

Reproducible Document Stack

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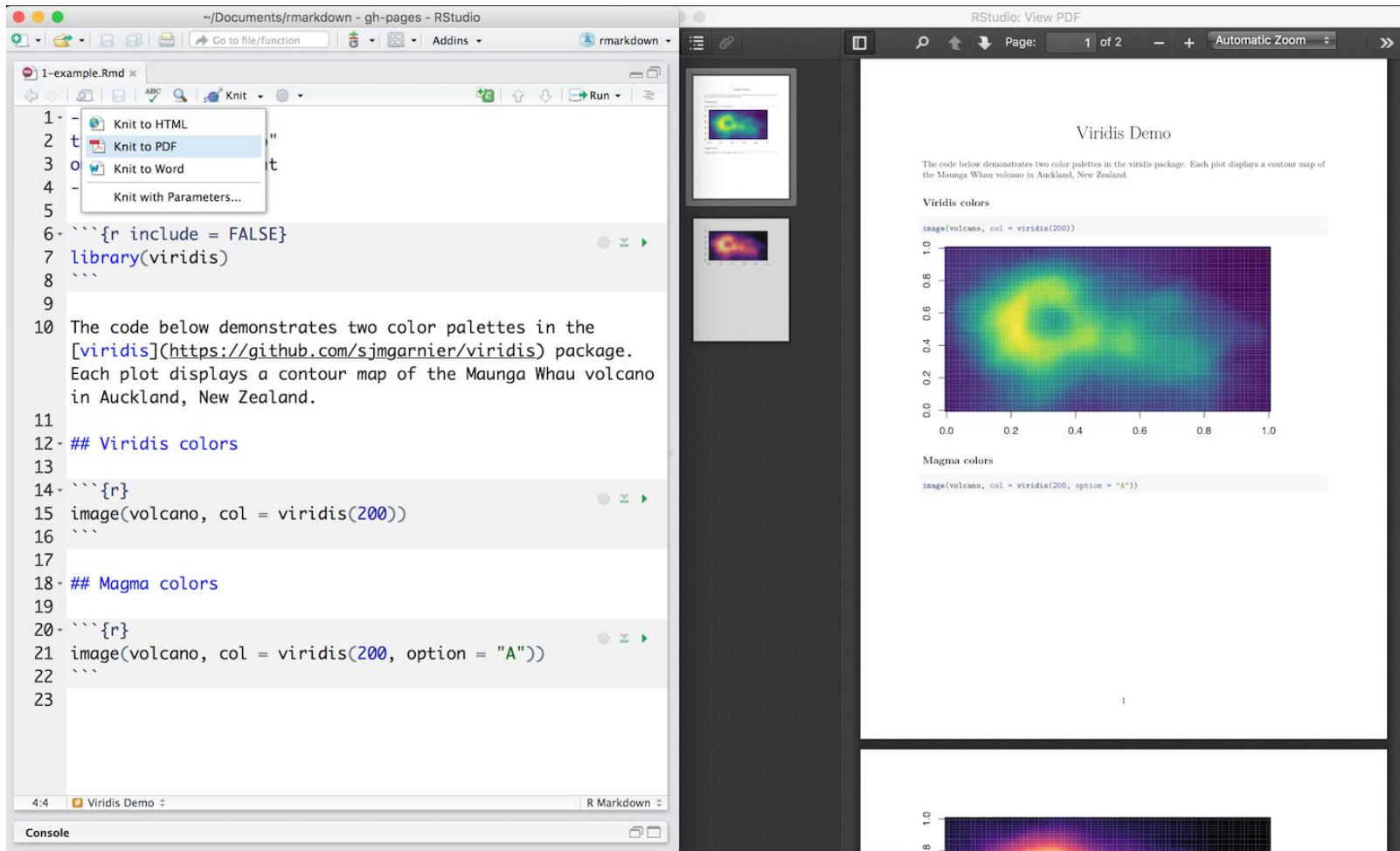


eLIFE

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operating a platform for research communication
that encourages and recognises the most
responsible behaviours in science

Is this you?

Is this you?



The screenshot displays the RStudio interface with an R Markdown document open. The editor shows the following code:

```
1- example.Rmd
2-
3-
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- ```
```

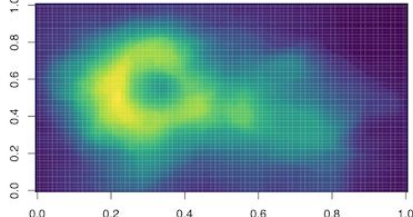
The right pane shows the rendered PDF output, titled "Viridis Demo". It contains the following text and plots:

Viridis Demo

The code below demonstrates two color palettes in the viridis package. Each plot displays a contour map of the Maunga Whau volcano in Auckland, New Zealand.


Viridis colors

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



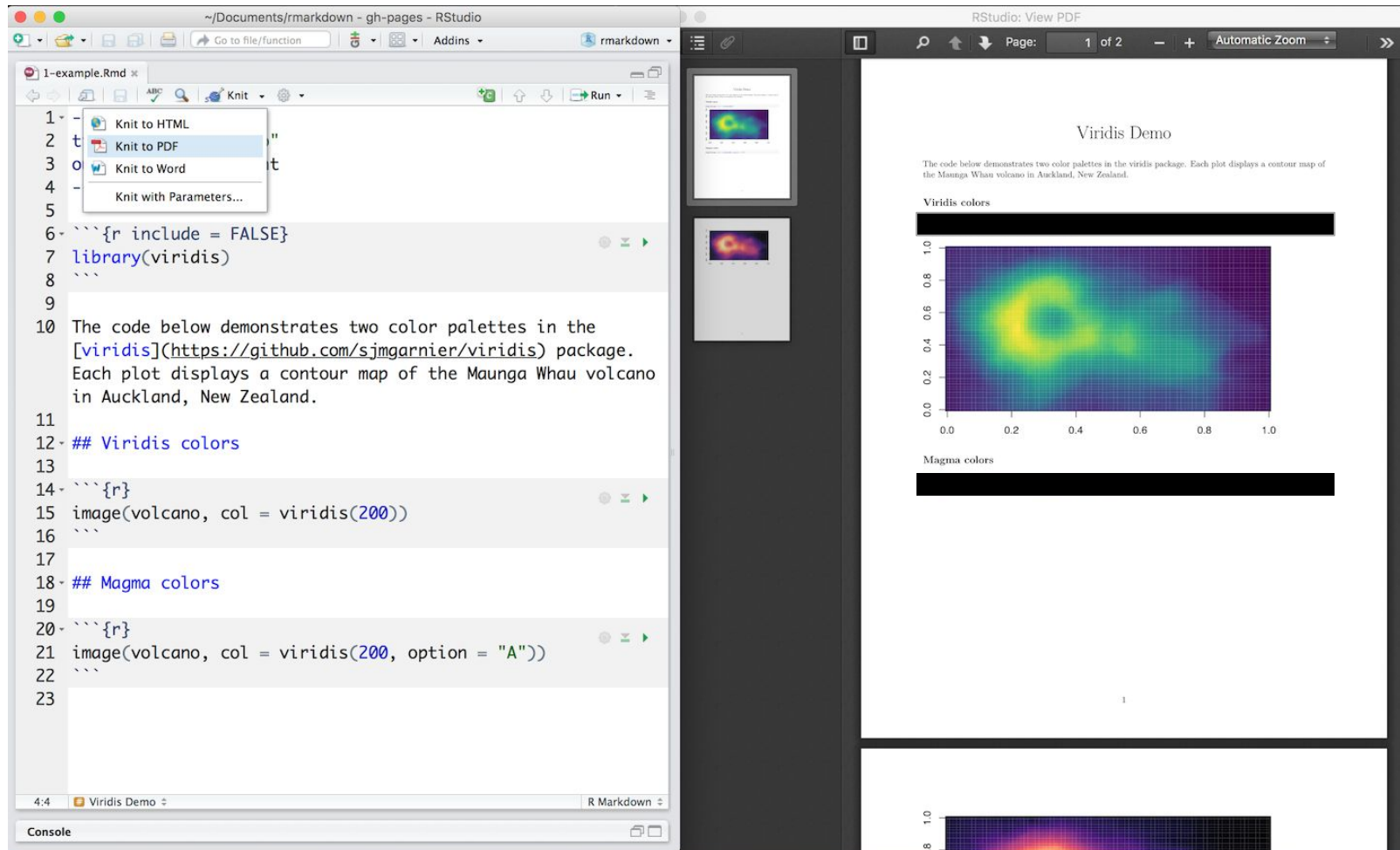
Magma colors

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



The rendered PDF shows two contour maps of the Maunga Whau volcano. The first map uses the viridis color palette, and the second map uses the viridis color palette with option "A".

Is this you?



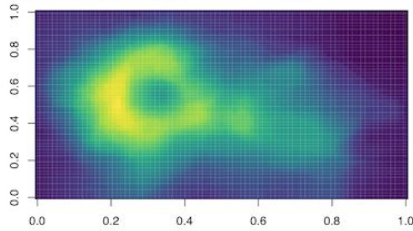
The screenshot displays the RStudio interface with a file named '1-example.Rmd' open. A context menu is visible over the code editor, offering options: 'Knit to HTML', 'Knit to PDF' (highlighted), 'Knit to Word', and 'Knit with Parameters...'. The code in the editor includes R Markdown syntax for including an R chunk, loading the 'viridis' library, and generating two contour plots of the Maunga Whau volcano using different color palettes. The rendered PDF on the right, titled 'Viridis Demo', shows the resulting output. It includes a title, a descriptive paragraph, and two sections: 'Viridis colors' and 'Magma colors', each featuring a contour plot. The 'Viridis colors' plot is a square with axes from 0.0 to 1.0, showing a color gradient from purple to yellow. The 'Magma colors' plot is a horizontal rectangle with axes from 0.0 to 1.0, showing a color gradient from purple to red. 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}
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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.

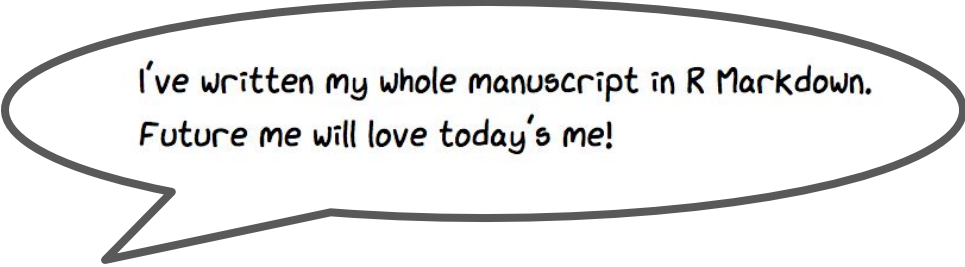
Viridis colors



Magma colors



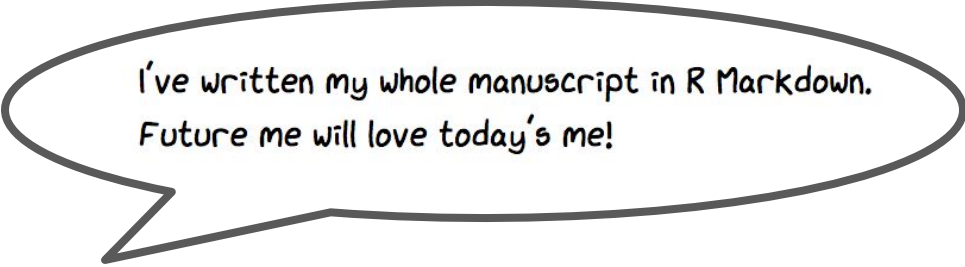
Is this you?




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?

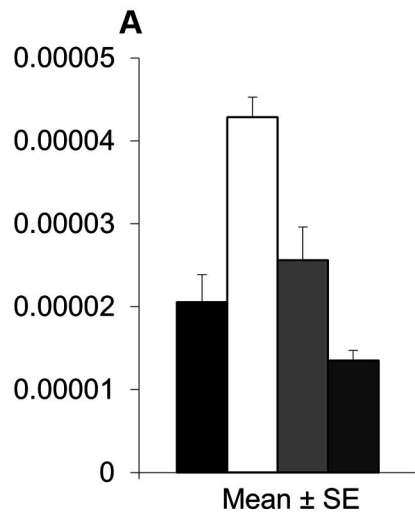
I've written my whole manuscript in R Markdown.
Future me will love today's me!

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

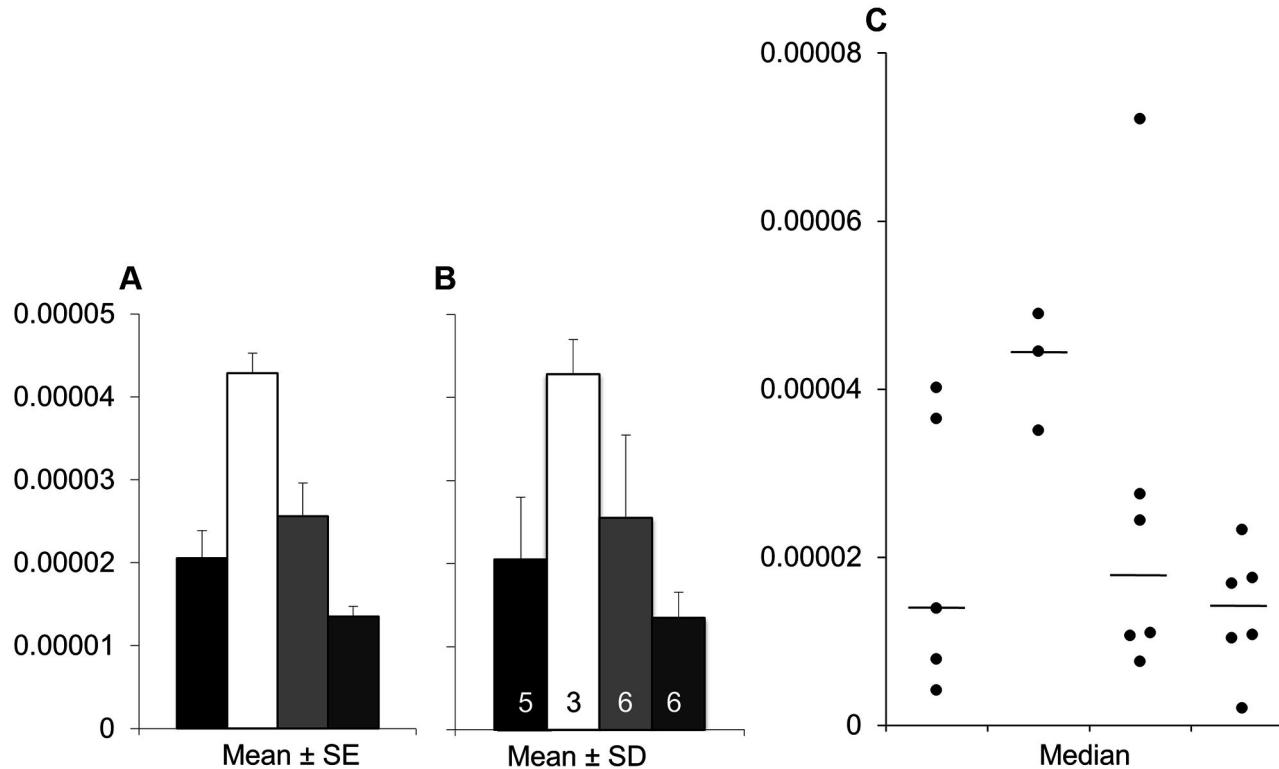
\$%*!@#!*\$&%



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



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- Play first in the browser before committing to local use
- Connect Excel clickers with iPython notebook coders



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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



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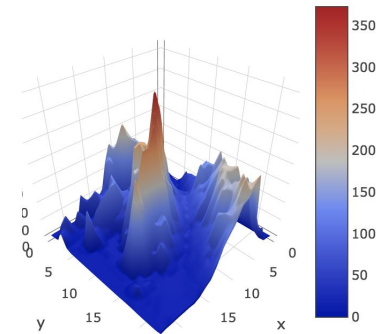
Recent progress

- 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
- 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/>.

You can help

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- Learn about progress and opportunities to help

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Supporting the open source community through the eLife Innovation Initiative

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Questions?

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