

Package ‘scientific’

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Type Package

Title Highly Customizable 'rmarkdown' Theme for Scientific Reporting

Version 2024.2

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Description Offers 'markdown' output formats designed with various scientific styles, allowing users to generate PDF and HTML outputs. The output has a contemporary appearance with vibrant visuals, providing numerous styles for effective highlighting. The package also includes additional features specifically tailored for front-page slides, enhancing the overall presentation and customization options. The package was created using the 'tufte' <<https://rstudio.github.io/tufte/>> package code as a starting point.

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URL <https://scientific.obianom.com>

BugReports <https://github.com/oobianom/scientific/issues>

Imports htmltools, knitr, rmarkdown, xfun

Suggests testthat

Encoding UTF-8

RoxygenNote 7.2.3

Config/testthat/edition 3

VignetteBuilder knitr

Language en-US

LazyData false

NeedsCompilation no

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Repository CRAN

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Description

Template for creating scientific handout

Usage

```
handout(  
  fig_width = 4,  
  fig_height = 2.5,  
  fig_crop = TRUE,  
  dev = "pdf",  
  highlight = "default",  
  ...  
)  
  
book(  
  fig_width = 4,  
  fig_height = 2.5,  
  fig_crop = TRUE,  
  dev = "pdf",  
  highlight = "default",  
  ...  
)  
  
html(...)  
  
newthought(text)  
  
margin_note(text, icon = "&#8853;")  
  
quote_footer(text)  
  
sans_serif(text)
```

Arguments

<code>fig_width</code>	Default width (in inches) for figures
<code>fig_height</code>	Default height (in inches) for figures
<code>fig_crop</code>	Whether to crop PDF figures with the command <code>pdftocrop</code> . This requires the tools <code>pdftocrop</code> and <code>ghostscript</code> to be installed. By default, <code>fig_crop = TRUE</code> if these two tools are available.
<code>dev</code>	Graphics device to use for figure output (defaults to pdf)

highlight	Syntax highlighting style passed to Pandoc. Supported built-in styles include "default", "tango", "pygments", "kate", "monochrome", "espresso", "zenburn", "haddock", and "breezedark". Two custom styles are also included, "arrow", an accessible color scheme, and "rstudio", which mimics the default IDE theme. Alternatively, supply a path to a '.theme' file to use a custom Pandoc style . Note that custom theme requires Pandoc 2.0+. Pass NULL to prevent syntax highlighting.
...	Other arguments to be passed to <code>pdf_document()</code> or <code>html_document()</code> (note you cannot use the <code>template</code> argument in <code>handout</code> or the <code>theme</code> argument in <code>html()</code> ; these arguments have been set internally)
text	A character string to be presented as a “new thought” (using small caps), or a margin note, or a footer of a quote
icon	A character string to indicate there is a hidden margin note when the page width is too narrow (by default it is a circled plus sign)

Details

`handout()` provides the PDF format

`html()` provides the HTML format based on the scientific CSS

`newthought()` can be used in inline R expressions in R Markdown

```
`r newthought(Some text)`
```

and it works for both HTML (`text`) and PDF (`\newthought{text}`) output.

`margin_note()` can be used in inline R expressions to write a margin note (like a sidenote but not numbered).

`quote_footer()` formats text as the footer of a quote. It puts text in `<footer></footer>` for HTML output, and after `\hfill` for LaTeX output (to right-align text).

`sans_serif()` applies sans-serif fonts to text.

Value

a PDF or HTML notebook output based on the R markdown document provided

Examples

```
## Not run:
# for Rmd to PDF
library(rmarkdown)
library(scientific)
rmdfile <- "input.Rmd"
rmarkdown::render(rmdfile,
  scientific::handout())

## End(Not run)
```

```
## Not run:  
# for Rmd to HTML  
library(rmarkdown)  
library(scientific)  
rmdfile <- "input.Rmd"  
rmarkdown::render(rmdfile,  
  scientific::html(  
    toc = TRUE,  
    toc_depth = 2))  
  
## End(Not run)  
newthought("In this section")
```

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