

# STAT 545A

## Class meeting #3

### Wednesday, September 11, 2013

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Department of Statistics and Michael Smith Laboratories



any questions from tutorial re: Basic care and feeding of data in R?

HW #1 “Dry Run”. Due as soon as possible. Publish 1 page of something -- practically anything -- to Rpubs, then share that link w/ rest of the class. Instructions/”assignment” is posted but will get some editing. Content not important.

HW #2 “Dry Run 2 + Hello Gapminder”. Due 9:30am Monday Sept. 16. Perform basic data intake and smell testing for the Gapminder data. Write it up via R Markdown. Compile, publish, share. Content can be drawn from tutorials, your job is to pull together. Instructions coming.

today

data analysis vs programming vs developing

combo lecture and hands on computing:

R objects, beyond the data.frame

students will complete on their own

R Markdown

three different modes of working with R:

data analysis

programming

developing

three different modes of working with R:

data analysis ← I am mostly trying to prepare  
you for this.

programming

developing

three different modes of working with R:

data analysis

programming

developing

But we need to talk .... about  
some things more related to this.



hands-on computing phase here

# JB's typology of R objects, i.e. "flavors"

a simple view of simple R objects that will get you pretty far

Simple view	Technically correct R view		
	mode	class	typeof
character	character	character	character
logical	logical	logical	logical
numeric	numeric	integer or numeric	integer or double
factor	numeric	factor	integer



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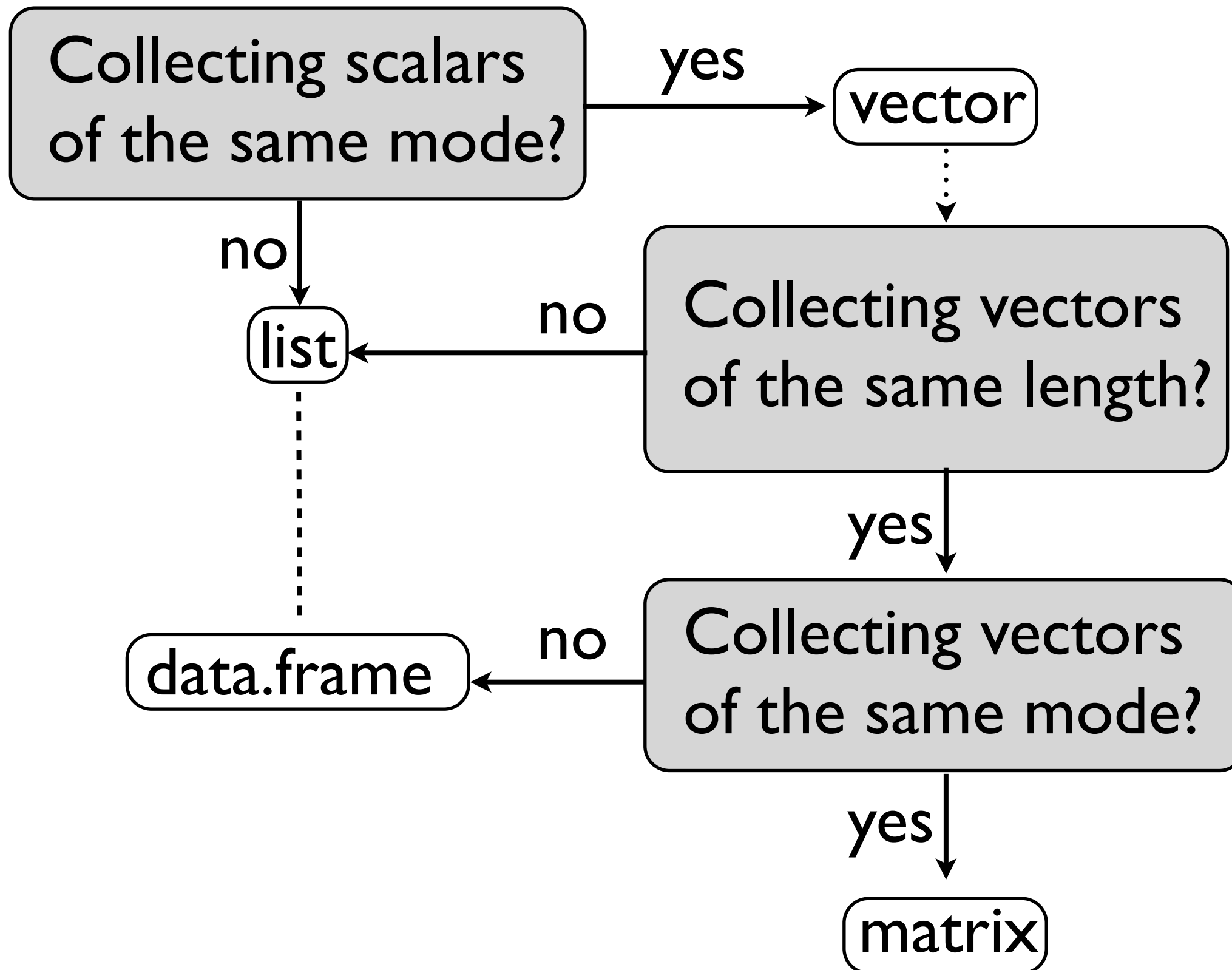
# JB's typology of R objects, i.e. "flavors"

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We will spend some time talking about factors in a future class.

# “Simple view” of data collections



# weak links in the chain: process, packaging and presentation



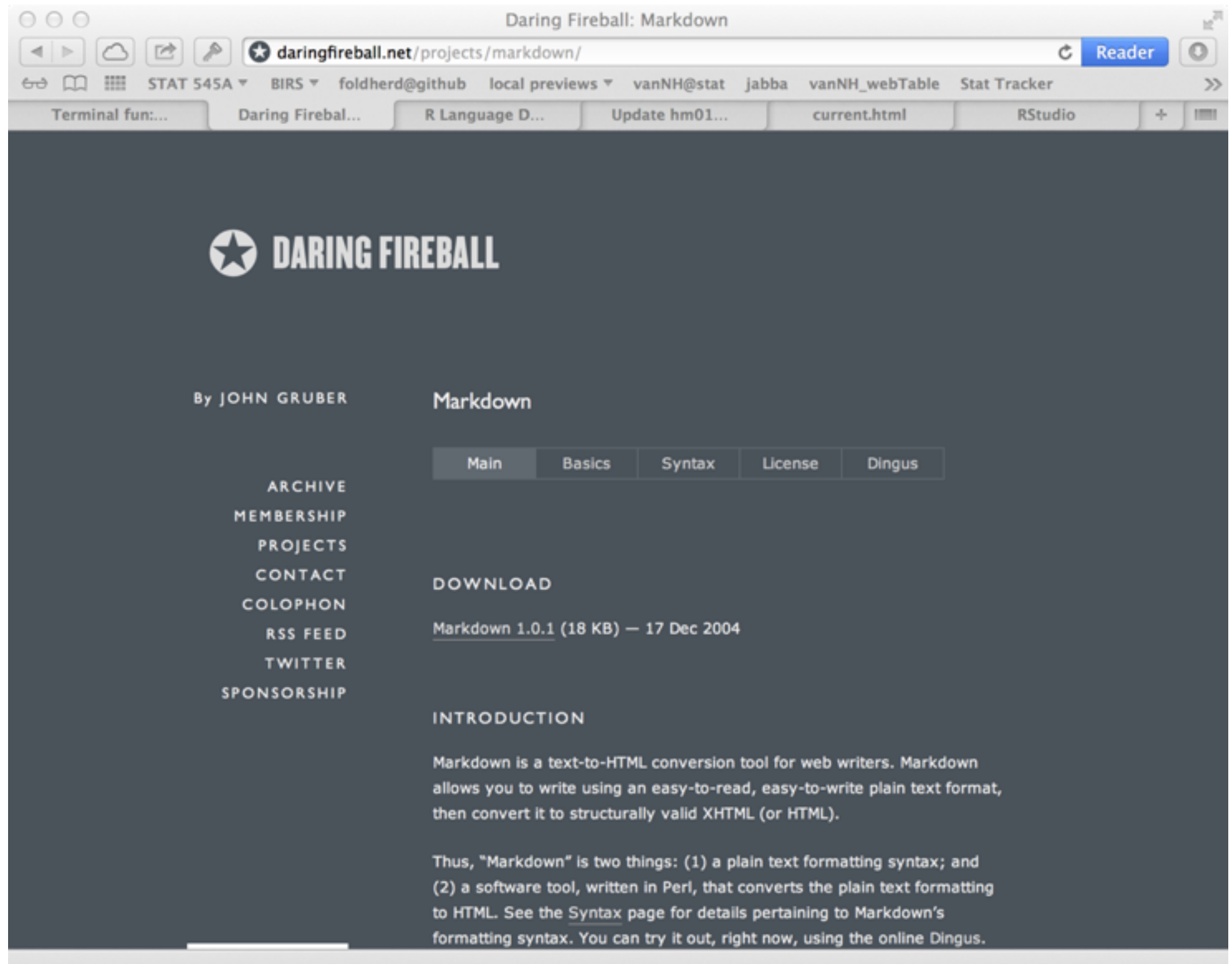
1. **Build your communication skills.** Unless you live in a plastic bubble, you are going to need to work with other people. You will be given tasks by other people, collaborate with other people to achieve those tasks, and ultimately have to report the results of your work to other people. You need to be able to speak clearly and concisely, listen carefully, write well (and quickly), and give informative and interesting presentations. Contrary to popular belief *the person who learns to do these things well will advance farther than someone who has better technical capabilities but poor communication skills!* Management usually can't tell the difference between a good statistician and a great one, but they can see immediately who communicates their results well and who does so poorly. Unfortunately, most university environments stress working alone and in isolation, completely the opposite of what life will be like on the other side of graduation.

**You need to take actions to ensure that your communication skills are sharp.** These actions can include: (i) Taking a writing class, especially one that stresses technical writing, which has a completely different flavour from essay writing; (ii) Taking a class in verbal communication, and in particular one that covers the fine art of making and delivering presentations; (iii) Taking business courses, especially those in business communication and organizational structure and behaviour, so that you can better understand your audience and learn to arrange your communications accordingly; (iv) Seeking out courses that expressly advertise group project work and/or presentations, even (*especially!*) if these things scare you. All of our speakers indicate that anything that you can do to practice your communication skills will have a positive effect on your employability and advancement.

Excerpt from “Real Advice from Real People”  
by Tom Loughin, Statistical Society of Canada  
Liaison, Vol. 22 No. 4 (November 2008).



<http://daringfireball.net/projects/markdown/>



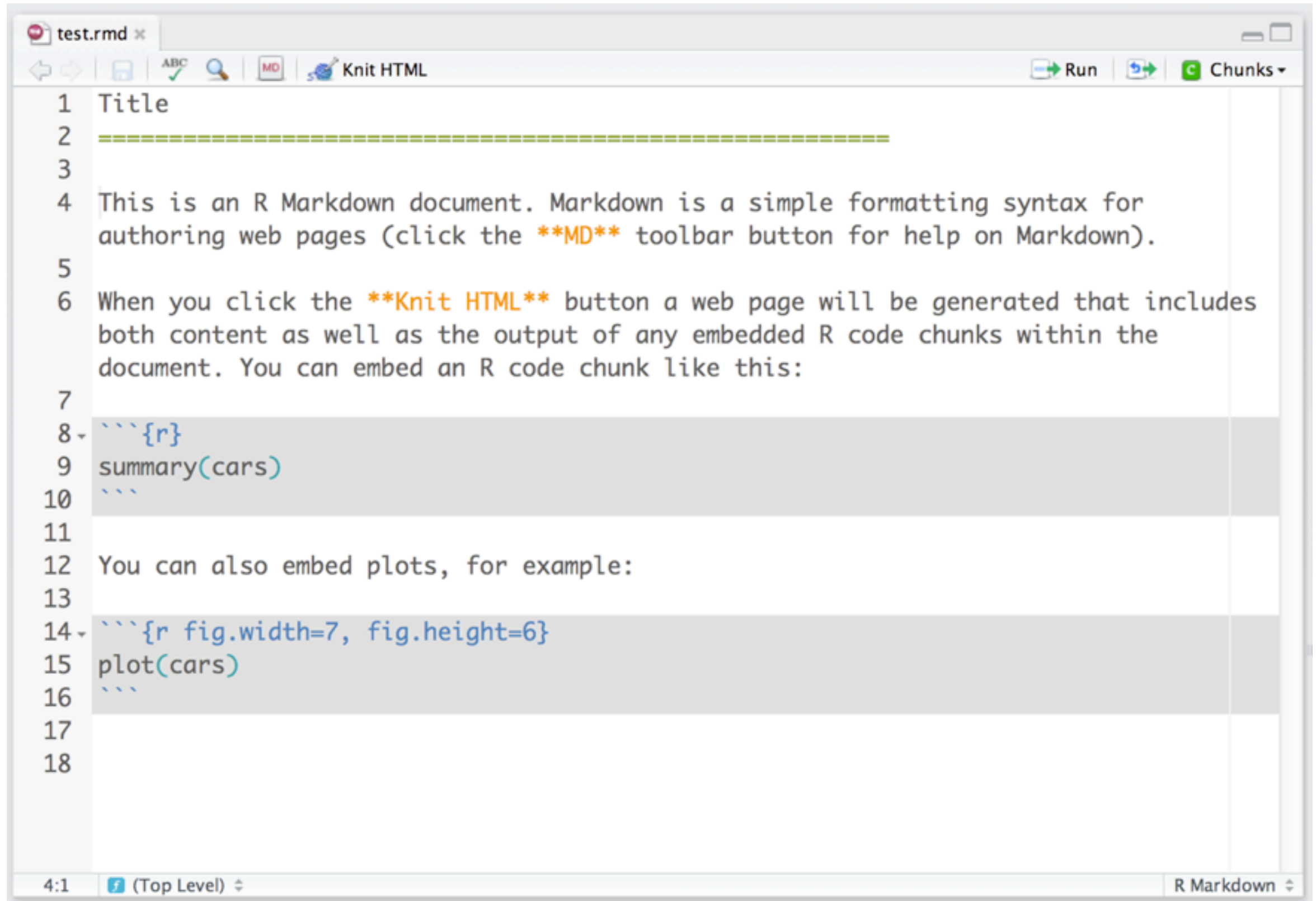
[http://www.rstudio.com/ide/docs/r\\_markdown](http://www.rstudio.com/ide/docs/r_markdown)

[http://www.rstudio.com/ide/docs/authoring/using\\_markdown](http://www.rstudio.com/ide/docs/authoring/using_markdown)

<http://yihui.name/knitr/>

# File > New > R Markdown

## Save As ... test.rmd



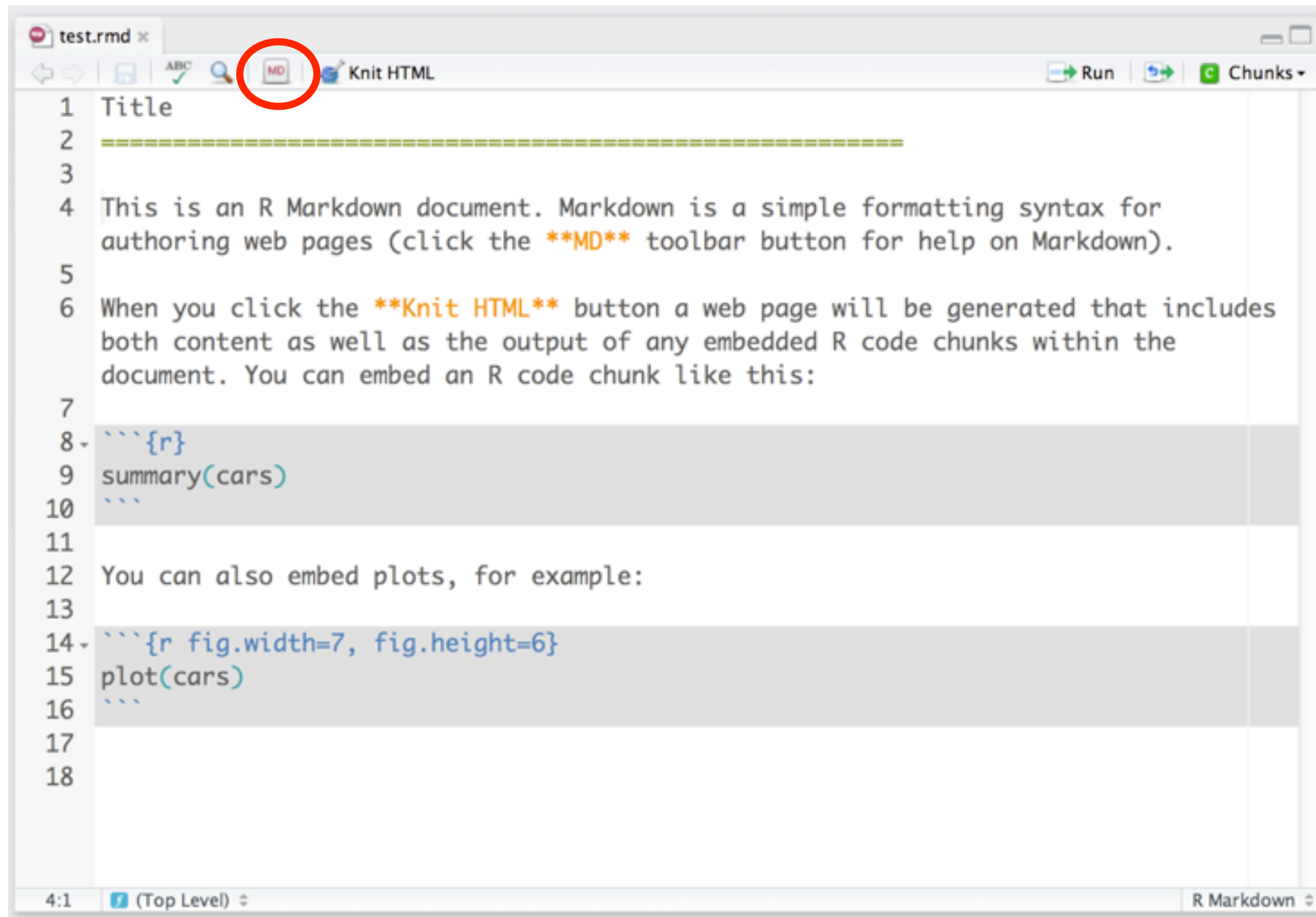
The screenshot shows an R Markdown editor window with a toolbar at the top containing icons for undo, redo, save, spell check, search, markdown, and knit HTML. On the right side of the toolbar are 'Run' and 'Chunks' buttons. The document content is as follows:

```
1 Title
2 =====
3
4 This is an R Markdown document. Markdown is a simple formatting syntax for
  authoring web pages (click the **MD** toolbar button for help on Markdown).
5
6 When you click the **Knit HTML** button a web page will be generated that includes
  both content as well as the output of any embedded R code chunks within the
  document. You can embed an R code chunk like this:
7
8 ```{r}
9 summary(cars)
10 ```
11
12 You can also embed plots, for example:
13
14 ```{r fig.width=7, fig.height=6}
15 plot(cars)
16 ```
17
18
```

The status bar at the bottom indicates the cursor is at line 4, column 1, and the document is at the top level. The file type is identified as R Markdown.

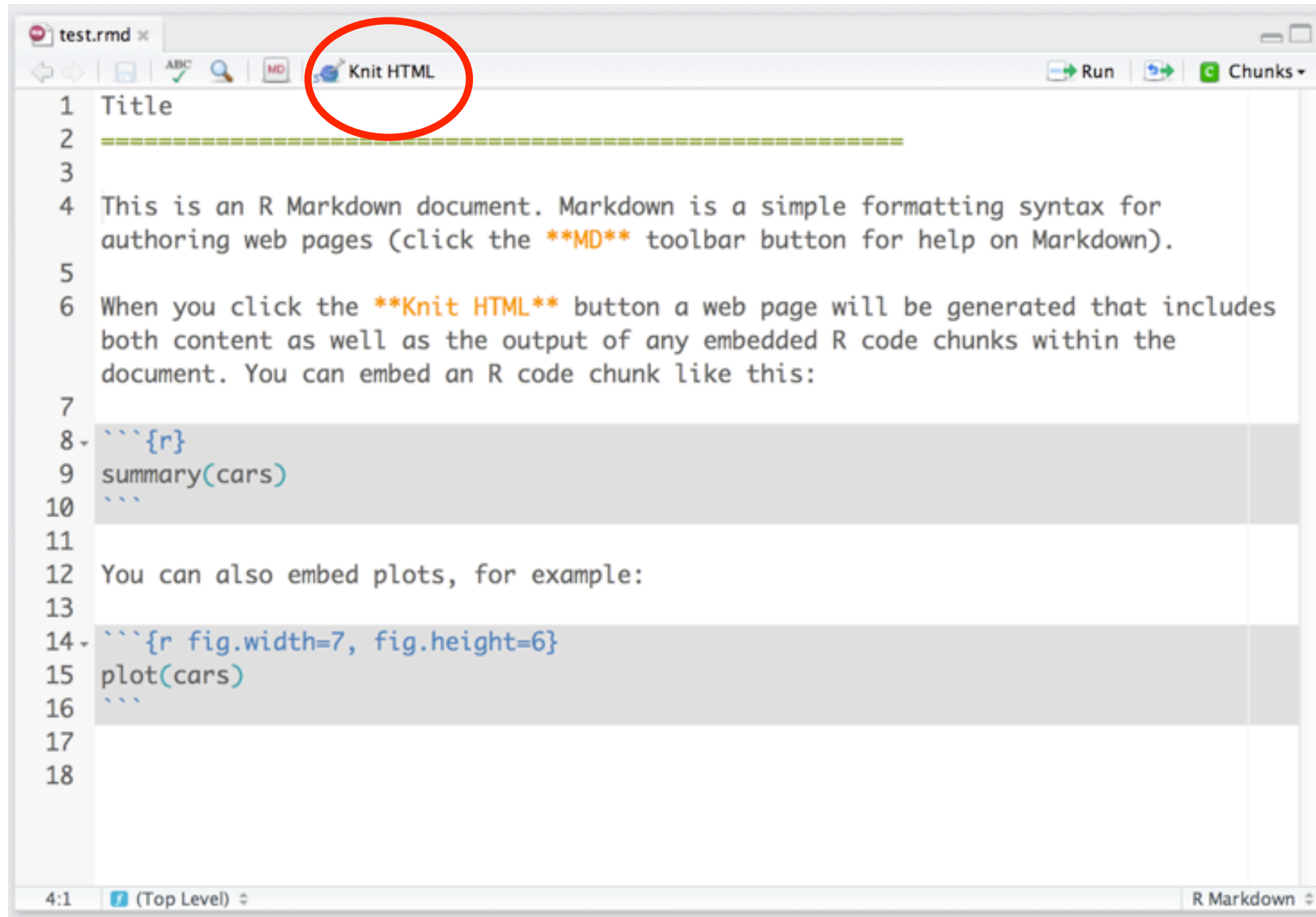


# Displays Markdown Quick Reference in the Help tab/pane



# File > New > R Markdown

## Save As ... test.rmd



## Knit HTML.

Yes this can be accomplished outside of RStudio, using knitr functions at the command line, so we are not creating unhealthy dependency on RStudio.

RStudio: Preview HTML

Preview: ~/teaching/STAT545A/2013\_STAT545A/test.html | Log | Save As | **Publish** | Find

## Title

This is an R Markdown document. Markdown is a simple formatting syntax for authoring web pages (click the **MD** toolbar button for help on Markdown).

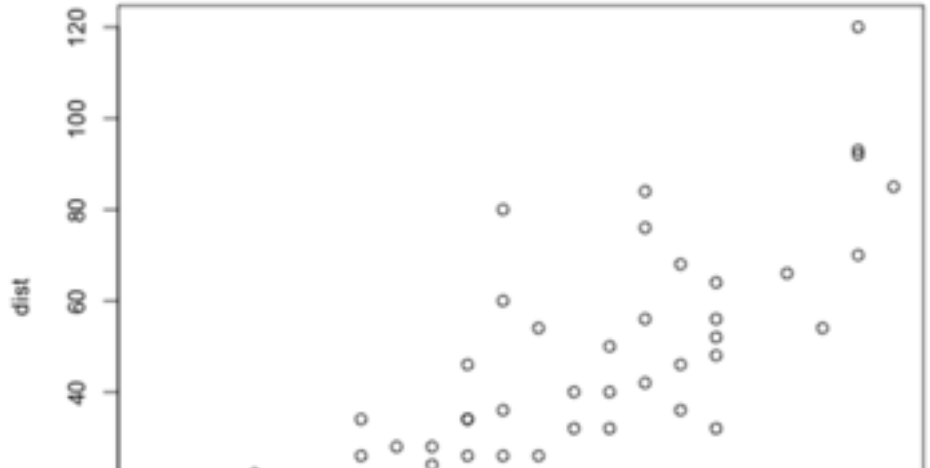
When you click the **Knit HTML** button a web page will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
summary(cars)
```

	speed	dist
## Min.	: 4.0	Min. : 2
## 1st Qu.	:12.0	1st Qu.: 26
## Median	:15.0	Median : 36
## Mean	:15.4	Mean : 43
## 3rd Qu.	:19.0	3rd Qu.: 56
## Max.	:25.0	Max. :120

You can also embed plots, for example:

```
plot(cars)
```



A scatter plot showing the relationship between speed (x-axis) and distance (y-axis) for the 'cars' dataset. The x-axis ranges from 0 to 25, and the y-axis ranges from 0 to 120. The plot shows a positive correlation, with distance increasing as speed increases. The data points are represented by open circles.

This is where you'll need that RPub account.

## What's actually happening?

<input type="checkbox"/>	 test.html	41.3 KB	Sep 10, 2013, 11:42 PM
<input type="checkbox"/>	 test.md	817 bytes	Sep 10, 2013, 11:42 PM
<input type="checkbox"/>	 test.rmd	542 bytes	Sep 10, 2013, 11:37 PM

You author this. You've seen it already.

<input type="checkbox"/>	 test.html	41.3 KB	Sep 10, 2013, 11:42 PM
<input type="checkbox"/>	 test.md	817 bytes	Sep 10, 2013, 11:42 PM
<input type="checkbox"/>	 test.rmd	542 bytes	Sep 10, 2013, 11:37 PM

knit() function from knitr package does R Markdown to Markdown conversion

R chunks are run and replaced by their output.  
Yes that's oversimplified but you get the idea.

# test.rmd

Title

=====

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```
```{r}
summary(cars)
```
```

You can also embed plots, for example:

```
```{r fig.width=7, fig.height=6}
plot(cars)
```
```

# test.md

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## 3rd Qu.:19.0      3rd Qu.: 56
## Max.    :25.0      Max.    :120
```
```

You can also embed plots, for example:

```
```r
plot(cars)
```
```

```
![plot of chunk unnamed-chunk-2](figure/unnamed-chunk-2.png)
```

|                          |   |           |                        |
|--------------------------|---|-----------|------------------------|
| <input type="checkbox"/> |  test.html | 41.3 KB   | Sep 10, 2013, 11:42 PM |
| <input type="checkbox"/> |  test.md   | 817 bytes | Sep 10, 2013, 11:42 PM |
| <input type="checkbox"/> |  test.rmd  | 542 bytes | Sep 10, 2013, 11:37 PM |

markdownToHTML() function from markdown package does  
Markdown to HTML conversion

Pandoc, the Swiss army knife of document conversion, is  
another option for Markdown to HTML conversion and,  
importantly, for Markdown to PDF conversion.

You can set things up so that RStudio and knitr use Pandoc  
for the second conversion step.

# test.md

Title

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## Max.   :25.0      Max.    :120
```
```

You can also embed plots, for example:

```
```r
plot(cars)
```
```

![plot of chunk unnamed-chunk-2](figure/unnamed-chunk-2.png)

```
page-break-inside: avoid;

pre {
  word-wrap: break-word;
}

</style>
```

# test.html

</head>

<body>  
<h1>Title</h1>

<p>This is an R Markdown document. Markdown is a simple formatting syntax for authoring web pages (click the **MD** toolbar button for help on Markdown).</p>

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<pre><code class="r">summary(cars)
</code></pre>
```

```
<pre><code>##           speed           dist
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## 1st Qu.:12.0      1st Qu.: 26
## Median :15.0      Median : 36
## Mean   :15.4      Mean    : 43
## 3rd Qu.:19.0      3rd Qu.: 56
## Max.   :25.0      Max.    :120
</code></pre>
```

<p>You can also embed plots, for example:</p>

```
<pre><code class="r">plot(cars)
</code></pre>
```

<p><img src="data:image/png;base64,iVBORw0KGgoAAAANSUUhEUgAAAFgAAAGwCAYAAABFI3d+AAAEJGldQ1BJQ0MgUHJvZmlsZQAABGFVd9v21QUPolVUqQWPyBYR4eKxa9VUlu5GxqtxgZJk6XtShal6dgqJOQ6N4mpGwfb6baqT3uBNwb8AUDZAw9IPCENBmJ72fbAtElThyqqSUh76MQPISbtBVXhu3ZiJ1PEXPX6yznfOec7517bRD1fabWaGVWIlquunc8klZOnFpSeTYrSs9RLA9Sr6U4tkcvNEi7BFffO6+EdigjL7ZHu/k72I796i9zRiSJPwG4VHX0Z+AxRzNRrtksUvwf7+Gm3BtzzHPDTNgQCqwKXfZwSeNHHJz10IT8JjtAq6xWtCLwGPLzYZi+3YV8DGMiT4VVuG7oiZpGzrZJhcs/hL49xtzH/Dy6bdfTsXYNY+5yluWO4D4neK/ZUvok/17X0HPBLsF+vuUlhfwX4j/rSfAJ4H1H0qZJ9dN7nR19frRTeBt4Fe9FwpwtN+2p1MXscGLHR9SXrmMgjONd1ZxKzpBeA71b4tNhj6JGoyFNp4GHgwUp9qplfmnFW5oTdy7NamcwCI49kv6fN5IAHgD



# Which would you rather read and write?

## test.rmd

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```
```



```
page-break-inside: avoid;
}
pre {
  word-wrap: break-word;
}
</style>
```

## test.html

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</code></pre>
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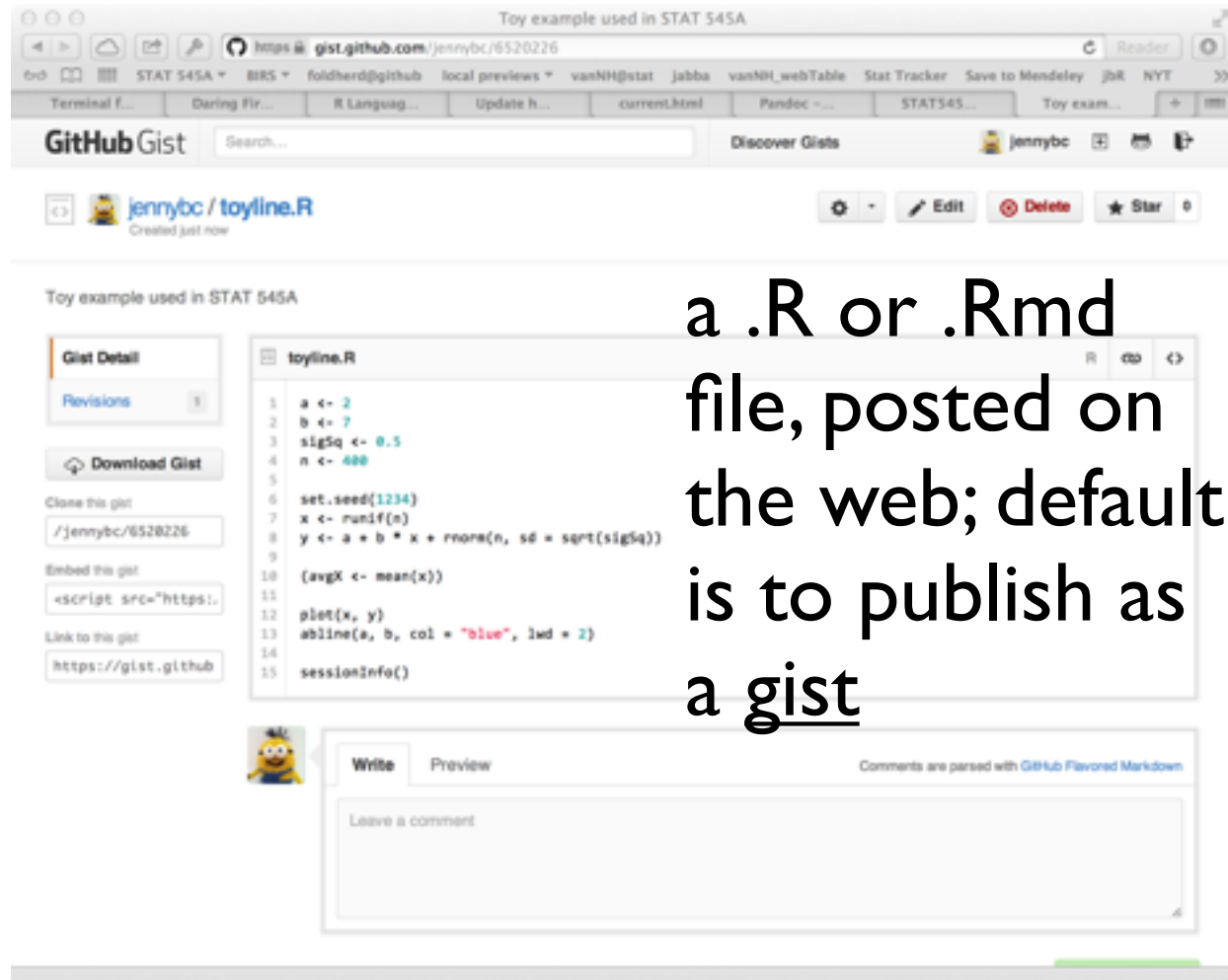


`knit2html()` function from knitr package does R Markdown to HTML conversion, i.e. hides the two separate steps

RStudio's Knit HTML button does practically the same thing

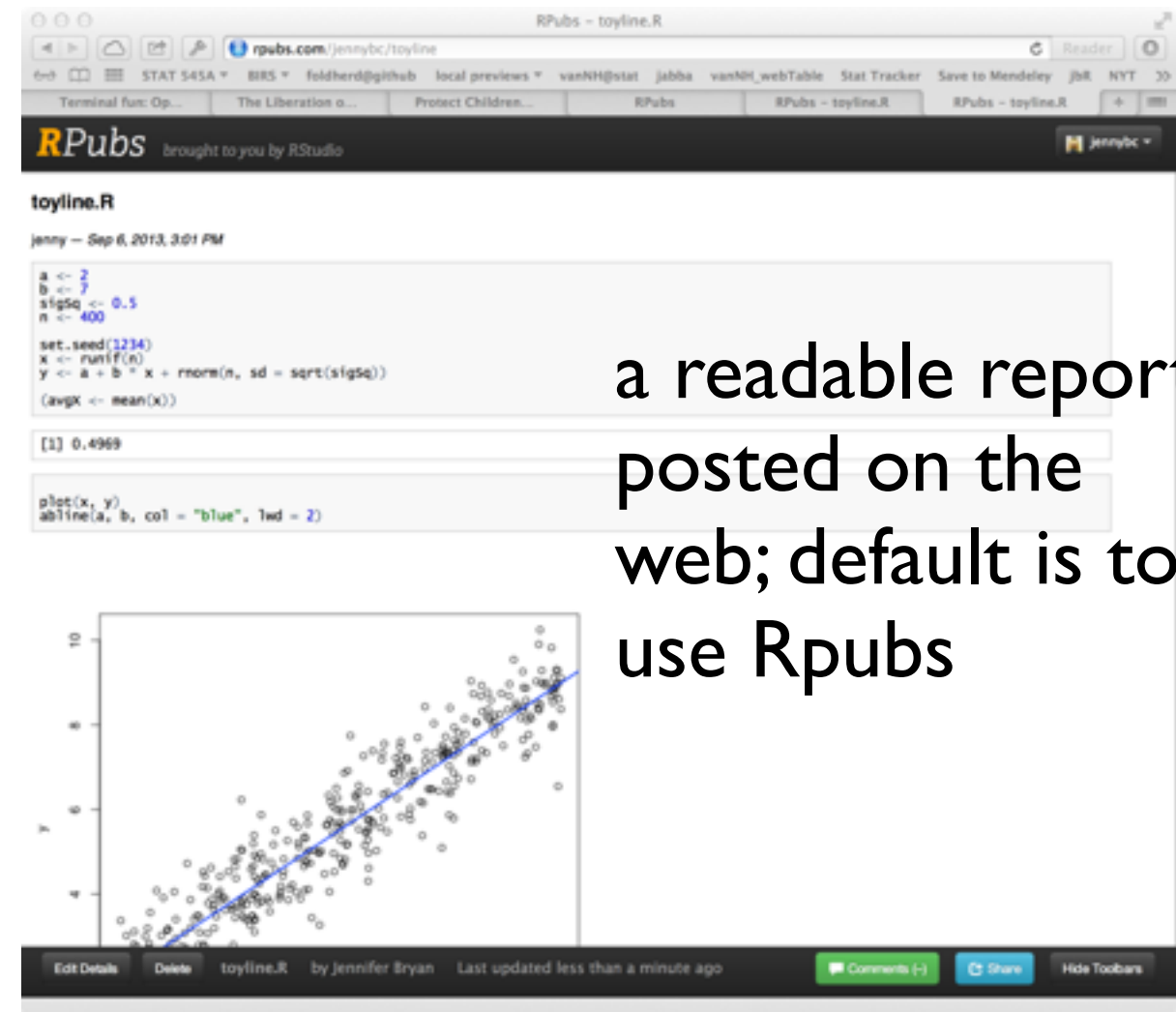
Admire, publish, etc. as you did before with the “notebook” you compiled from a plain .R script.

# What a finished piece of coursework will look like:



a .R or .Rmd  
file, posted on  
the web; default  
is to publish as  
a gist

+



a readable report,  
posted on the  
web; default is to  
use Rpubs

+

```
21 Please add your link in this bulleted list:
22
23 * Jenny Bryan: toylne \[script\]\(https://gist.github.com/jennybc/6520226\) | \[report\]\(http://rpubs.com/jennybc/toylne\)
24 * Matthew Gingerich: \[TestFile\]\(http://rpubs.com/majugi/TestFile\)
25 * Justin Chu: \[toylneTest\]\(http://rpubs.com/cjustin/8316\)
```

Please add your link in this bulleted list:

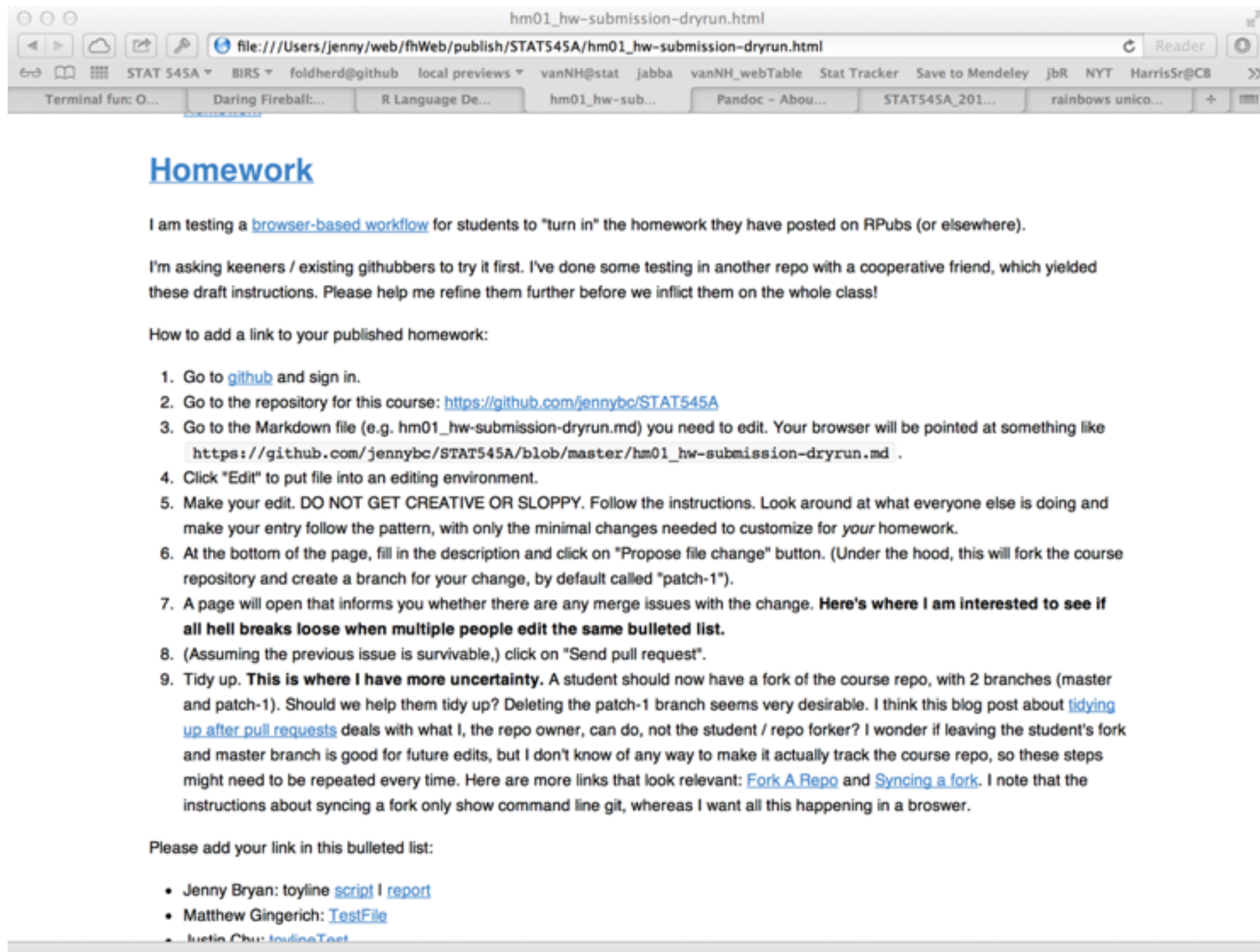
- Jenny Bryan: toylne [script](https://gist.github.com/jennybc/6520226) | [report](http://rpubs.com/jennybc/toylne)
- Matthew Gingerich: [TestFile](http://rpubs.com/majugi/TestFile)
- Justin Chu: [toylneTest](http://rpubs.com/cjustin/8316)

a line in course  
repository  
giving your  
name and links

=

# Instructions for how to submit coursework.

## Draft exists. Will be revised today.



The screenshot shows a web browser window with the title 'hm01\_hw-submission-dryrun.html'. The address bar shows the file path: 'file:///Users/jenny/web/fhWeb/publish/STAT545A/hm01\_hw-submission-dryrun.html'. The browser has several tabs open, including 'STAT 545A', 'BIRS', 'foldherd@github', 'local previews', 'vanNH@stat', 'jabba', 'vanNH\_webTable', 'Stat Tracker', 'Save to Mendeley', 'jbR', 'NYT', and 'HarrisSr@CB'. The main content area has a blue heading 'Homework'. Below the heading, the text reads: 'I am testing a [browser-based workflow](#) for students to "turn in" the homework they have posted on RPubS (or elsewhere). I'm asking keeners / existing githubbers to try it first. I've done some testing in another repo with a cooperative friend, which yielded these draft instructions. Please help me refine them further before we inflict them on the whole class! How to add a link to your published homework: 1. Go to [github](#) and sign in. 2. Go to the repository for this course: <https://github.com/jennybc/STAT545A> 3. Go to the Markdown file (e.g. hm01\_hw-submission-dryrun.md) you need to edit. Your browser will be pointed at something like `https://github.com/jennybc/STAT545A/blob/master/hm01_hw-submission-dryrun.md` . 4. Click "Edit" to put file into an editing environment. 5. Make your edit. DO NOT GET CREATIVE OR SLOPPY. Follow the instructions. Look around at what everyone else is doing and make your entry follow the pattern, with only the minimal changes needed to customize for *your* homework. 6. At the bottom of the page, fill in the description and click on "Propose file change" button. (Under the hood, this will fork the course repository and create a branch for your change, by default called "patch-1"). 7. A page will open that informs you whether there are any merge issues with the change. **Here's where I am interested to see if all hell breaks loose when multiple people edit the same bulleted list.** 8. (Assuming the previous issue is survivable,) click on "Send pull request". 9. Tidy up. **This is where I have more uncertainty.** A student should now have a fork of the course repo, with 2 branches (master and patch-1). Should we help them tidy up? Deleting the patch-1 branch seems very desirable. I think this blog post about [tidying up after pull requests](#) deals with what I, the repo owner, can do, not the student / repo forker? I wonder if leaving the student's fork and master branch is good for future edits, but I don't know of any way to make it actually track the course repo, so these steps might need to be repeated every time. Here are more links that look relevant: [Fork A Repo](#) and [Syncing a fork](#). I note that the instructions about syncing a fork only show command line git, whereas I want all this happening in a browser. Please add your link in this bulleted list: • Jenny Bryan: [toylne script](#) | [report](#) • Matthew Gingerich: [TestFile](#) • Justin Chu: [toylneTest](#)