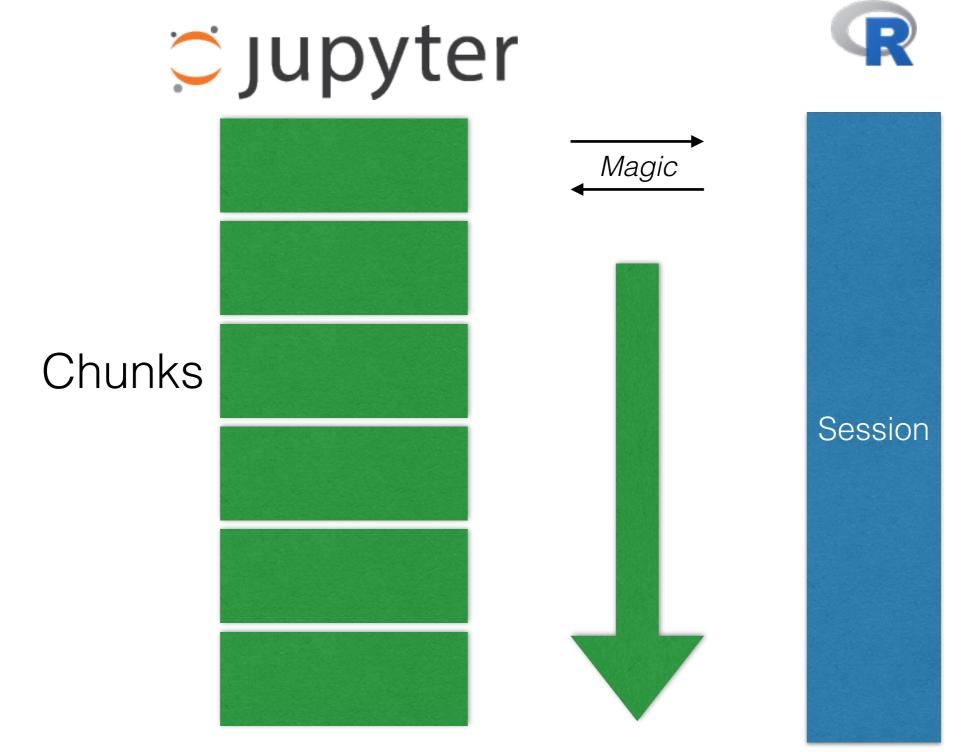
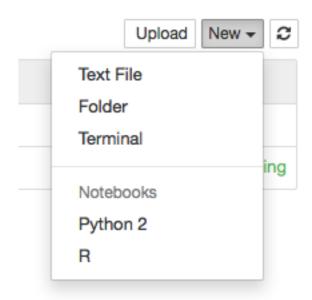
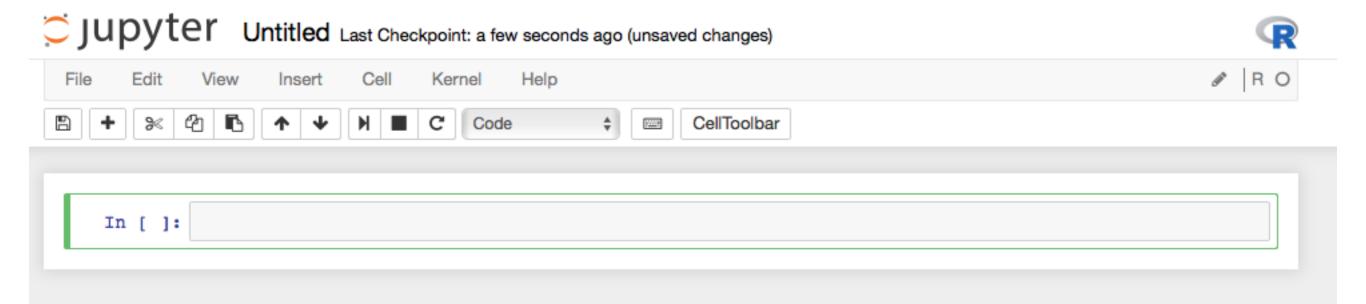
Jupyter

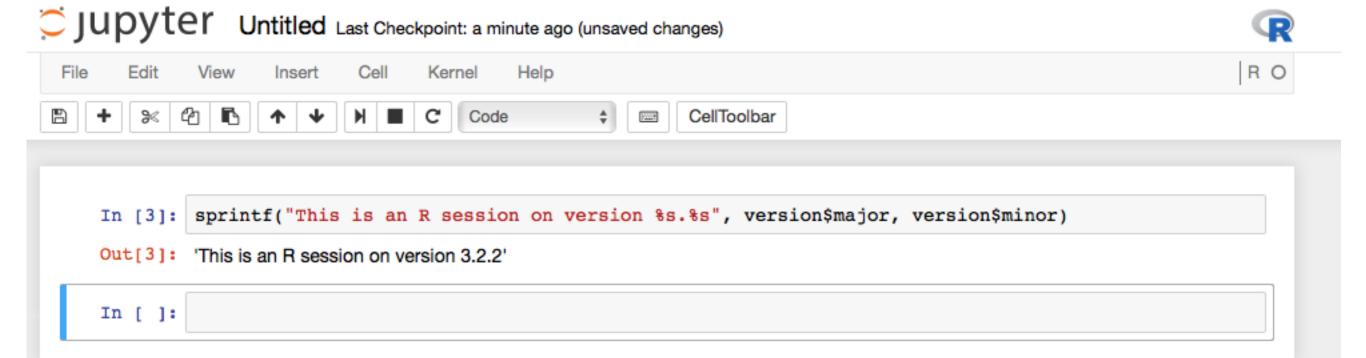


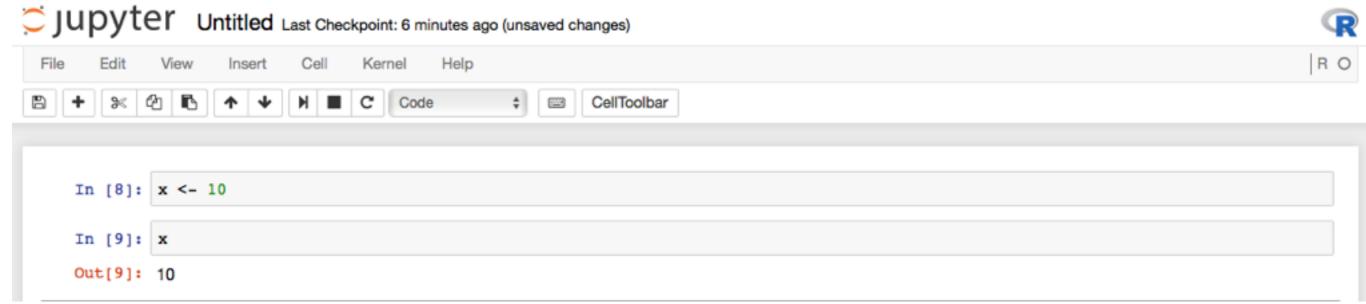








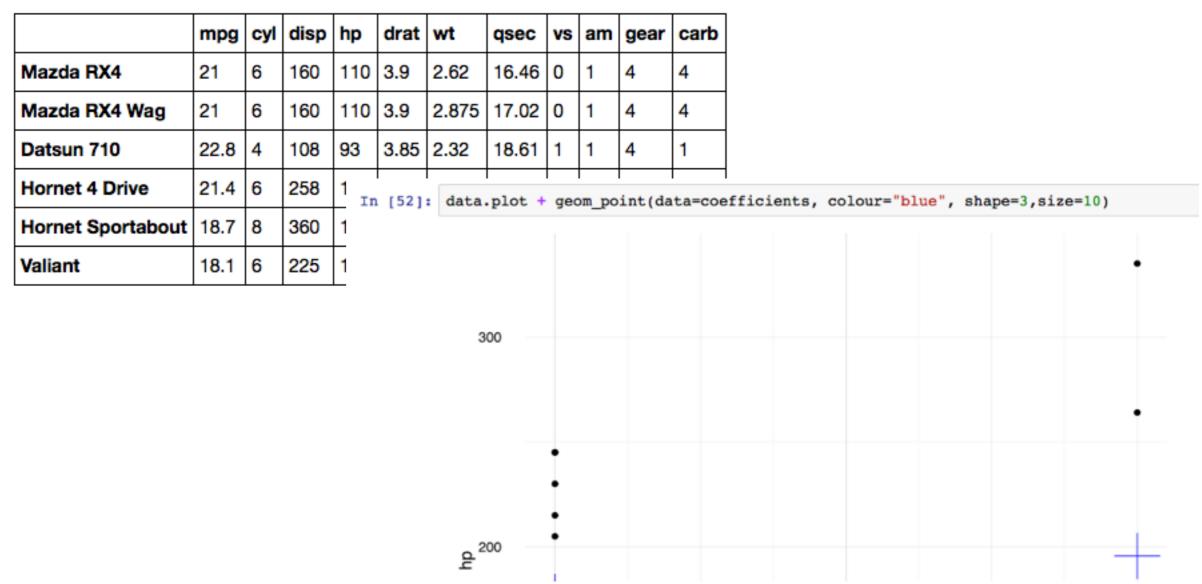




Demo time

Sharing

Out[11]:



notebook files include the output so users don't need R

Sharing



JUPYTER FAQ

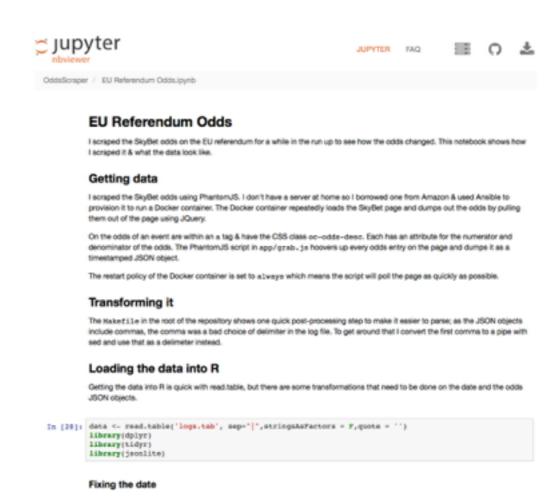
nbviewer

A simple way to share Jupyter Notebooks

URL | GitHub username | GitHub username/repo | Gist ID

Got

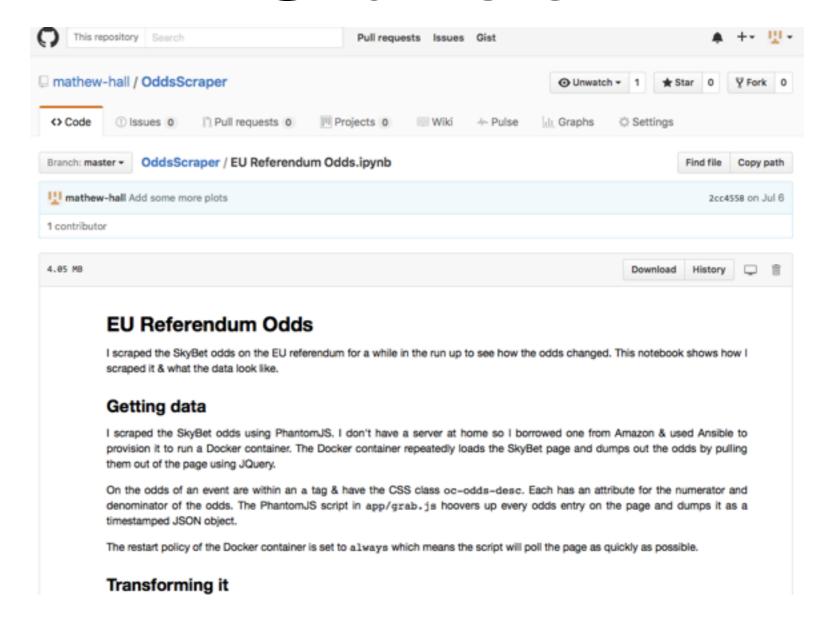
nbviewer.org



The date needs to be parsed first. The datestams include two different timezones (with the same offset) which mean the offset can't

Allows anyone to view a notebook online

GitHub



notebooks in a GitHub repo show nice previews

RStudio + RMarkdown

Jupyter Notebooks

RStudio + Source Files

Mix code & data	Yup	Yup	Not really
Generate publication-ready docs	Yup	Not really	Not really
Share repeatable, remixable analyses	Yup	Yup	Not really
Write testable code	Not really	Not really	Yup
Super interactive	Yup*	Yup	Not really

Use cases

Prototyping analyses

Automatic data cleaning workflows (with bonus reports)

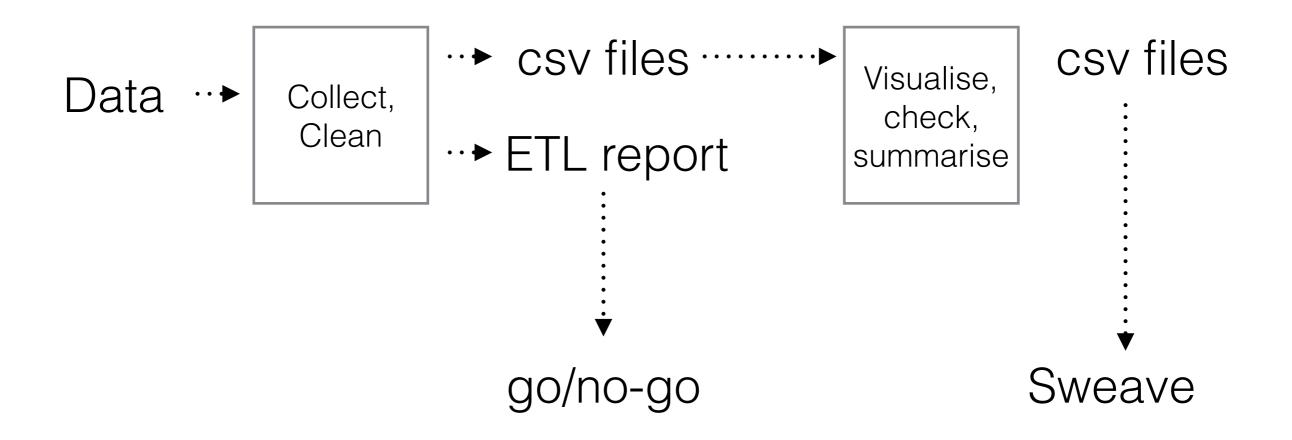
Generating data for use in reports

Dialogs with collaborators (exploratory analyses)

Should I use it?

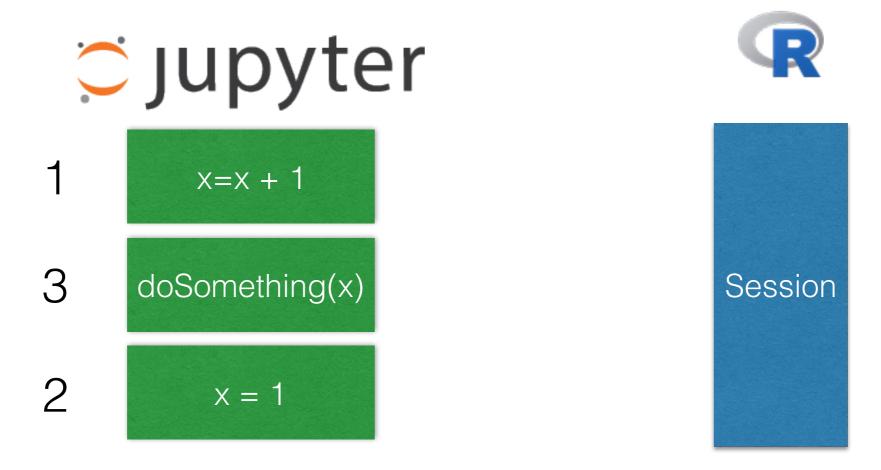
Just running some R commands	R GUI/RStudio	Yeah
Prototyping an analysis	RMarkdown	Yeah
Preparing analysis for publication	RMarkdown, Sweave	Maybe
Data cleaning	RMarkdown? R GUI/ RStudio	Yeah
Collaboration with a lot of people using Git	RMarkdown ,R Package	Probably not

Example



Used as a script with a nicer UI

Developed interactively & deployed as-is

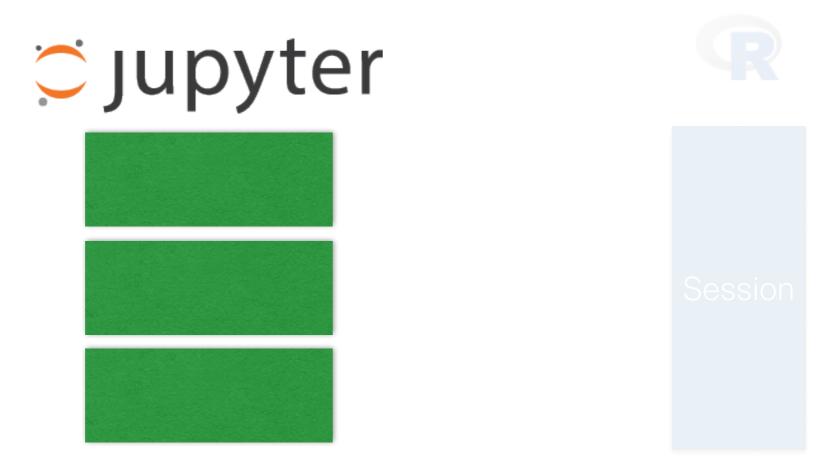


State of the R Session depends on the order chunks run. It's easy to write non-linear notebooks by accident

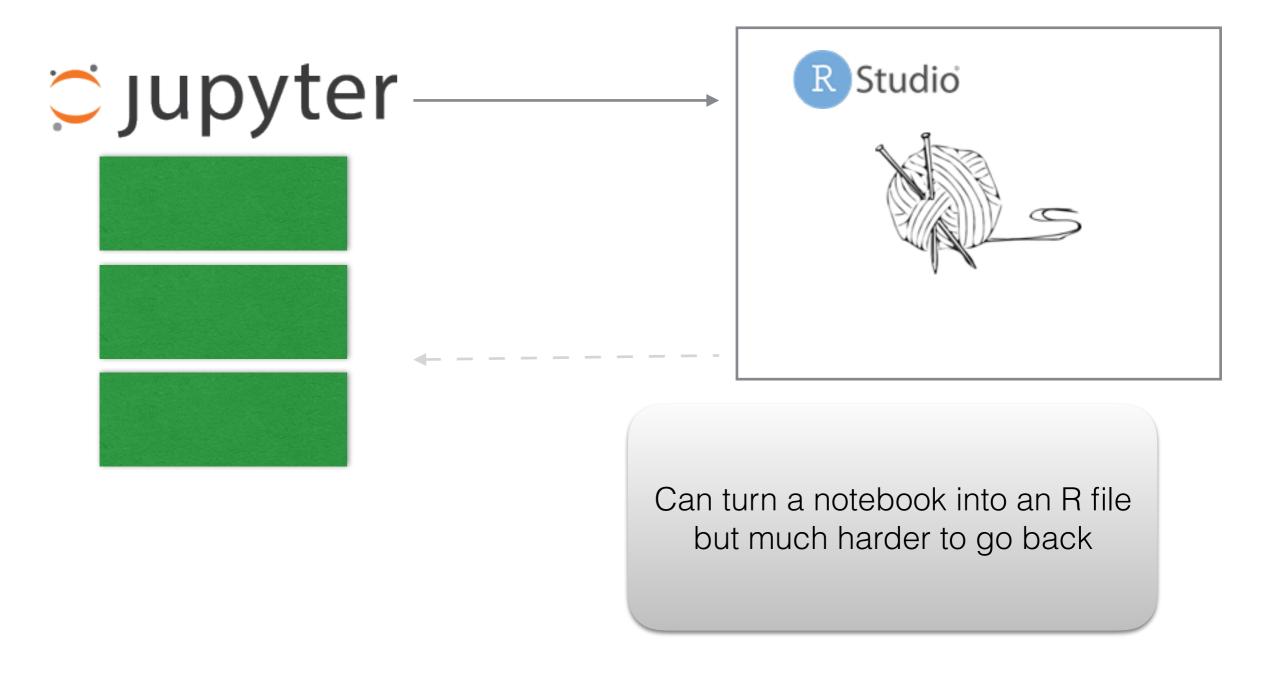
```
In [53]: library(superAwesomePackage)
```

Error in library(superAwesomePackage): there is no package called 'superAwesomePackage'

Need a mechanism to fetch dependencies



State of the R Session is not persistent, only the stored outputs are kept in the notebook file



http://stackoverflow.com/questions/32183164/best-practices-for-turning-jupyter-notebooks-into-python-scripts

Tips

Don't get attached to your R session

Routinely rerun **all** chunks

Don't write enormous notebooks

Migrate code to packages when sensible

Don't silently depend on data/packages

Document dependencies/ automate fetching them

Resist temptation to write spaghetti

Refactor as if the notebook were a source file

Setup



1. Install Jupyter



2. Install IRKernel package



3. Register the kernel in Jupyter

Installing

- Recommended: use Anaconda (https://www.continuum.io/downloads) and R Essentials bundle (http://anaconda.org/r/r-essentials)
- Otherwise: pip install jupyter and follow instructions on (<u>https://irkernel.github.io</u>)
 - Will probably need to install zmq
- Option 3: docker run -d -p 8888:8888 jupyter/rnotebook