
stitch Documentation

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`stitch` is a library for making reproducible reports. It takes a markdown source file, executes the code chunks, captures the output, and stitches the output into the destination file.

Those familiar with `knitr` and `RMarkdown` will recognize it as a python clone of those great libraries. It's also heavily influenced by `knitpy`.

While `stitch` is written in python, it can be used for any of the dozens of `Jupyter kernels`.

Examples

See the [example site](#) for a side-by-side comparison of the markdown source and rendered output. Links to a more complicated document rendered in various formats is also provided.

Install

At the moment, the name `stitch` is taken on PyPI via an inactive project. You can install `stitch` from PyPI via

```
pip install knotr
```

I know, it's confusing. I've filed a claim for `stitch` on PyPI, but I think the people working that support queue are over-worked. Once that gets processed, I'll put it up on conda-forge as well. If you need a mnemonic, it's "I want knitr, but *not* the one written in *R*. Also I wanted to confuse R users. And knots are kind of like a buggy version of knits.

But to be clear the package name and command-line tool is `stitch`.

You'll also need to have a recent version of `pandoc`. Either use your system package manager, or use the `pypandoc` provided on conda-forge, which includes `pandoc`.

Usage

You write a markdown file, and include code chunks that look like

```
```{kernel_name, [chunk_name], **kwds}  
your code here
```
```

The `kernel_name` is required (see `jupyter kernelspec list`). The `chunk_name` is optional; it controls things like the name assigned to plots if they're saved to disk.

The supported keyword arguments are

- `eval`: bool, whether to execute the code chunk
- `echo`: bool, whether to include the input code chunk in the output

More options will be added.

The command-line interface is essentially the same as `pandocs`. For the most part you call

```
stitch input_file.md -o output_file.html
```

You can use `-t` for the output type, or infer it from the file extension of `-op`. All other options are passed through to `pandoc`.

`stitch` defines a few new options that control stitch-specific features

- `--no-standalone`
- `--no-self-contained`

What's in a Name?

The name `stitch` has a couple meanings. Like R's `knit`, we are taking a source document, executing code chunks, and knitting or stitching the output back into the document.

The second meaning is for `stitch` bringing together a bunch of great libraries, minimizing the work we have to do ourselves. `stitch` uses

- `Pandoc` markdown parsing and conversion to the destination output
- `jupyter`, specifically `jupyter-client` for managing kernels, passing code to kernels, and capturing the output
- `pandocfilters` for converting code-chunk output to pandoc's AST
- `py pandoc` for communicating with pandoc
- `click` for the command-line interface.

`stitch` itself is fairly minimal. The main tasks are

- processing code-chunk arguments
- passing the correct outputs from the jupyter kernel to `pandocfilters`
- assembling all the chunks of text, code, and output in a sensible way
- making things look nice

Contents:

4.1 API

4.1.1 Chunk Options

Code chunks are blocks that look like

```
```{kernel_name, [chunk_name], **kwargs}
code
```
```

The `kernel_name` is required, and `chunk_name` is optional. All parameters are separated by a comma.

kernel_name (*name: str*)

Name of the kernel to use for executing the code chunk. Required. See `jupyter kernelspec list`.

chunk_name (*chunk_name: str*)

Name for the chunk. Controls the filename for supporting files created by that chunk. Optional.

echo (*echo=True*)

whether to include the input-code in the rendered output. Default True.

eval (*eval=True*)

Whether to execute the code cell. Default True.

results (*s*)

str; how to display the results

- hide: hide the chunk output (but still execute the chunk)

width (*w*)

Width for output figure. See <http://pandoc.org/MANUAL.html#images>

Warning: This will probably change to `fig.width` in a future release.

height (*w*)

Height for output figure. See <http://pandoc.org/MANUAL.html#images>

Warning: This will probably change to `fig.height` in a future release.

4.2 Whatsnew

4.2.1 Version 0.3.3

- Included `default.css` in the source and binary distributions (GH26).
- Fixed not handling output from IPython's various `display` methods (GH27).

Indices and tables

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