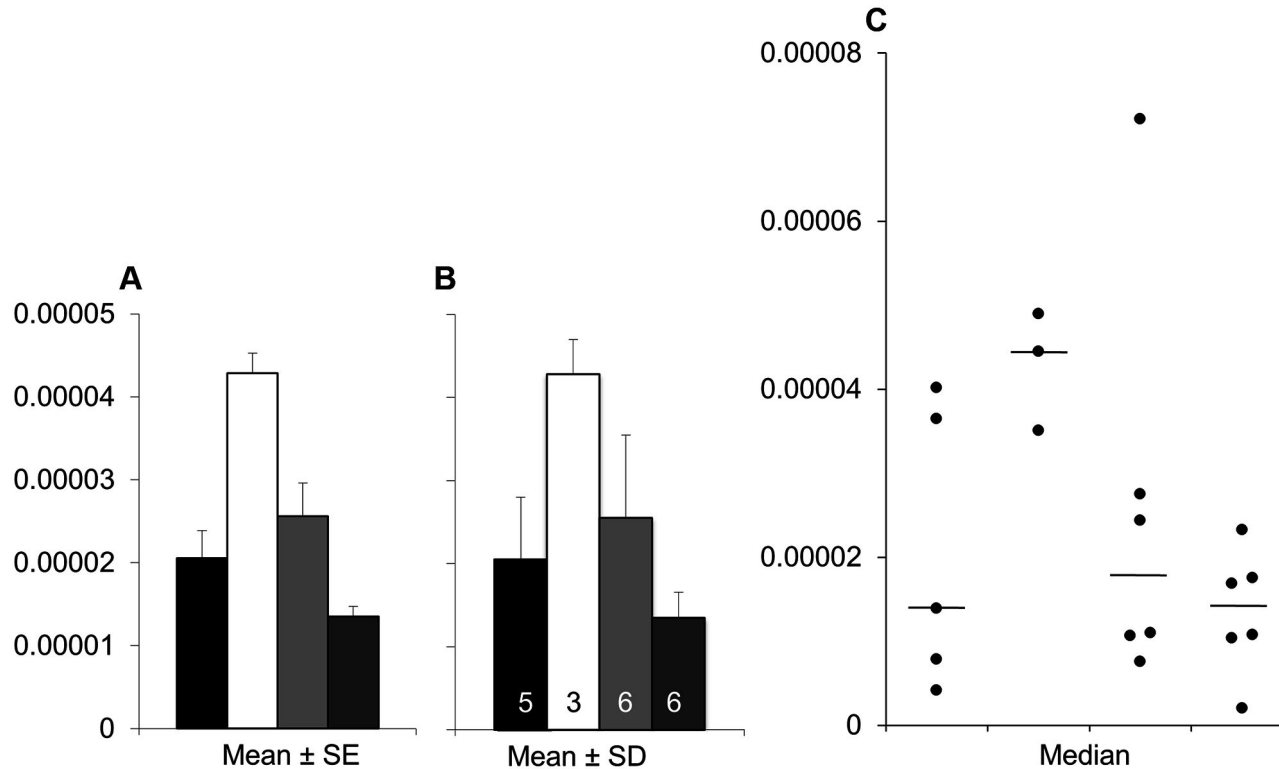


`$%*!@#!*$&%`

# Is this you?



# Is this you?



# Is this you?

LIKELIHOOD YOU WILL GET CODE WORKING  
BASED ON HOW YOU'RE SUPPOSED TO INSTALL IT:



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## Recent progress

- Stencila have improved support for R and Python code in reproducible articles and R Markdown and Jupyter converters
- Substance have built the first reader interface; eLife are conducting user tests now



# DEMO





Try Stencila: <https://stenci.la/>

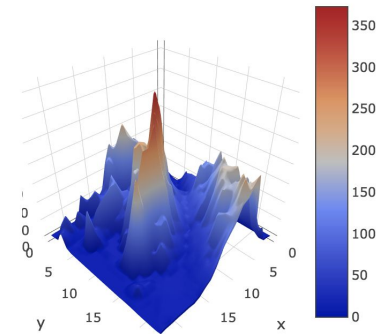
# Recent progress

- Stencila have improved support for R and Python code in reproducible articles and R Markdown and Jupyter converters
- Substance have built the first reader interface; eLife are conducting user tests now
- The first reproducible article is being prepared (see <https://github.com/stencila/examples/tree/master/elife-30274>)

Figure 2

Formula for: Plotly

status: **ready**



## Digital gene expression analysis.

P493-6 cells grown in the presence of tetracycline (Tet) for 72 hr for repression of the conditional *pmyc-tet* construct, were switched into Tet-free growth medium to induce c-Myc expression. Cells were cultured in two separate lots of serum. Transcripts/cell estimates from NanoString nCounter gene expression assays (1369 genes assay) for active (left) and silent (right) genes at 0, 1, and 24 hr after release from Tet. Active genes expressed greater than one transcript/cell. Silent genes expressed less than 0.5 transcript/cell. Box and whisker plots with median represented as the line through the box and whiskers representing values within 1.5 IQR of the first and third quartile. Cells grown in serum lot one: active genes = 708, silent genes = 580. Cells grown in serum lot two: active genes = 719, silent genes = 573. Confirmatory analysis is reported in [Table 1](#) and exploratory statistical analysis is reported in [Table 2](#) and [Table 3](#). Additional details for this experiment can be found at <https://osf.io/fn2y4/>.

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- Share your use case
- Provide feedback
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# Questions?

Email: [n.penfold@elifesciences.org](mailto:n.penfold@elifesciences.org)

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