

Introduction to R / R Studio

MATH 5305, Fall 2019
Department of mathematics, UTA

What is R?

- ❖ Programming Language S = developed at Bell Labs for statistics, simulation, graphics (Becker and Chambers, 1984)
 - ❖ S-PLUS: commercial implementation
 - ❖ R: implementation under GPL (GNU General Public License), open source.
- ❖ For R related tutorials and/or resources see the following links:
 - ❖ <https://cran.r-project.org/doc/manuals/R-intro.pdf>
 - ❖ <https://github.com/berkeley-scf>

Why R?

- ❖ R is widely used and has the widest statistical functionality. Wide usage helps to improve quality and reduce bugs.
- ❖ R is free and available on all major platforms
- ❖ Users add functionality via packages all the time, and access to fast growing number of packages
 - ❖ CRAN (>3500 packages) general data analysis
 - ❖ Bioconductor (>600 packages) bioscience data analysis
- ❖ As a scripting language, R is very powerful, flexible, and easy to use
- ❖ Efficient functions and data structures for data analysis; Powerful graphics.
- ❖ Standard for data mining and biostatistical analysis
- ❖ R is built on C and can call user-written and external C code and packages

Why not R?

- ❖ Other software is better than R at various tasks.
 - ❖ Python is very good for interacting with the operating system, and as a glue for tying together various applications/software in a workflow.
 - ❖ R can be much slower than compiled languages.
 - ❖ R's packages are only as good as the person who wrote them.
- ❖ Other programming languages
 - ❖ Python
 - ❖ Matlab/Octave
 - ❖ Julia
 - ❖ C/C++
 - ❖ SAS

R Studio

- ❖ R Studio allows the user to run R in a more user-friendly environment.
- ❖ Open-source and available at <https://www.rstudio.com>
- ❖ R Studio desktop can be installed in your local machine. R Studio server can be accessed using your web browser.
- ❖ Nice GUI for
 - ❖ File management/navigation, code highlighting, auto completion, R package management/development, version control, debugging, etc.

- ❖ R Markdown is an extension to the Markdown markup language that makes it easy to write HTML in a simple plain text format.

- ❖ <http://rmarkdown.rstudio.com/>

- ❖ R Sweave enables integration of R code into LaTeX documents. The purpose is to create dynamic reports, which can be updated automatically if data or analysis change.

- ❖ <http://yihui.name/knitr/>

- ❖ Version Control with Git and SVN

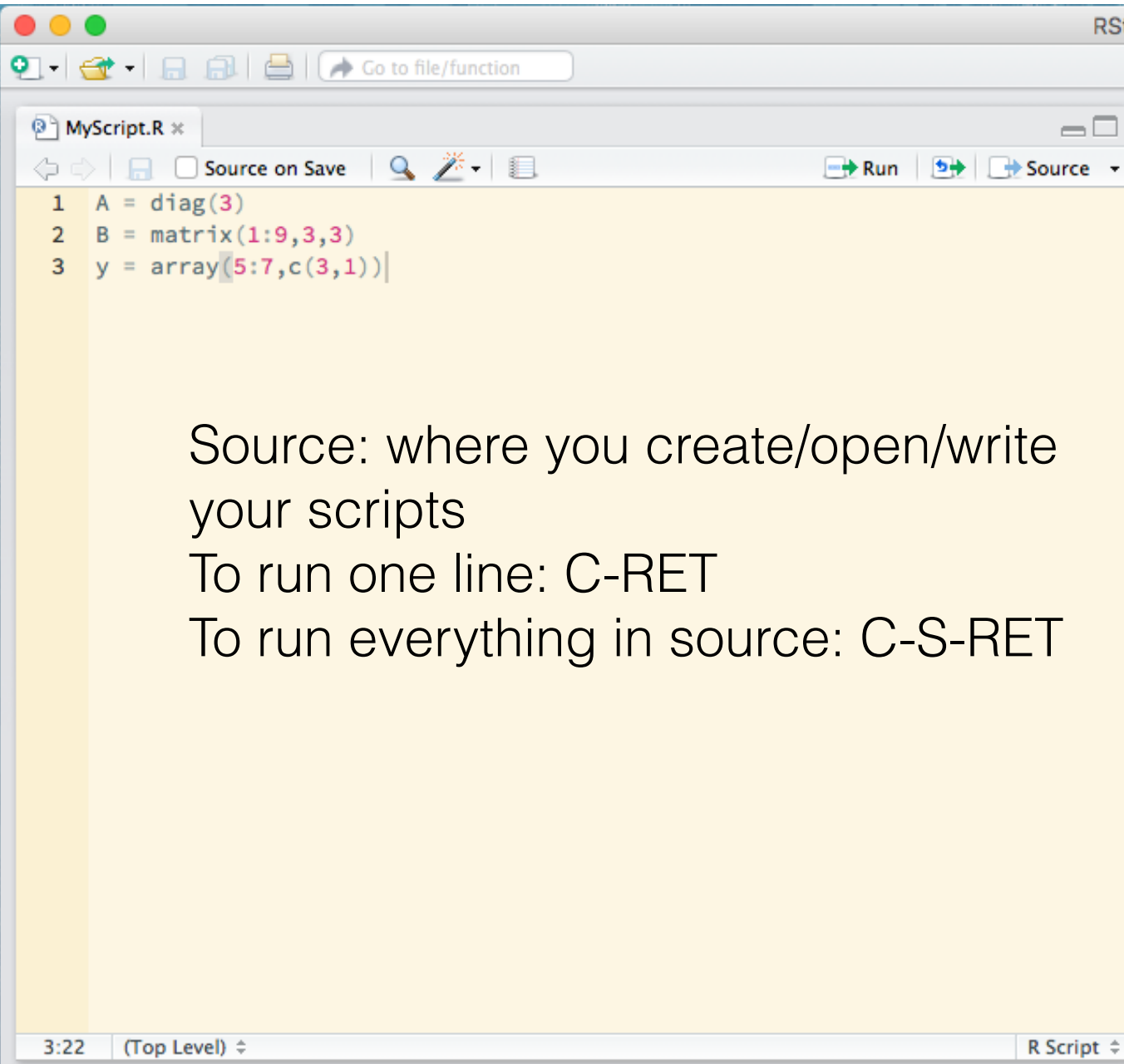
- ❖ <https://support.rstudio.com/hc/en-us/articles/200532077-Version-Control-with-Git-and-SVN/>

- ❖ R Studio cheat sheet

- ❖ <https://www.rstudio.com/resources/cheatsheets/>

- ❖ R for Matlab/Octave users

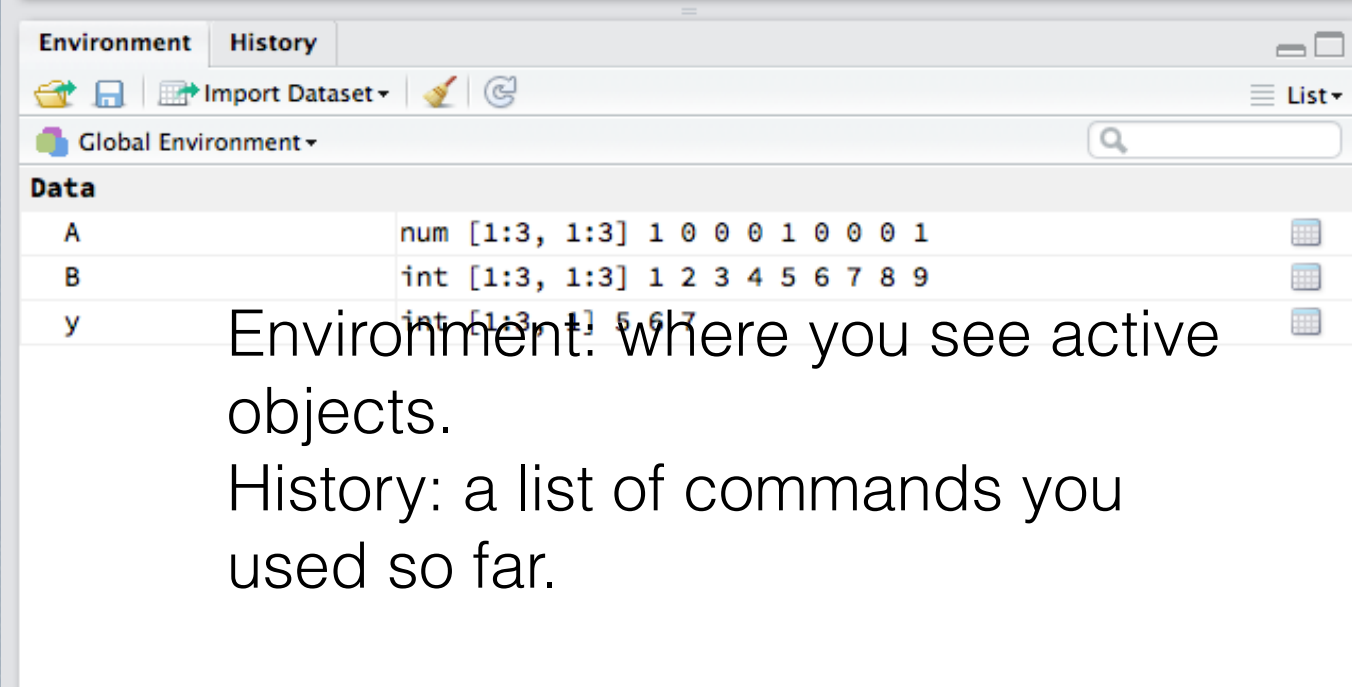
- ❖ <http://mathesaurus.sourceforge.net/octave-r.html>



Source: where you create/open/write your scripts

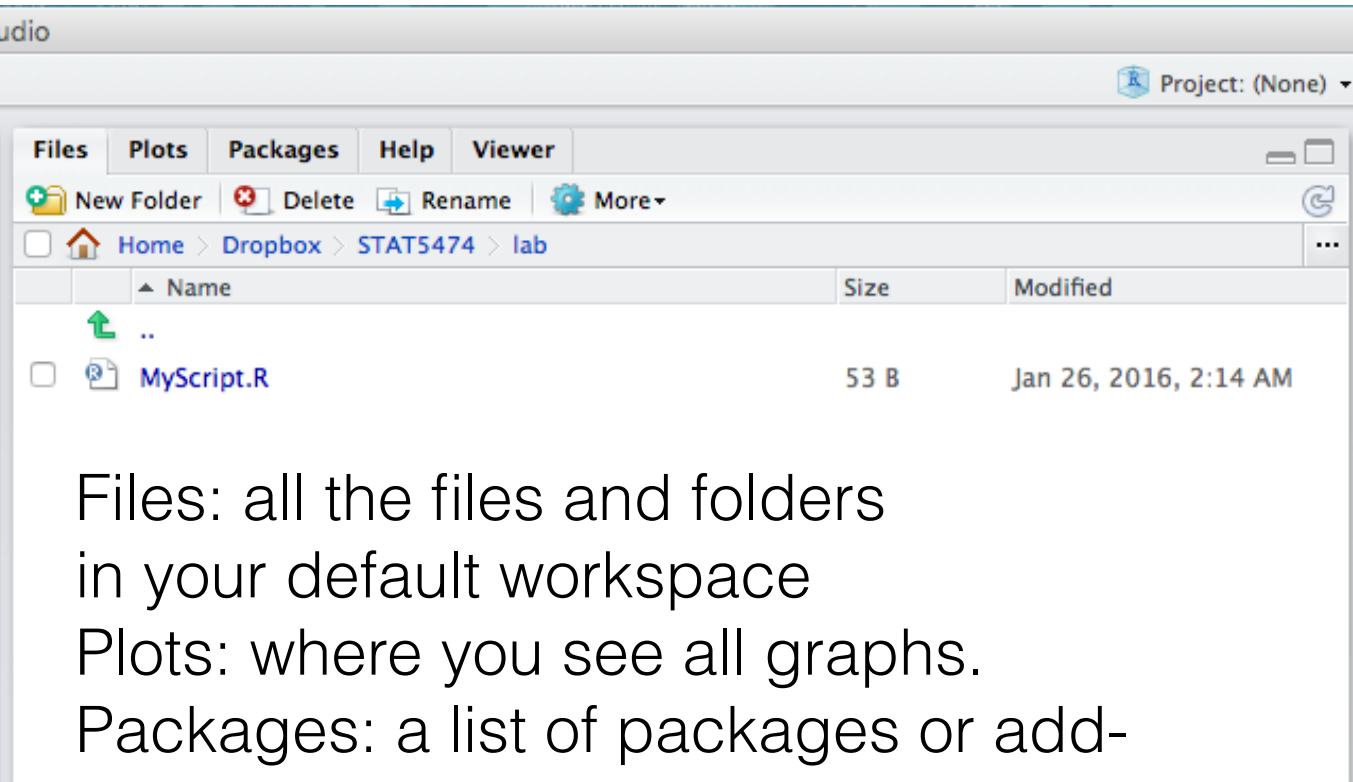
To run one line: C-RET

To run everything in source: C-S-RET



Environment: where you see active objects.

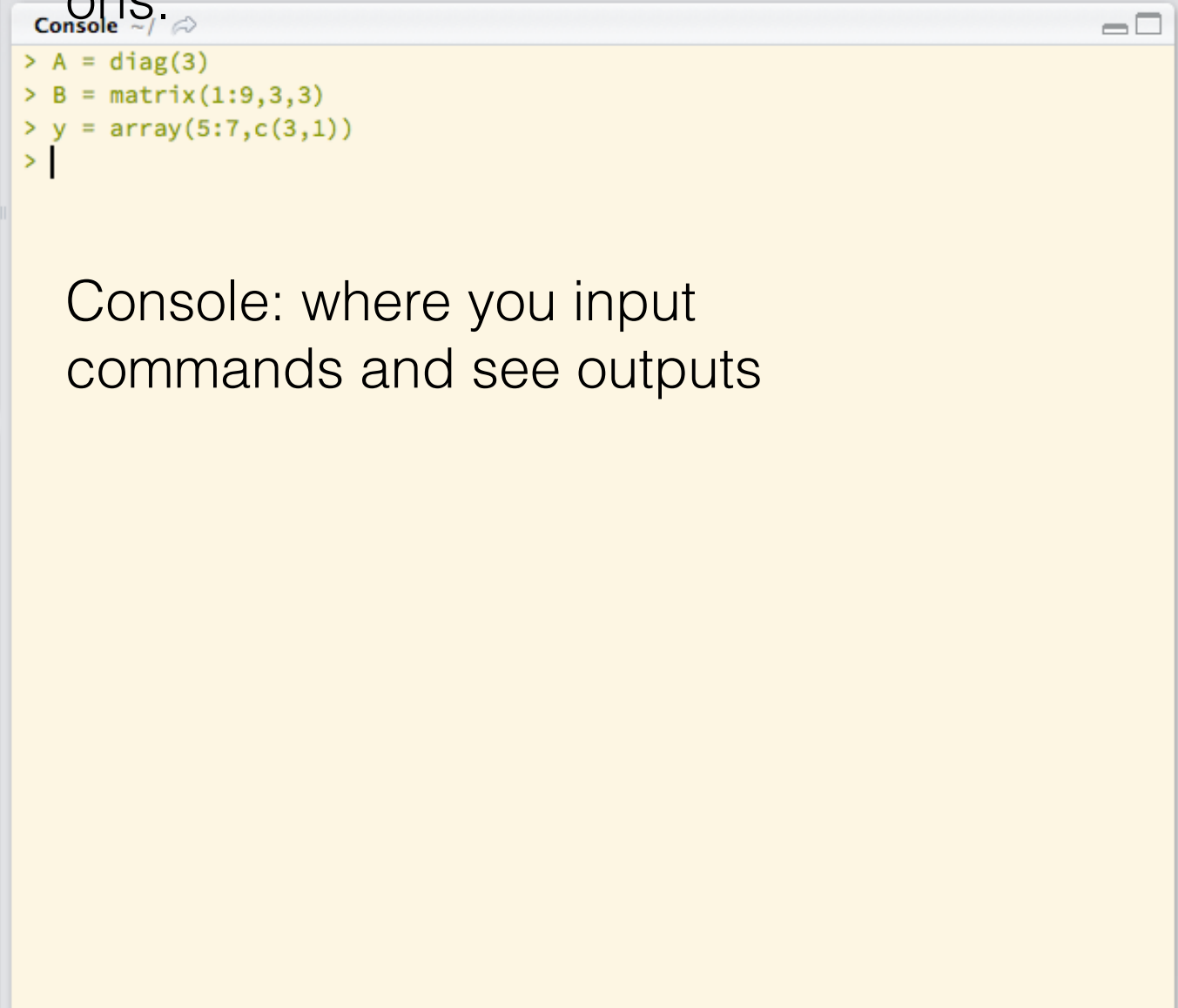
History: a list of commands you used so far.



Files: all the files and folders in your default workspace

Plots: where you see all graphs.

Packages: a list of packages or add-ons.



Console: where you input commands and see outputs

RStudio

Go to file/function

MyScript.R x

```
1 A = diag(3)
2 B = matrix(1:9,3,3)
3 y = array(5:7,c(3,1))
```

Run Source

Files Plots Packages Help Viewer

New Folder Delete Rename More

Home > Dropbox > STAT5474 > lab

Name	Size	Modified
..		
.Rhistory	304 B	Jan 26, 2016, 7:09 PM
	53 B	Jan 26, 2016, 2:14 AM

Options

Choose the layout of the panes in RStudio by selecting from the controls in each quadrant.

General

Code

Appearance

Pane Layout

Packages

Sweave

Spelling

Git/SVN

Publishing

Source

Files, Plots, Packages, Help, Vi

Environment

History

Files

Plots

Packages

Help

Build

VCS

Viewer

Environment, History, Build, VC

Console

Environment

History

Import Dataset

Global Environment

Data

A	num [1:3, 1:3]	1 0 0 0
B	int [1:3, 1:3]	1 2 3 4
y	int [1:3, 1]	5 6 7

Layout can be changed from tools -> global options -> pane layout

RStudio

Project: (None)

MyScript.R x A x

Filter

	V1	V2	V3
1	1	0	0
2	0	1	0
3	0	0	1

Showing 1 to 3 of 3 entries

Environment History

Global Environment

Data

A	num [1:3, 1:3]	1 0 0 0 1 0 0 0 1
B	int [1:3, 1:3]	1 2 3 4 5 6 7 8 9
y	int [1:3, 1]	5 6 7

Files Plots Packages Help Viewer

New Folder Delete Rename More

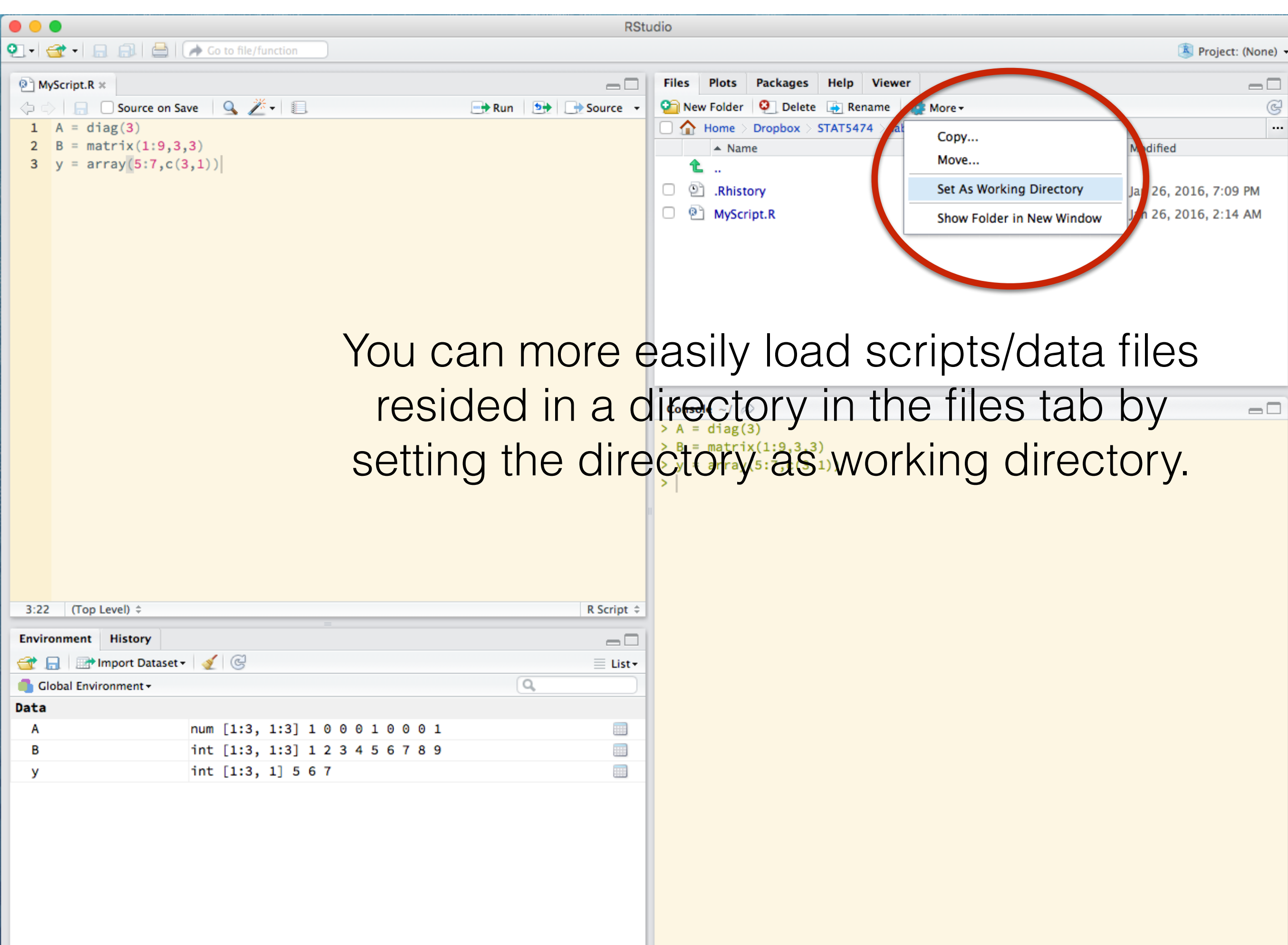
Home > Dropbox > STAT5474 > lab

	Name	Size	Modified
	..		
	.Rhistory	304 B	Jan 26, 2016, 7:09 PM
	MyScript.R	53 B	Jan 26, 2016, 2:14 AM

Console ~/

```
> A = diag(3)
> B = matrix(1:9,3,3)
> y = array(5:7,c(3,1))
> View(A)
> |
```

click one of these, and you can see objects in the source window



You can more easily load scripts/data files resided in a directory in the files tab by setting the directory as working directory.

The screenshot shows the RStudio interface. In the top right, the 'Project' dropdown is set to '(None)'. The 'Packages' tab is active, displaying a list of installed and available packages. The 'Install' button, located in the top toolbar of the Packages tab, is circled in red. A dialog box titled 'Install Packages' is open in the foreground, showing the 'Repository (CRAN)' dropdown, a text field for 'Packages (separate multiple with space or comma):', the 'Install to Library:' dropdown set to '/usr/local/lib/R/3.2/site-library [Default]', and the 'Install dependencies' checkbox checked. The 'Install' button in the dialog is highlighted.

MyScript.R x

Source on Save Run Source Install Update

Files Plots Packages Help Viewer

System Library

Name	Description	Version
<input type="checkbox"/> abind	Combine Multidimensional Arrays	1.4-3
<input type="checkbox"/> acepack	ace() and avas() for selecting regression transformations	1.3-3.3
<input type="checkbox"/> annotate	Annotation for microarrays	1.46.1
<input type="checkbox"/> AnnotationDbi	Annotation Database Interface	1.30.1
<input type="checkbox"/> arm	Data Analysis Using Regression and Multilevel/Hierarchical Models	1.8-6
<input type="checkbox"/> Biobase	Biobase: Base functions for Bioconductor	2.28.0
<input type="checkbox"/> BiocGenerics	S4 generic functions for Bioconductor	0.14.0
<input type="checkbox"/> BiocInstaller	Install/Update Bioconductor and CRAN Packages	1.18.5
<input type="checkbox"/> Bioconductor	Bioconductor facilities for parallel evaluation	1.2.22

Install Packages

Install from: ? Configuring Repositories

Repository (CRAN)

Packages (separate multiple with space or comma):

Install to Library:

/usr/local/lib/R/3.2/site-library [Default]

☒ Install dependencies

Install Cancel

3:22 (Top Level) ↕

Environment History

Global Environment

Data

Variable	Type	Value
A	num [1:3, 1:3]	1 0 0 0 1 0 0 0 1
B	int [1:3, 1:3]	1 2 3 4 5 6 7 8 9
y	int [1:3, 1]	5 6 7

In the package tab, click install, and you can pop-up a window to install R packages.

RStudio

Project: (None)

MyScript.R x

```
1 A = diag(3)
2 B = matrix(1:9,3,3)
3 y = array(5:7,c(3,1))
```

Files Plots Packages Help Viewer

Save as Image...
Save as PDF...
Copy to Clipboard...

Console ~/Dropbox/STAT5474/lab/

```
> prostate=read.table("prostate.txt")
> plot(prostate)
> |
```

3:22 (Top Level) R Script

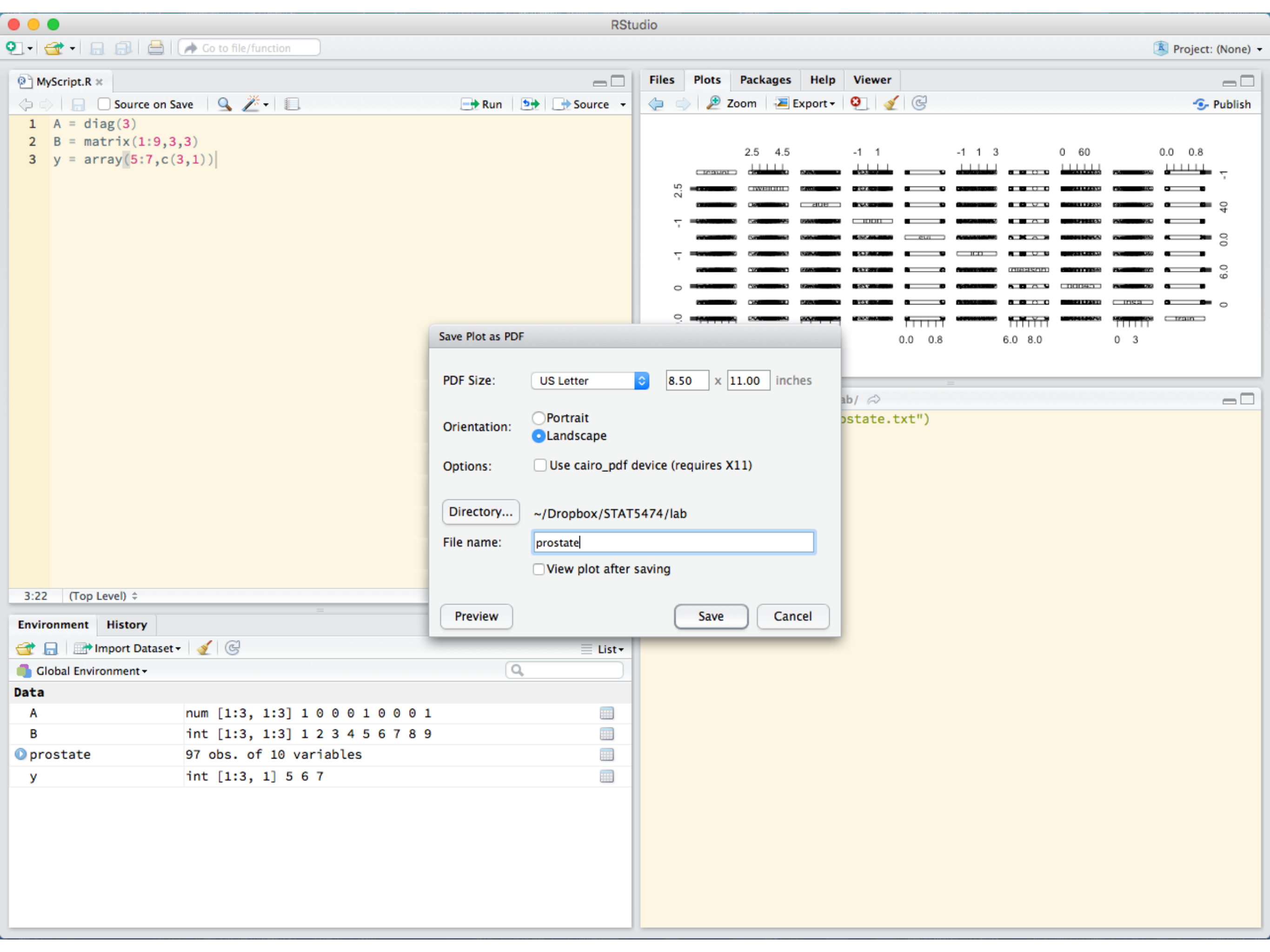
Environment History

Global Environment

Data

A	num [1:3, 1:3]	1 0 0 0 1 0 0 0 1
B	int [1:3, 1:3]	1 2 3 4 5 6 7 8 9
prostate	97 obs. of 10 variables	
y	int [1:3, 1]	5 6 7

From the plots tab, export -> save as image/
PDF lets you save a plot as a file



R Markdown

- ❖ R Markdown is a format that enables easy authoring of reproducible reports from R.
- ❖ It combines the core syntax of Markdown with embedded R code chunks that are run so their output can be included in the final document. i.e. you do not need to copy and paste plots or text from R to the final document
- ❖ R Markdown can produce HTML/PDF/WORD document, but it is optimized for HTML.

Rmd is the extension of R Markdown files. You can open/compile Rmd files to generate the final document in R Studio.

The screenshot displays the RStudio environment with the following components:

- Editor Pane:** Contains an R Markdown file named `Rbasics.Rmd`. The code includes a YAML header, an R chunk with arithmetic and function examples, and a text block. A red circle highlights the `Knit HTML` menu option, which also shows `Knit PDF`, `Knit Word`, `View in Pane`, and `View in Window`.
- Files Pane:** Shows the project directory structure: `Home > Dropbox > STAT5474 > lab`. It lists files including `.Rhistory`, `MyScript.R`, `prostate.txt`, and `Rbasics.Rmd`.
- Console:** Shows the current directory path: `~/Dropbox/STAT5474/lab/`.
- Environment:** The `Global Environment` is currently empty.