Flex Knit

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Synopsis

This document presents the R-function `myknit` that enables dynamically changing the output files of the knit-process by using the undocumented `knit` yaml option.

Acknowledgement

The idea for manipulating yaml information in the way done here was found in a Stack Overflow article (option 2) authored by *mathematical-coffee*.

Introduction

When creating a new document I always have to try various versions of the yaml metadata block before I have a version that works for me. I need an easy way to keep these various inputs and outputs separated. The standard way (the `knit` button) always creates/overwrites the same output file. The `rmarkdown::render` function that is called as a result of clicking the button allows specifying a name for the output file. I found (see acknowledgement) a method to call this function with the undocumented (why?) `knit` yaml option. The `knit` option option specifies what is executed when the `knit` button is clicked: a direct call to `rmarkdown::render` or a user function.
I use the construction `knit:  (function (...) { source('myknit.r'); myknit(...) })` with the user written function `myknit`. The `myknit` function supports some additional options in the yaml metadata block and uses these to construct output names.

Functionality of the `myknit` function

By specifying additional options in the yaml metadata block the user can do one or more of the following things:

- specify an alternative name for the output file
- append a version indicator to the name of the output file
- force that the proper extension is given to the name of the output file (when not specified)
- create a file with the yaml metadata block that was specified with a name including the version indicator
- create a file with the yaml metadata block after processing with a name including the version indicator
- create a copy of the input file with a name including the version indicator

How to use the functionality of the `myknit` function

The `myknit` function is controlled by additional options in the yaml metadata block. To distinguish these options from the regular ones, they are prefixed with `hoqc_`. The additional options (with defaults in parentheses) are:

- `hoqc_output`   (`''`) : the name of the output document file
- `hoc_version`   (`''`) : suffix to be given to files to indicate version
- `hoqc_force_ext` (F) : should a filename extension be forced for document and yaml files? (TRUE or FALSE)
- `hoqc_yaml`   (`''`) : the name of file containing the original yaml metadata block
- `hoqc_yaml_new`   (`''`) : the name of file containing the new (modified) yaml metadata block

- `hoqc_rmd_in`   ('') : the name of file containing a copy of the input file
- `hoqc_rmd_out`   ('') : the name of file of the processed input file (just before rendering)

The output document will in principle be named `hoqc_output` if this is specified; otherwise it is named after the input file. If `hoqc_output` is specified without an extension and `hoqc_force_ext == T` then the output name will be given an extension of `pdf` or `html` depending on the (first encountered, not commented) document_type. `hoc_version` will be inserted after the basename of the file.

The orginal yaml metadata block will be written to an output file with name `hoqc_yaml` if this is specified. After reading the orginal yaml metadata block comment lines (beginning with `#`) are removed. The modified yaml metadata block will be written to an output file with name `hoqc_yaml_new` if this is specified. In both yaml file names the `hoc_version` will be inserted and a `txt` extension will be given when an extension was not specified and `hoqc_force_ext == T`.

A copy of the input file is created with name `hoqc_rmd_in` if this is specified. The `hoc_version` will be inserted and a `Rmd` extension will be given when an extension was not specified and `hoqc_force_ext == T`. When `hoqc_rmd_out` is specified the same holds for the processed input file that is used for the actual rendering.

In the modified metadata block the additional options are included (added if it already exists) in the `params` block. The name of the output files are expanded to the full path. The next paragraph shows how the `params` block can be used to include e.g. the yaml metadata block in the document by specifying `r params$hoqc_yaml` within backticks.

A usage scenario for working with LaTeX could be:

- set `hoqc_output` and `hoqc_copy` e.g. to `./output/flexkit` and `keep_tex` to `yes`
- set `hoc_version` to `_v1` and knit the file.
- set `hoc_version` to `_v2`, make some changes and knit again

Intermediate folders will be created if necessary.

In the `output` subfolder one now can find for both versions the saved inputs (`flexknit_v1.Rmd` and `flexknit_v2.Rmd`) and the corresponding `tex` files and `pdf` files (with version indicators). When something is not working as expected one can compare the various versions to see what is wrong. In this way I saw that the yaml options for `geometry` and `params` do not allow blanks between the options and the colon.

Listing of the yaml metadata blocks

As an example the original yaml (in D:/data/R/rmd_pdf_examples/output/yaml_v1.txt) for this document will be shown:

```
cat(paste0(readLines(params$hoqc_yaml),collapse = '\n'))
```

```
    ---
    title            : "Flex Knit"
    author           : "Han Oostdijk"
    date             : "file created : r format(Sys.time(), '%d%b%Y')"
    output           :
      pdf_document    :
        keep_tex      : yes
      # html_document   :
      #   keep_md      : yes
    classoption      : portrait # or landscape
    # do not place blanks between geometry and ':' or the option will be ignored !
    geometry: [
      left=0.6in ,
      right=0.5in ,
```

```
      top=1in ,
      bottom=1in
      ]
    # comment following lines if you do not want to use sans serif monotype
    header-includes    : [
      '\usepackage[scaled]{DejaVuSansMono}' ,
      '\renewcommand*{\familydefault}{\sfdefault}'
      ]
    urlcolor            : blue
    hoqc_version        : _v1
    # hoqc_ filenames can have a path
    hoqc_output         : './output/Flex Knit'
    hoqc_yaml           : ./output/yaml
    hoqc_yaml_new       : ./output/yaml_new
    hoqc_rmd_in         : ./output/flexknit.Rmd
    hoqc_force_ext      : yes
    knit                : (function (...) { source('myknit.R'); myknit(...) })
    # do not place blanks between params and ':' !
    params:
      parm1             : myfirstparm
    ---
```

and also the modified yaml (in D:/data/R/rmd_pdf_examples/output/yaml_new_v1.txt) :

```
cat(paste0(readLines(params$hoqc_yaml_new),collapse = '\n'))
```

```
    ---
    title               : "Flex Knit"
    author              : "Han Oostdijk"
    date                : "file created : r format(Sys.time(), '%d%b%Y')"
    output              :
      pdf_document      :
        keep_tex        : yes
    classoption         : portrait # or landscape
    geometry: [
      left=0.6in ,
      right=0.5in ,
      top=1in ,
      bottom=1in
      ]
    header-includes     : [
      '\usepackage[scaled]{DejaVuSansMono}' ,
      '\renewcommand*{\familydefault}{\sfdefault}'
      ]
    urlcolor            : blue
    knit                : (function (...) { source('myknit.R'); myknit(...) })
    params:
      parm1             : myfirstparm
      hoqc_output     : 'D:/data/R/rmd_pdf_examples/output/Flex Knit_v1.pdf'
      hoqc_yaml       : 'D:/data/R/rmd_pdf_examples/output/yaml_v1.txt'
      hoqc_yaml_new   : 'D:/data/R/rmd_pdf_examples/output/yaml_new_v1.txt'
      hoqc_force_ext  : 'yes'
      hoqc_version    : '_v1'
```

```
  hoqc_rmd_in    : 'D:/data/R/rmd_pdf_examples/output/flexknit_v1.Rmd'
---
```

Session Info

```
R version 3.4.1 (2017-06-30)
Platform: x86_64-w64-mingw32/x64 (64-bit)
Running under: Windows 10 x64 (build 16299)

Matrix products: default

locale:
[1] LC_COLLATE=English_United States.1252
[2] LC_CTYPE=English_United States.1252
[3] LC_MONETARY=English_United States.1252
[4] LC_NUMERIC=C
[5] LC_TIME=English_United States.1252

attached base packages:
[1] stats     graphics  grDevices utils     datasets  methods   base

other attached packages:
[1] stringr_1.3.0 magrittr_1.5  fs_1.1.0

loaded via a namespace (and not attached):
 [1] compiler_3.4.1   backports_1.1.2  rprojroot_1.3-2  htmltools_0.3.6
 [5] tools_3.4.1      yaml_2.1.16      Rcpp_0.12.14     rmarkdown_1.8.10
 [9] stringi_1.1.6    knitr_1.20       digest_0.6.14    evaluate_0.10.1
```