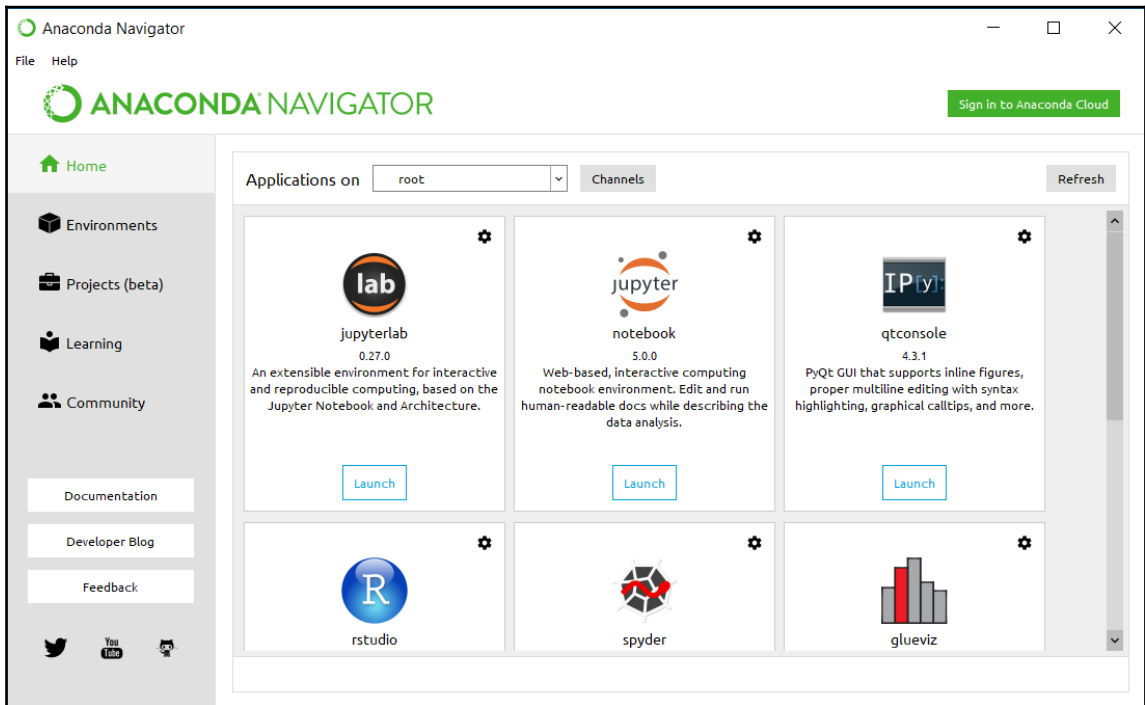
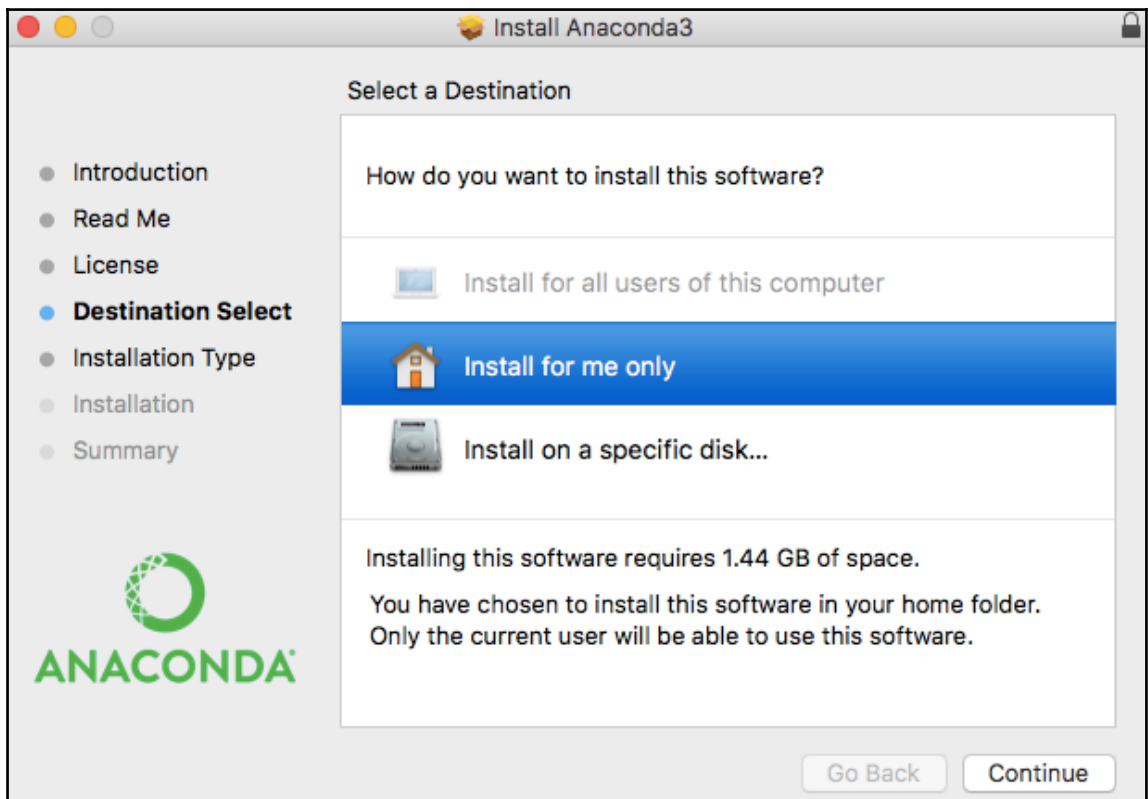
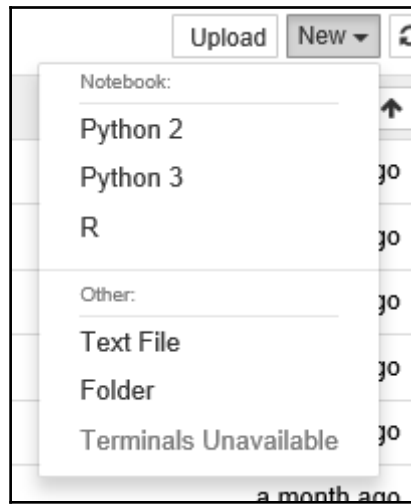


Chapter 1: Installation and Setting up the Environment



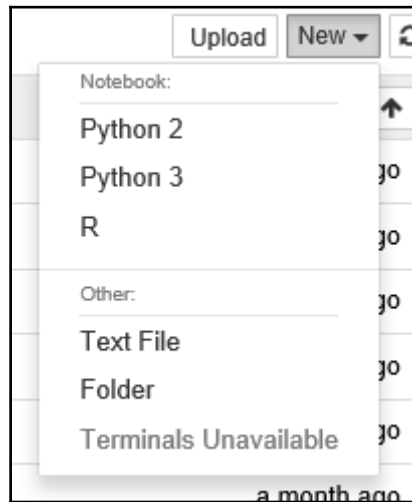
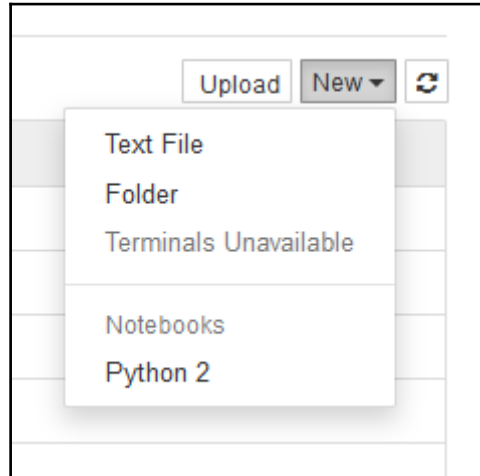


The screenshot shows a Jupyter Notebook window titled "Number of Users (autosaved)". The interface includes a menu bar with "File", "Edit", "View", "Insert", "Cell", "Kernel", "Widgets", and "Help". The current kernel is "Python 2". Below the menu bar is a toolbar with icons for file operations, navigation, and execution. The main area contains a code cell with the following Python code:

```
In [3]: from ipywidgets import interact
def myfunction(x):
    return x
interact(myfunction, x= "Hello World ");
```

Below the code, an interactive widget is displayed. It consists of a label "x" followed by a text input field containing the letter "A". Below the input field, the output of the function is shown as "u'A'".


Chapter 2: Adding an Engine

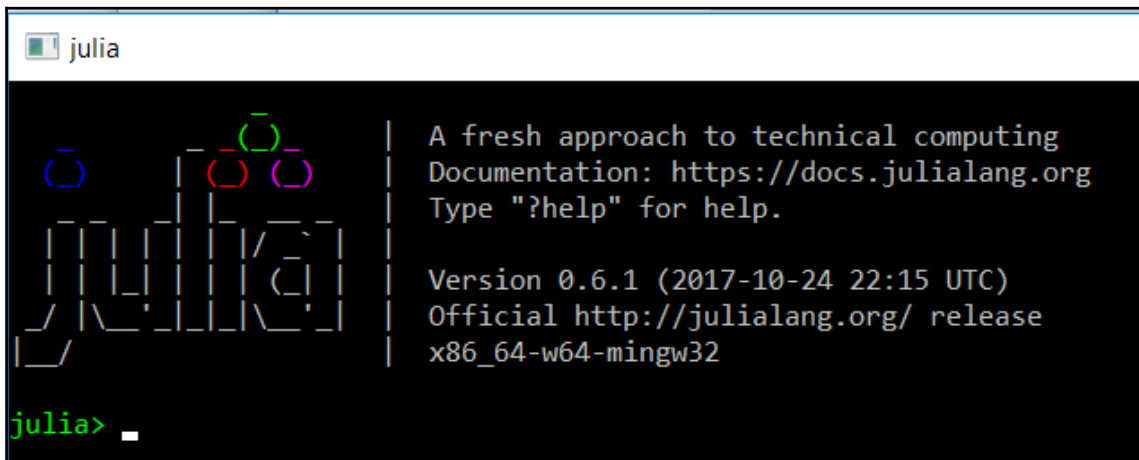
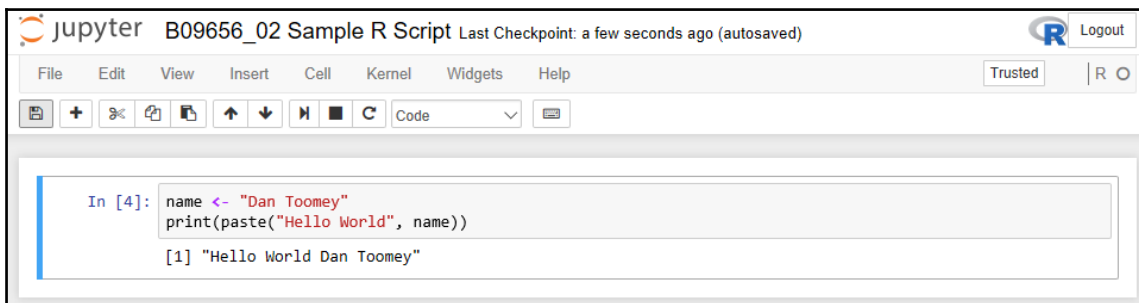
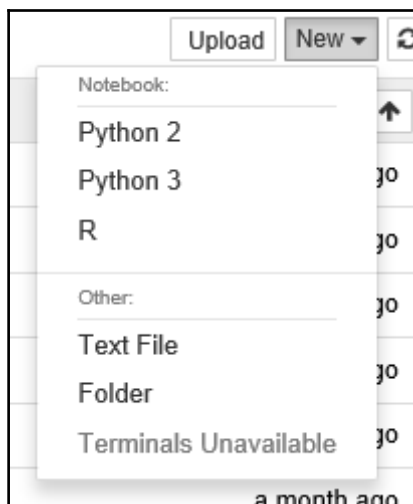


jupyter B09565_02 Python 3 Example Last Checkpoint: a few seconds ago (unsaved changes) Logout

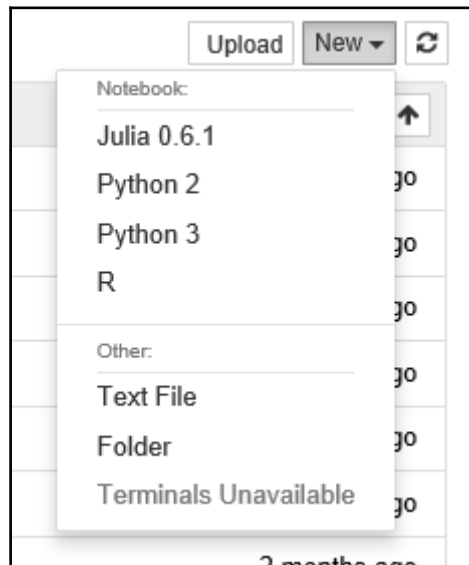
File Edit View Insert Cell Kernel Widgets Help Trusted Python 3

In [1]: `print("Hello World")`
Hello World


rstudio
1.0.136
A set of integrated tools designed to help you be more productive with R. Includes R essentials and notebooks.
Launch



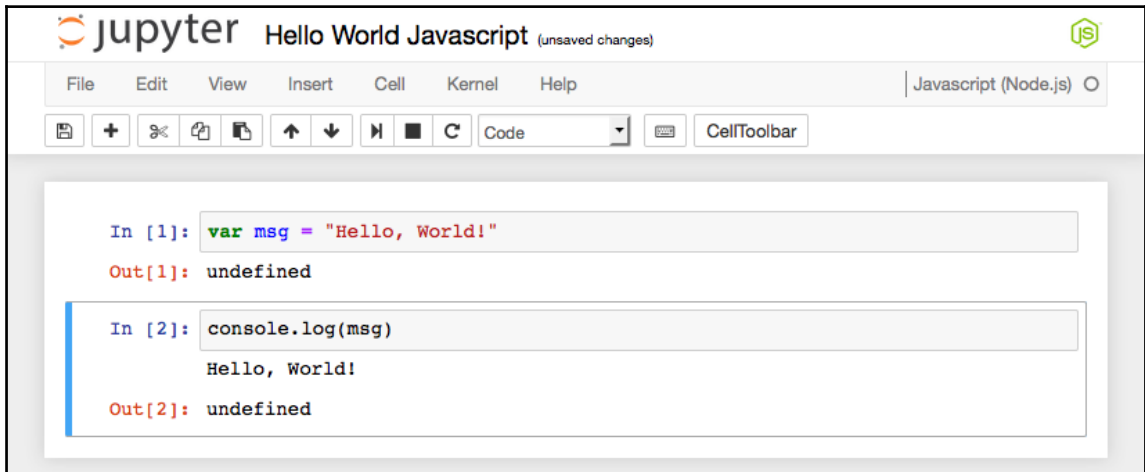
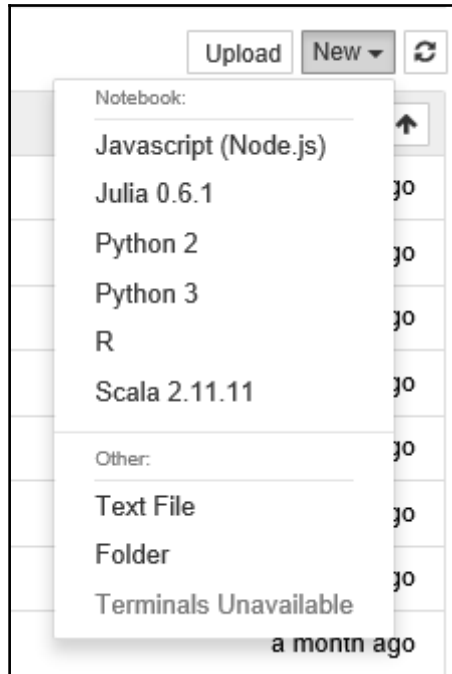
```
julia> Pkg.add("IJulia")
INFO: Initializing package repository C:\Users\Dan\.julia\v0.6
INFO: Cloning METADATA from https://github.com/JuliaLang/METADATA.jl
INFO: Cloning cache of BinDeps from https://github.com/JuliaLang/BinDeps.jl.git
INFO: Cloning cache of BufferedStreams from https://github.com/BioJulia/BufferedStreams.jl.git
INFO: Cloning cache of Compat from https://github.com/JuliaLang/Compat.jl.git
INFO: Cloning cache of Conda from https://github.com/JuliaPy/Conda.jl.git
INFO: Cloning cache of IJulia from https://github.com/JuliaLang/IJulia.jl.git
INFO: Cloning cache of JSON from https://github.com/JuliaIO/JSON.jl.git
INFO: Cloning cache of LibExpat from https://github.com/JuliaIO/LibExpat.jl.git
INFO: Cloning cache of Libz from https://github.com/BioJulia/Libz.jl.git
INFO: Cloning cache of MbedTLS from https://github.com/JuliaWeb/MbedTLS.jl.git
INFO: Cloning cache of SHA from https://github.com/staticfloat/SHA.jl.git
INFO: Cloning cache of URIParser from https://github.com/JuliaWeb/URIParser.jl.git
INFO: Cloning cache of WinRPM from https://github.com/JuliaPackaging/WinRPM.jl.git
INFO: Cloning cache of ZMQ from https://github.com/JuliaInterop/ZMQ.jl.git
INFO: Installing BinDeps v0.7.0
INFO: Installing BufferedStreams v0.2.2
```

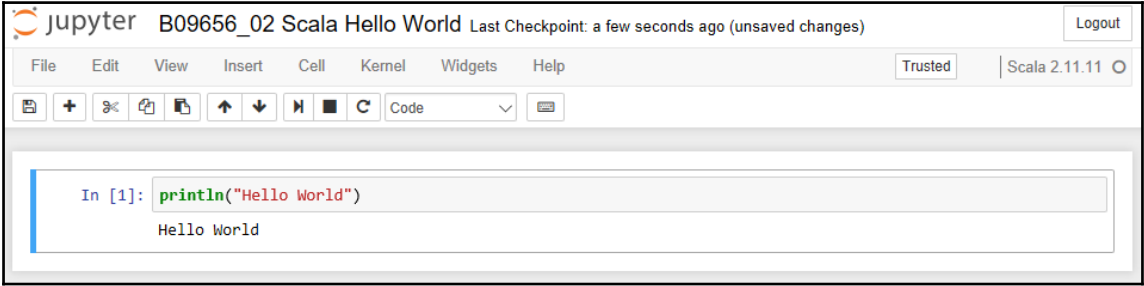
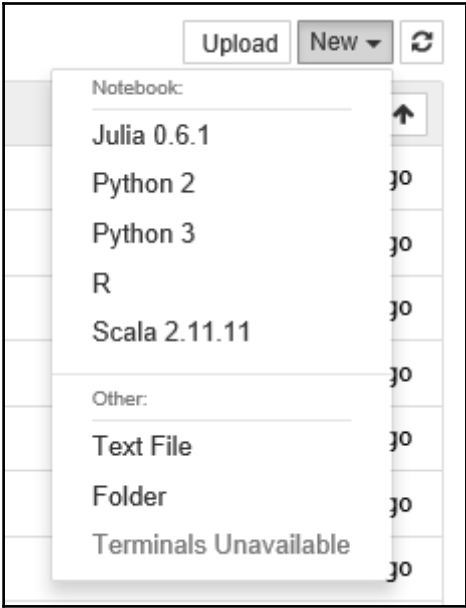


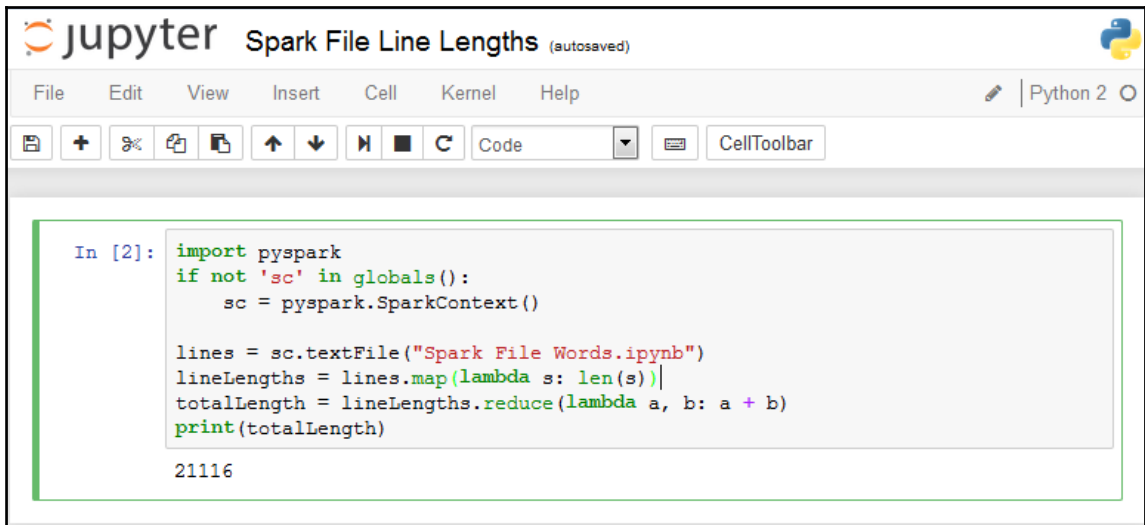
```
In [2]: Pkg.add("RDatasets")
        using RDatasets
        describe(dataset("datasets", "iris"))
```

INFO: Cloning cache of DataArrays from <https://github.com/Julia>
INFO: Cloning cache of DataFrames from <https://github.com/Julia>
INFO: Cloning cache of DataStructures from <https://github.com/Julia>
INFO: Cloning cache of FileIO from <https://github.com/JuliaIO/FileIO>
INFO: Cloning cache of GZip from <https://github.com/JuliaIO/GZip>
INFO: Cloning cache of RData from <https://github.com/JuliaStats/RData>
INFO: Cloning cache of RDatasets from <https://github.com/johnmyers>
INFO: Cloning cache of Reexport from <https://github.com/simonstuart>
INFO: Cloning cache of SortingAlgorithms from <https://github.com/JuliaStats>
INFO: Cloning cache of SpecialFunctions from <https://github.com/JuliaMath>
INFO: Cloning cache of StatsBase from <https://github.com/JuliaStats>
INFO: Installing DataArrays v0.6.2
INFO: Installing DataFrames v0.10.1
INFO: Installing DataStructures v0.7.2
INFO: Installing FileIO v0.5.2
INFO: Installing GZip v0.3.0
INFO: Installing RData v0.1.0
INFO: Installing RDatasets v0.2.0
INFO: Installing Reexport v0.0.3
INFO: Installing SortingAlgorithms v0.1.1
INFO: Installing SpecialFunctions v0.3.4
INFO: Installing StatsBase v0.19.0
INFO: Building SpecialFunctions
INFO: Package database updated
INFO: Precompiling module Reexport.
INFO: Precompiling module FileIO.
INFO: Precompiling module DataFrames.
INFO: Precompiling module RData.

Sepallength
Summary Stats:
Mean: 5.843333
Minimum: 4.300000
1st Quartile: 5.100000
Median: 5.800000
3rd Quartile: 6.400000
Maximum: 7.900000
Length: 150
Type: Float64
Number Missing: 0
% Missing: 0.000000







The image shows a Jupyter Notebook window titled "Spark File Line Lengths (autosaved)". The interface includes a menu bar (File, Edit, View, Insert, Cell, Kernel, Help) and a toolbar with icons for file operations and execution. The code cell contains the following Python code:

```
In [2]: import pyspark
if not 'sc' in globals():
    sc = pyspark.SparkContext()

lines = sc.textFile("Spark File Words.ipynb")
lineLengths = lines.map(lambda s: len(s))
totalLength = lineLengths.reduce(lambda a, b: a + b)
print(totalLength)
```

The output of the code cell is the number 21116.

Chapter 3: Accessing and Retrieving Data

In [2]: `head(heating)`

X	idcase	depvar	ic.gc	ic.gr	ic.ec	ic.er	ic.hp	oc.gc	oc.gr	...	oc.hp	income	agehed
1	1	gc	866.00	962.64	859.90	995.76	1135.50	199.69	151.72	...	237.88	7	25
2	2	gc	727.93	758.89	796.82	894.69	968.90	168.66	168.66	...	199.19	5	60
3	3	gc	599.48	783.05	719.86	900.11	1048.30	165.58	137.80	...	171.47	4	65
4	4	er	835.17	793.06	761.25	831.04	1048.70	180.88	147.14	...	222.95	2	50
5	5	er	755.59	846.29	858.86	985.64	883.05	174.91	138.90	...	178.49	2	25
6	6	gc	666.11	841.71	693.74	862.56	859.18	135.67	140.97	...	209.27	6	65

In [7]: `head(heating)`

X	system	install_cost	annual_cost	income	agehed	rooms	region	ratio_annual_install
1	gc	866.00	199.69	7	25	6	ncostl	4.336722
2	gc	727.93	168.66	5	60	5	scostl	4.315961
3	gc	599.48	165.58	4	65	2	ncostl	3.620486
4	er	835.17	180.88	2	50	4	scostl	4.617260
5	er	755.59	174.91	2	25	6	valley	4.319879
6	gc	666.11	135.67	6	65	7	scostl	4.909781

```
{
  "Models": [
    {
      "model_name": "021 C",
      "model_make_id": "ford"
    },
    {
      "model_name": "12 M",
      "model_make_id": "ford"
    },
  ],
}
```

```
//load the JSON dataset
//http://www.carqueryapi.com/api/0.3/?callback=?&cmd=getModels&make=ford
var fords = require('/Users/dtoomey/fords.json');

//display how many Ford models are in our data set
console.log("There are " + fords.Models.length + " Ford models in the data set");

//loop over the set
var index = 1
for(var i=0; i<fords.Models.length; i++) {

  //get this model
  var model = fords.Models[i];

  //pull it's name
  var name = model.model_name;

  //if the model name does not have numerics in it
  if(! name.match(/[0-9]/i)) {
    //display the model name
    console.log("Model " + index + " is a " + name);
    index++;
  }

  //only display the first 5
  if (index>5) break;
}
```

```
There are 147 Ford models in the data set
Model 1 is a Aerostar
Model 2 is a Anglia
Model 3 is a Artic
Model 4 is a Aspire
Model 5 is a Bantam
```

```
Out[27]: 5
```

```
 Sepal.Length Sepal.Width Petal.Length Petal.Width
Min. :4.300 Min. :2.000 Min. :1.000 Min. :0.100
1st Qu.:5.100 1st Qu.:2.800 1st Qu.:1.600 1st Qu.:0.300
Median :5.800 Median :3.000 Median :4.350 Median :1.300
Mean :5.843 Mean :3.057 Mean :3.758 Mean :1.199
3rd Qu.:6.400 3rd Qu.:3.300 3rd Qu.:5.100 3rd Qu.:1.800
Max. :7.900 Max. :4.400 Max. :6.900 Max. :2.500
Species
setosa :50
versicolor:50
virginica :50
```

```
# Source: table<iris> [?? x 5]
# Database: sqlite 3.19.3 []
 Sepal.Length Sepal.Width Petal.Length Petal.Width Species
      <dbl>      <dbl>      <dbl>      <dbl>      <chr>
1         5.1         3.5         1.4         0.2 setosa
2         4.9         3.0         1.4         0.2 setosa
3         4.7         3.2         1.3         0.2 setosa
4         4.6         3.1         1.5         0.2 setosa
5         5.0         3.6         1.4         0.2 setosa
6         5.4         3.9         1.7         0.4 setosa
7         4.6         3.4         1.4         0.3 setosa
8         5.0         3.4         1.5         0.2 setosa
9         4.4         2.9         1.4         0.2 setosa
10        4.9         3.1         1.5         0.1 setosa
# ... with more rows
```

```
In [21]: head(iris_db, n = 10)

# Source:   lazy query [?? x 5]
# Database: sqlite 3.19.3 []
   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
   <dbl>         <dbl>         <dbl>         <dbl>    <chr>
1         5.1         3.5           1.4         0.2  setosa
2         4.9         3.0           1.4         0.2  setosa
3         4.7         3.2           1.3         0.2  setosa
4         4.6         3.1           1.5         0.2  setosa
5         5.0         3.6           1.4         0.2  setosa
6         5.4         3.9           1.7         0.4  setosa
7         4.6         3.4           1.4         0.3  setosa
8         5.0         3.4           1.5         0.2  setosa
9         4.4         2.9           1.4         0.2  setosa
10        4.9         3.1           1.5         0.1  setosa
# ... with more rows
```

```
In [19]: show_query(head(iris_db, n = 10))

<SQL>
SELECT *
FROM `iris`
LIMIT 10
```

```
In [20]: # pull data back into R using collect
my_iris <- iris_db %>% collect()
my_iris
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa



baseball.txt - Notepad

File Edit Format View Help

```
4ansonca0118711RC125120
44forceda0118711WS332162
68mathebo0118711FW119 89
99startjo0118711NY233161
102suttoez0118711CL129128
106whitede0118711CL129146
113yorkto01 18711TR029145
121ansonca0118721PH146217
143burdoja0118721BR237174
167forceda0118721TR025130
168forceda0118722BL119 95
186hinespa0118721WS411 49
209mathebo0118721BL150223
226nelsoca0118721TRO 4 20
227nelsoca0118722BR118 76
229orourji0118721MID23101
249startjo0118721NY255282
252suttoez0118721CL122107
259whitede0118721CL122109
```

Out[1]:

	row	id	year	stint	team	g	ab
0	4	ansonca01	1871	1	RC1	25	120
1	44	forceda01	1871	1	WS3	32	162
2	68	mathebo01	1871	1	FW1	19	89
3	99	startjo01	1871	1	NY2	33	161
4	102	suttoez01	1871	1	CL1	29	128
5	106	whitede01	1871	1	CL1	29	146
6	113	yorkto01	1871	1	TRO	29	145
7	121	ansonca01	1872	1	PH1	46	217
8	143	burdoja01	1872	1	BR2	37	174
9	167	forceda01	1872	1	TRO	25	130
10	168	forceda01	1872	2	BL1	19	95
11	186	hinespa01	1872	1	WS4	11	49
12	209	mathebo01	1872	1	BL1	50	223
13	226	nelsoca01	1872	1	TRO	4	20
14	227	nelsoca01	1872	2	BR1	18	76
15	229	orourji01	1872	1	MID	23	101
16	249	startjo01	1872	1	NY2	55	282
17	252	suttoez01	1872	1	CL1	22	107
18	259	whitede01	1872	1	CL1	22	109

	Length	Class	Mode
1	2	PlainTextDocument	list
2	2	PlainTextDocument	list
3	2	PlainTextDocument	list
4	2	PlainTextDocument	list
5	2	PlainTextDocument	list

<<SimpleCorpus>>

Metadata: corpus specific: 1, document level (indexed): 0

Content: documents: 1

[1] For the second time in a month,<U+00A0>President Trump discussed his plans to change the tax code. This is an annotated transcript of his remarks, which took place at the Indiana Farm Bureau Building.<U+00A0>Click on the highlighted statements below to see a brief summary of statements that we and others have fact-checked.<U+00A0>For more information, please read our story, <U+0093><U+0091>Death Tax<U+0092> Talking Point Won<U+0092>t Die.<U+0094>

<<SimpleCorpus>>

Metadata: corpus specific: 1, document level (indexed): 0

Content: documents: 1

[1] for the second time in a month, president trump discussed his plans to change the tax code. this is an annotated transcript of his remarks, which took place at the indiana farm bureau building. click on the highlighted statements below to see a brief summary of statements that we and others have fact-checked. for more information, please read our story, <U+0093><U+0091>death tax<U+0092> talking point won<U+0092>t die.<U+0094>

<<DocumentTermMatrix (documents: 94, terms: 1048)>>

Non-/sparse entries: 3092/95420

Sparsity : 97%

Maximal term length: 15

Weighting : term frequency (tf)

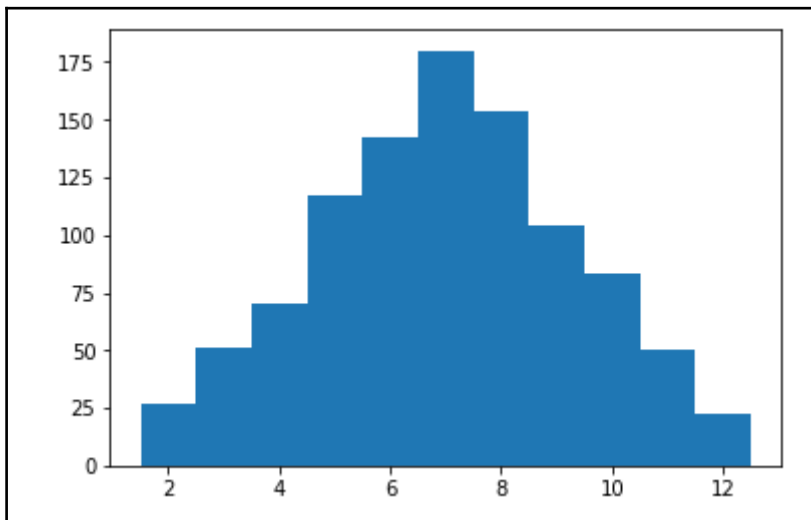
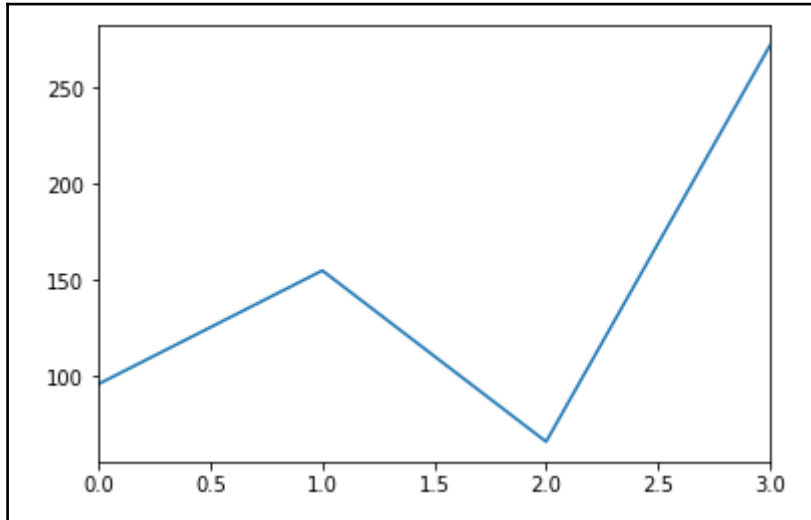
Sample :

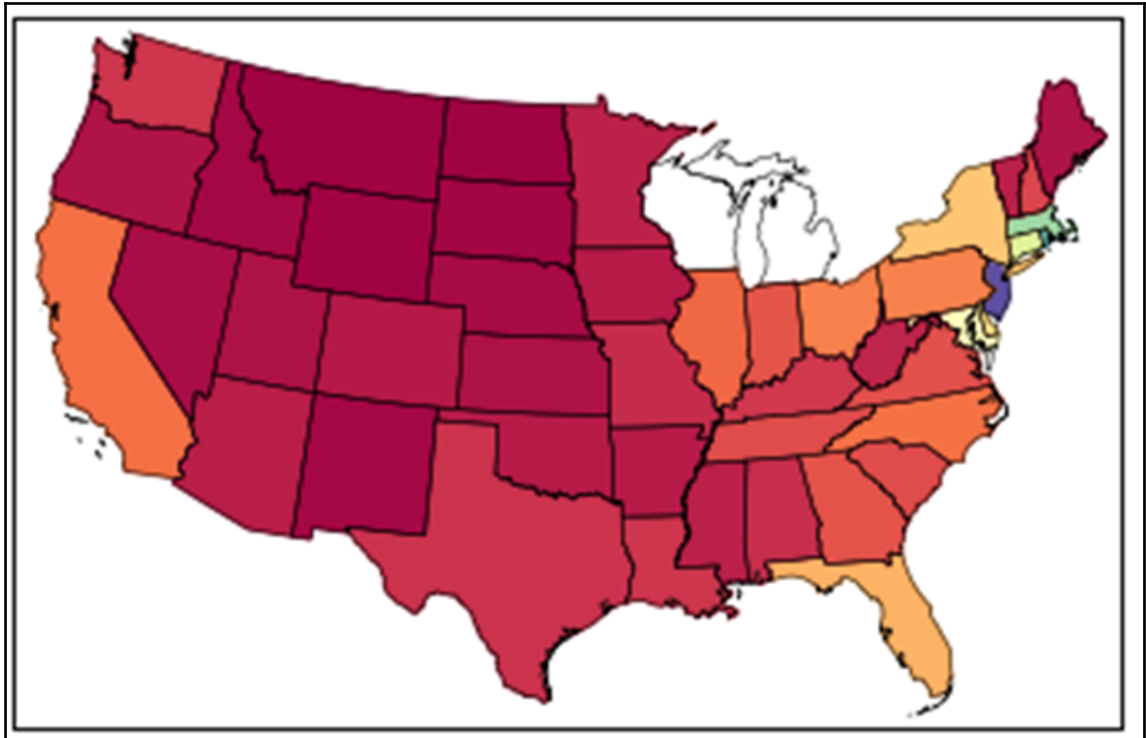
	Terms									
Docs	and	applause	for	have	our	tax	that	the	will	you
10	2	1	1	2	2	1	1	7	0	0
11	11	0	0	3	0	0	3	6	0	1
45	7	1	3	3	0	3	4	8	0	1
47	2	2	1	0	0	2	4	2	0	0
51	4	1	2	1	2	0	1	1	0	1
64	4	2	1	0	1	0	1	5	2	0
67	2	1	1	2	0	0	2	3	1	3
73	5	1	3	0	2	1	0	4	0	0
80	5	1	0	3	0	3	0	6	2	2
9	5	0	0	3	2	0	4	4	0	1

\$tax =

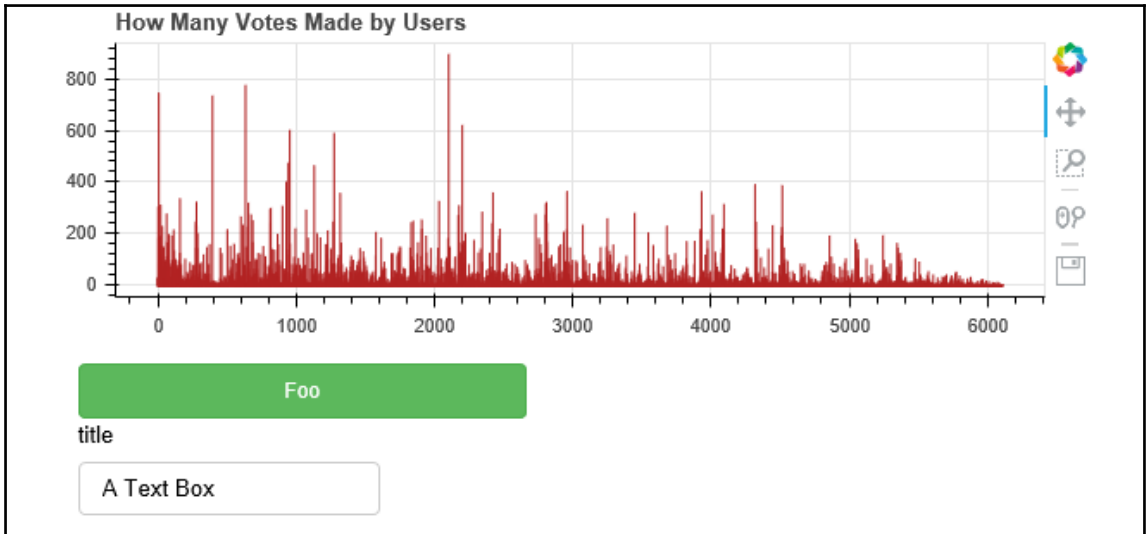
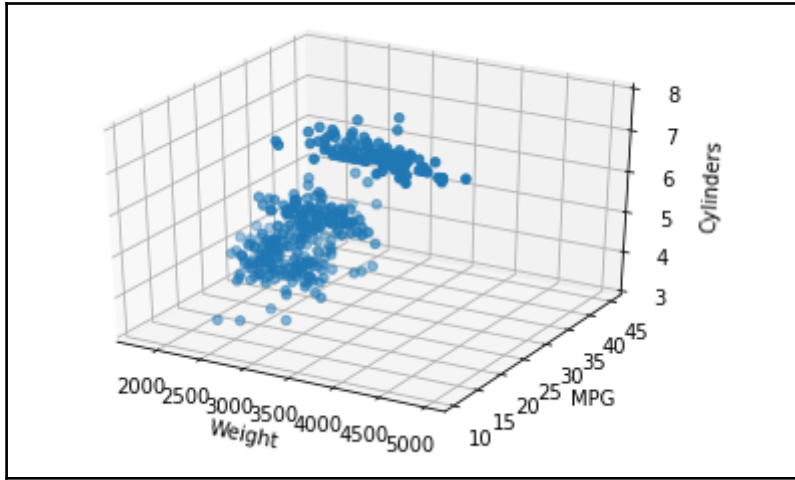
the	0.44
cuts	0.43
pay	0.37
system	0.32
lower	0.3
significant	0.3
breaks	0.3
farms	0.3
competitors	0.3
numerous	0.3
death	0.29
tremendous	0.29
small	0.29
because	0.27
reform	0.27
calculate	0.27

Chapter 4: Visualizing Your Analytics

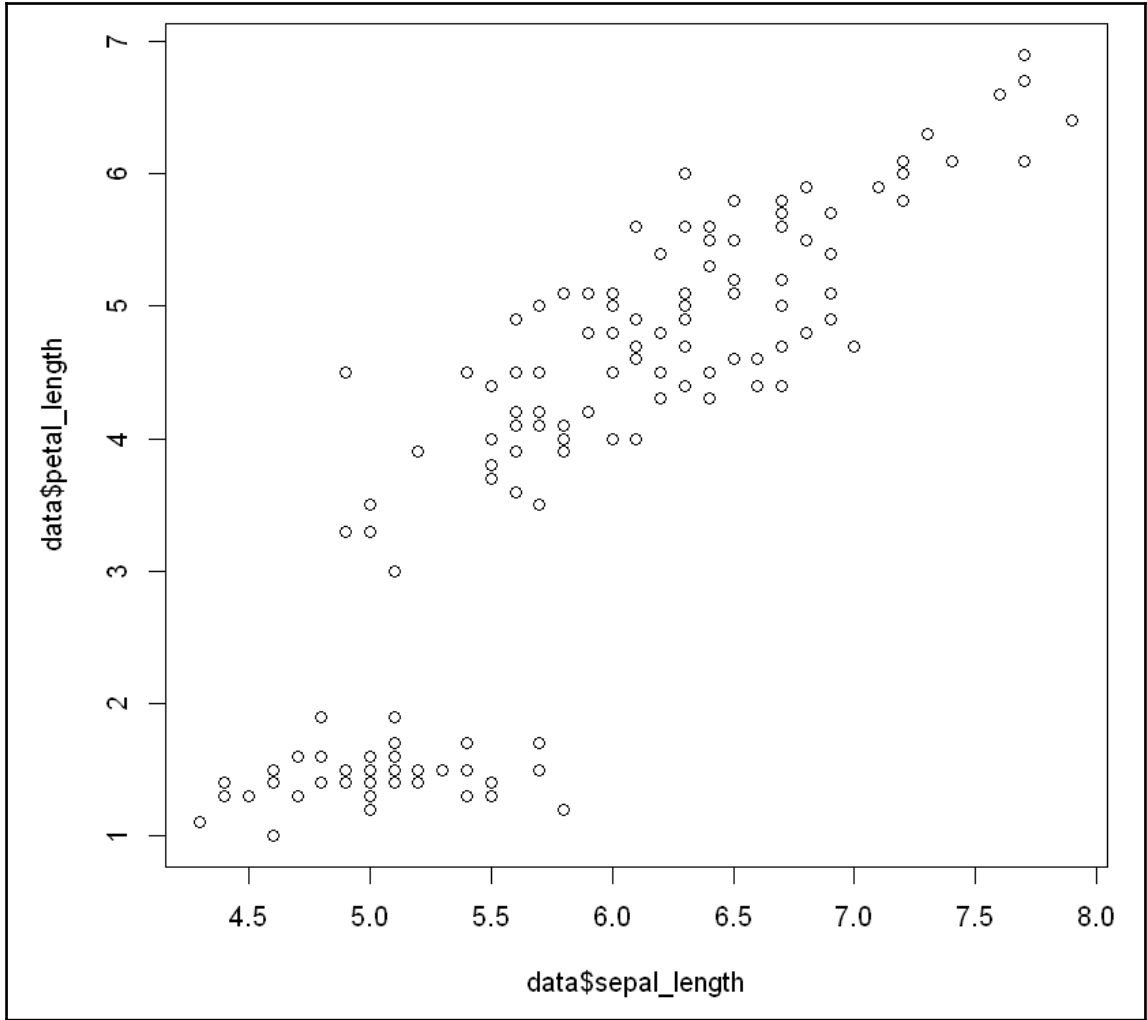


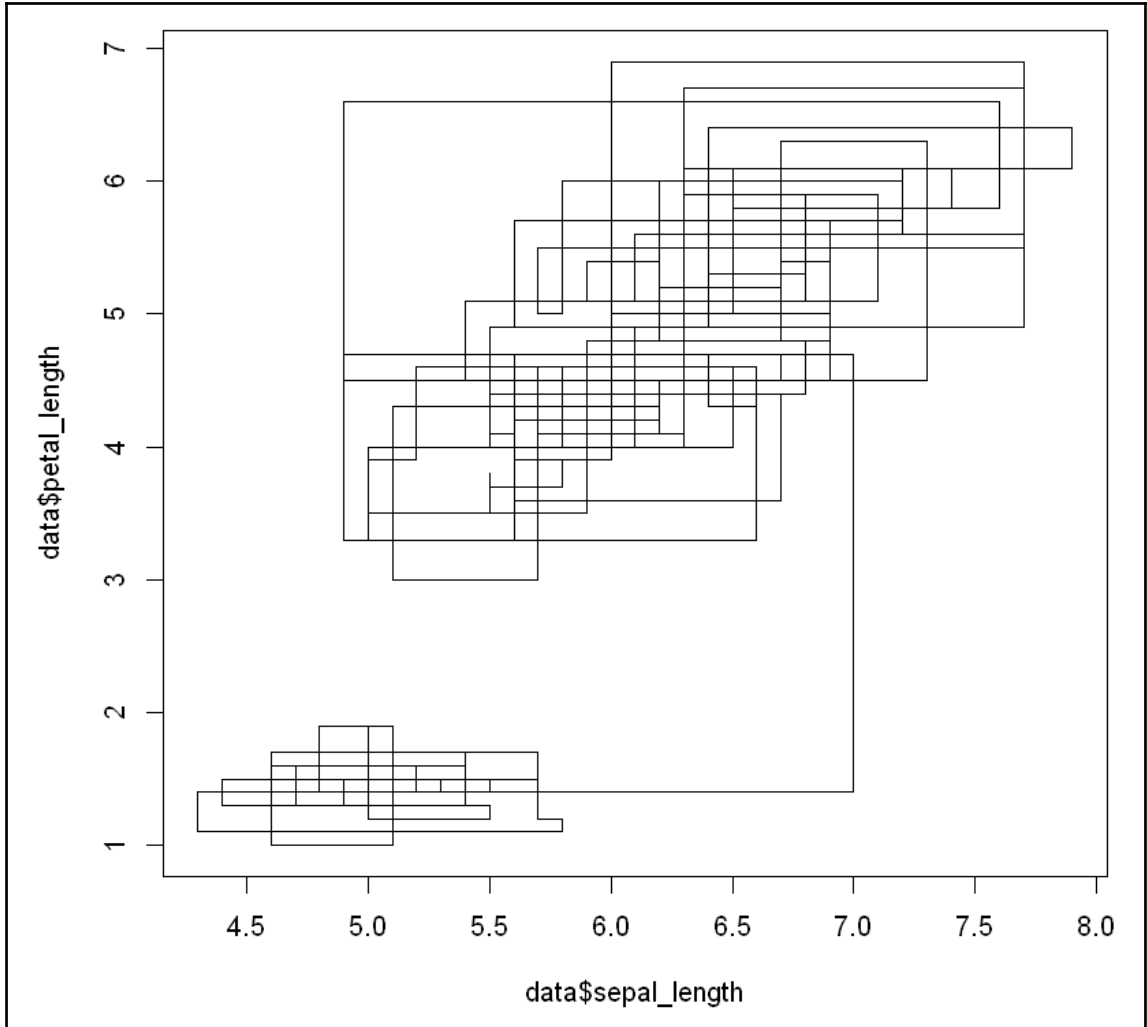


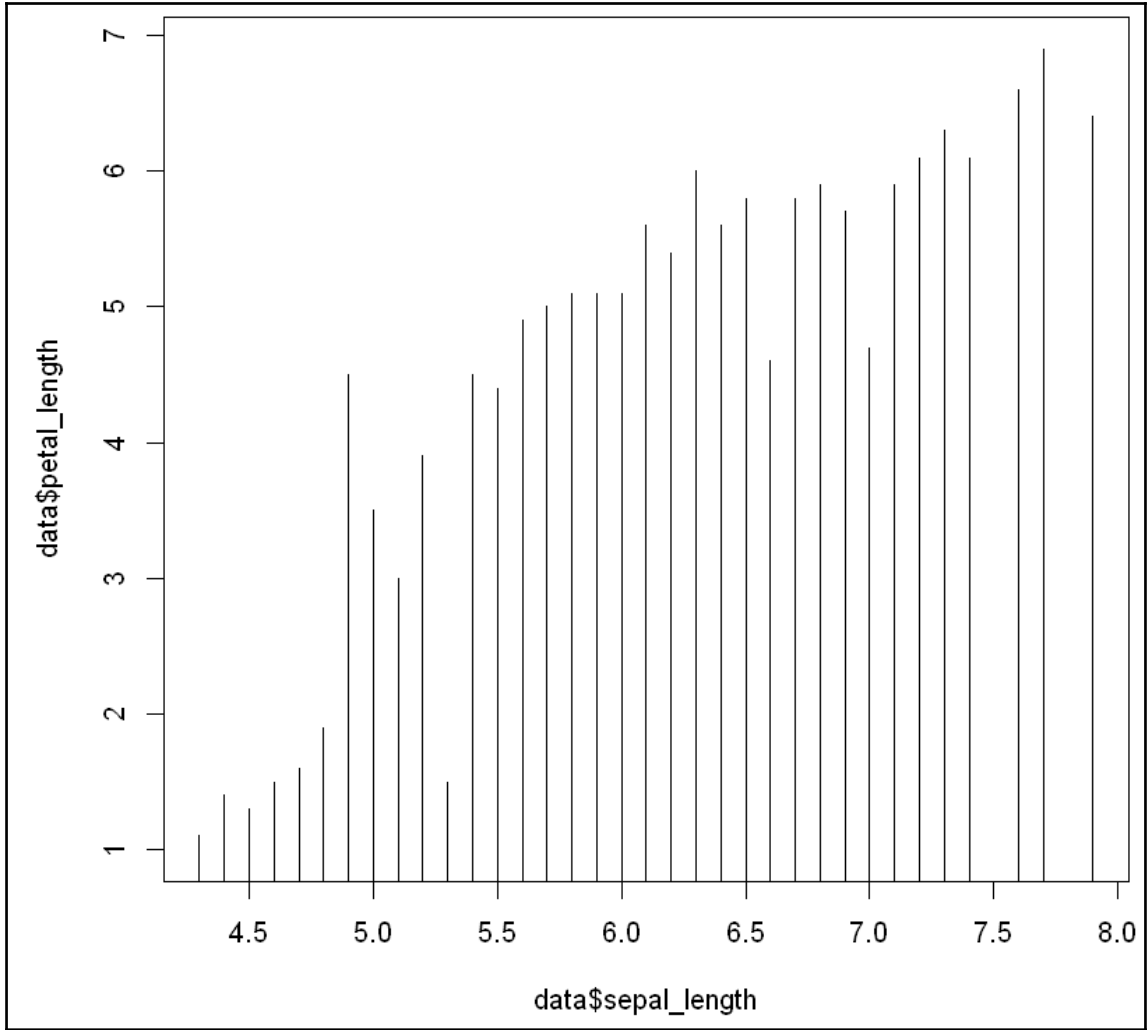
mpg	cylinders	displacement	horsepower	weight	acceleration	year	origin \
18.0	8	307.0	130.0	3504.0	12.0	70	1
15.0	8	350.0	165.0	3693.0	11.5	70	1
18.0	8	318.0	150.0	3436.0	11.0	70	1
16.0	8	304.0	150.0	3433.0	12.0	70	1
17.0	8	302.0	140.0	3449.0	10.5	70	1
							name
18.0							chevrolet chevelle malibu
15.0							buick skylark 320
18.0							plymouth satellite
16.0							amc rebel sst
17.0							ford torino

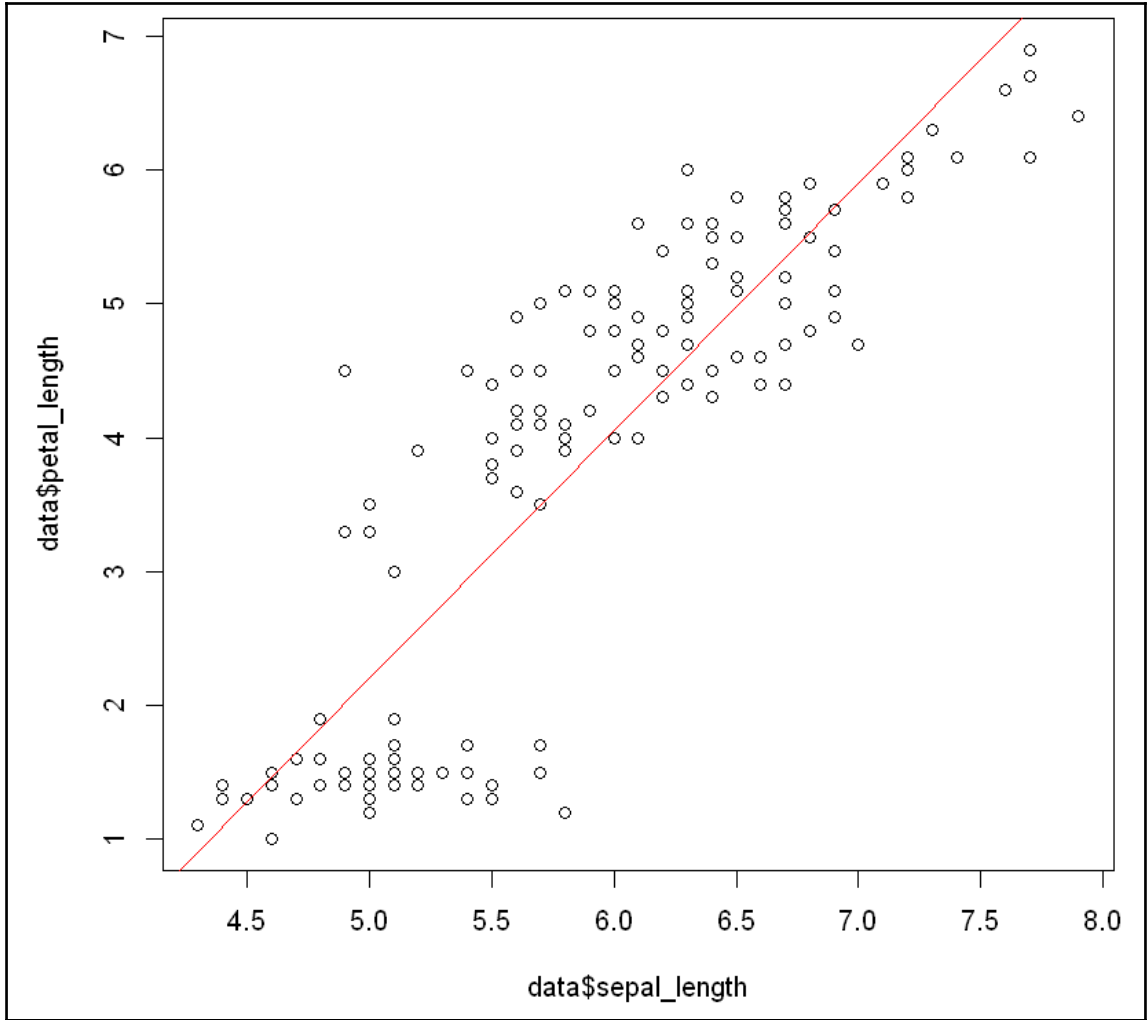


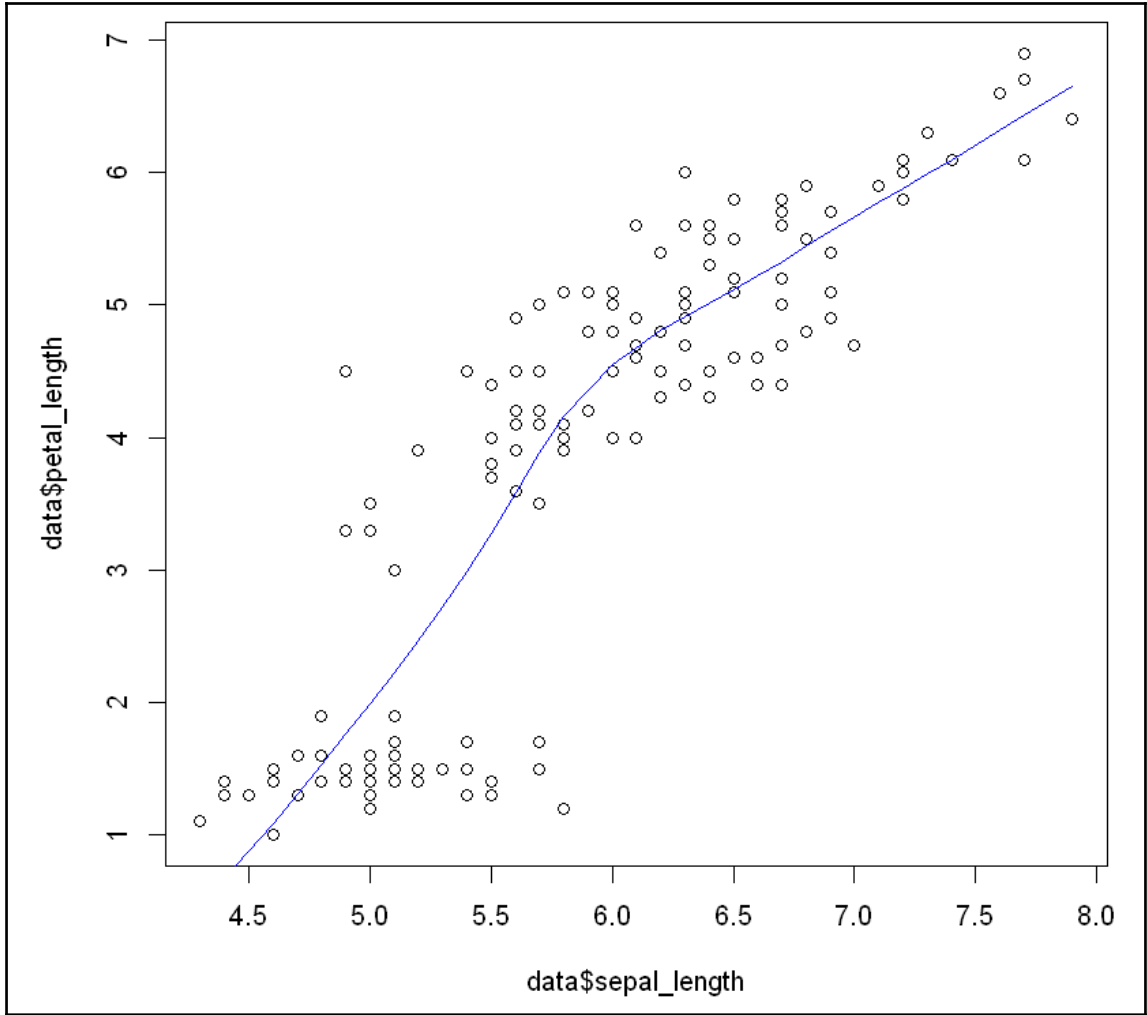
sepal_length	sepal_width	petal_length	petal_width
Min. :4.300	Min. :2.000	Min. :1.000	Min. :0.100
1st Qu.:5.100	1st Qu.:2.800	1st Qu.:1.600	1st Qu.:0.300
Median :5.800	Median :3.000	Median :4.400	Median :1.300
Mean :5.848	Mean :3.051	Mean :3.774	Mean :1.205
3rd Qu.:6.400	3rd Qu.:3.300	3rd Qu.:5.100	3rd Qu.:1.800
Max. :7.900	Max. :4.400	Max. :6.900	Max. :2.500
species			
Iris-setosa	:49		
Iris-versicolor	:50		
Iris-virginica	:50		

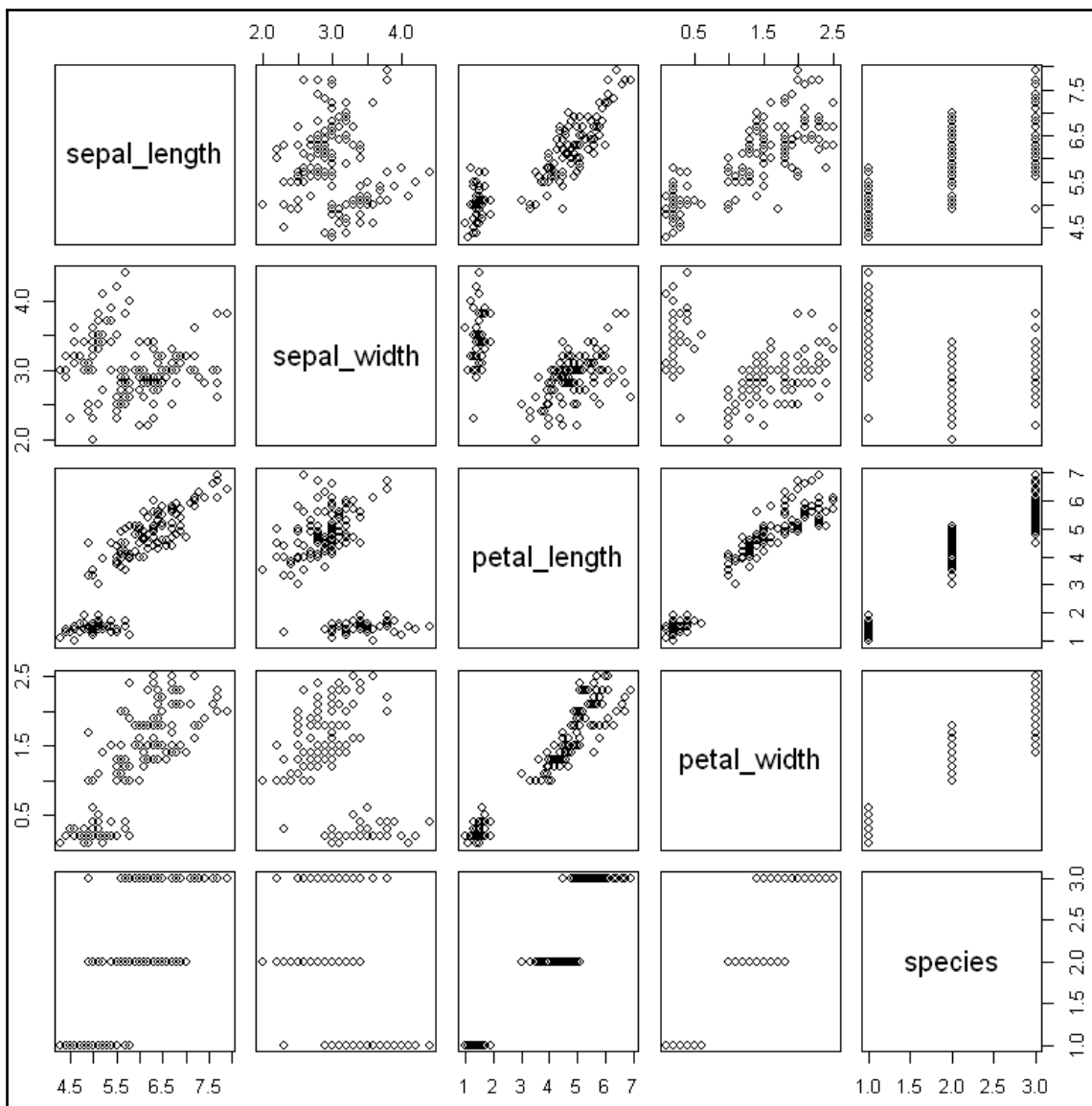












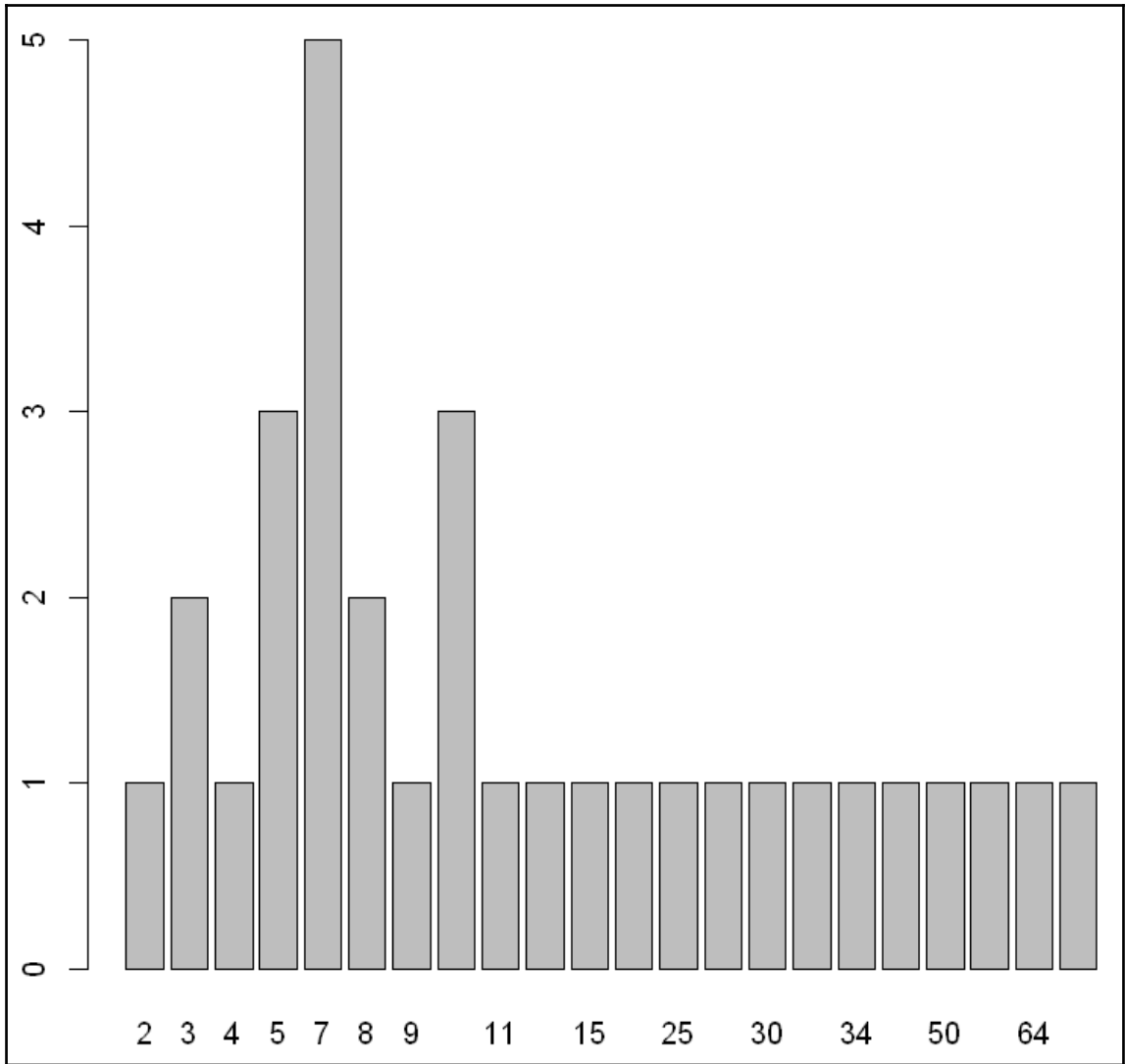
Number of cases in table: 592
 Number of factors: 3
 Test for independence of all factors:
 $\text{Chisq} = 164.92, \text{df} = 24, \text{p-value} = 5.321\text{e-}23$
 Chi-squared approximation may be incorrect

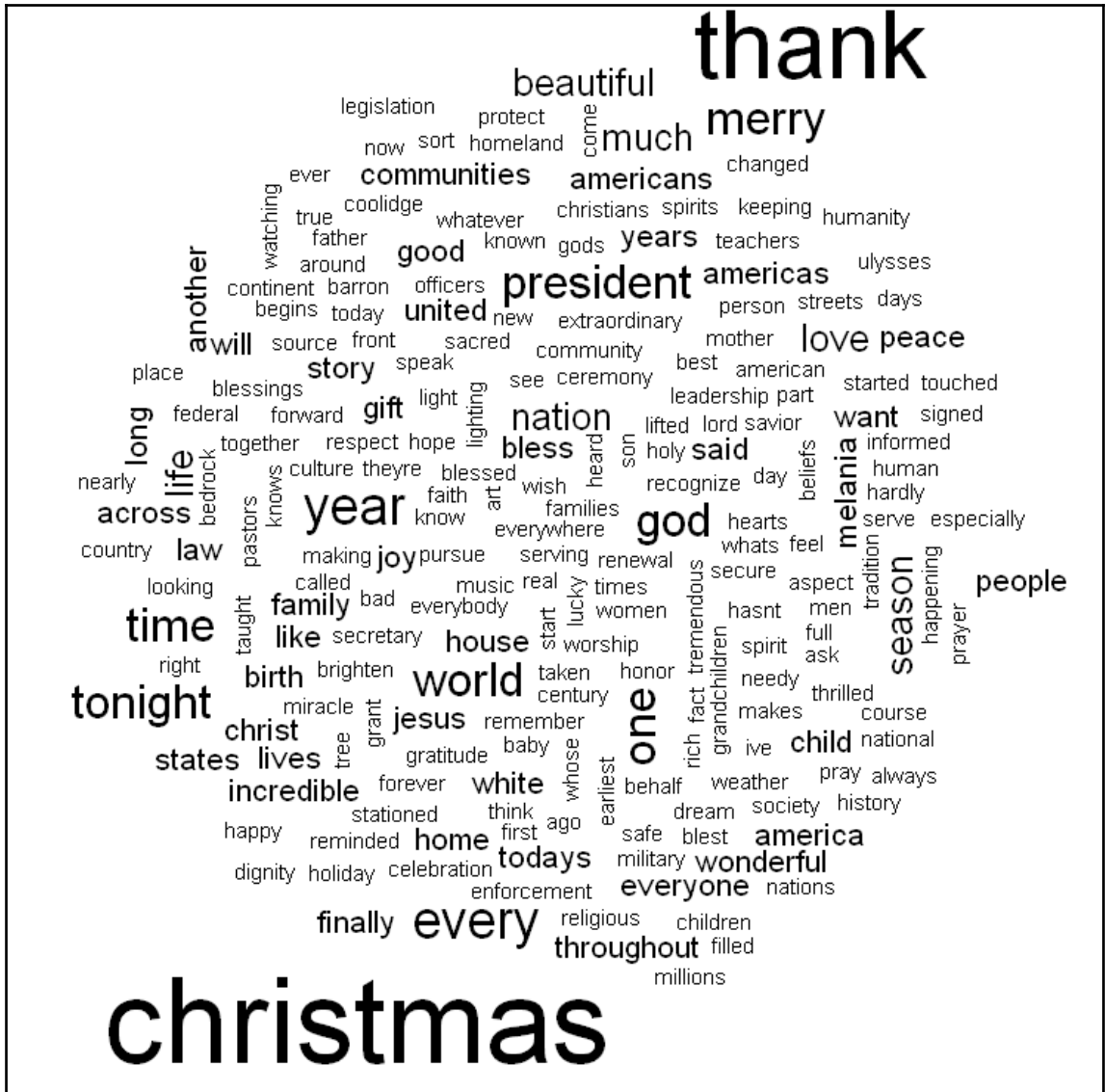
, , Sex = Male

	Eye			
Hair	Brown	Blue	Hazel	Green
Black	32	11	10	3
Brown	53	50	25	15
Red	10	10	7	7
Blond	3	30	5	8

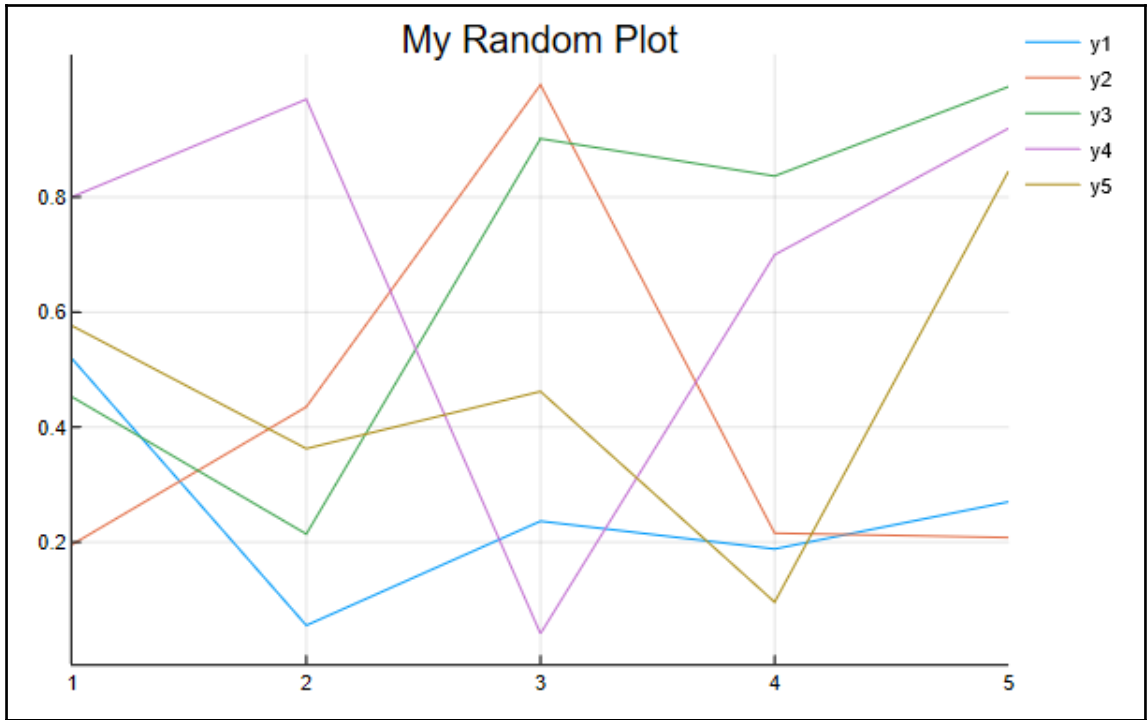
, , Sex = Female

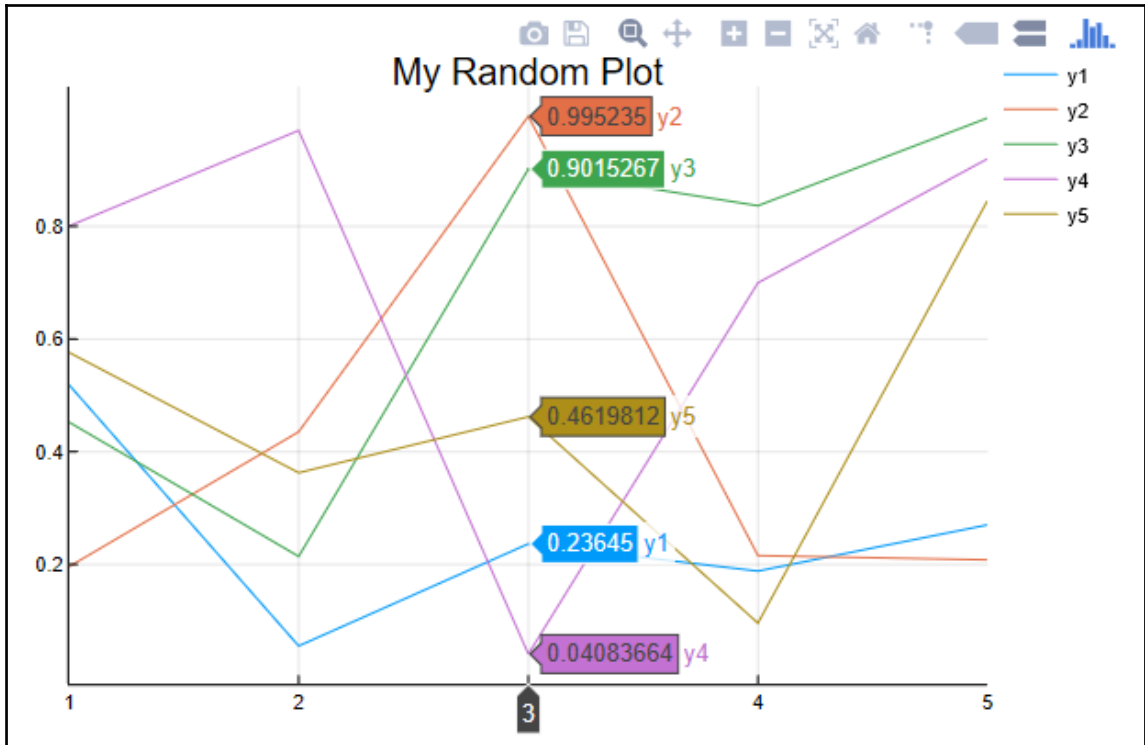
	Eye			
Hair	Brown	Blue	Hazel	Green
Black	36	9	5	2
Brown	66	34	29	14
Red	16	7	7	7
Blond	4	64	5	8





```
Pkg.update()
INFO: Cloning cache of AxisAlgorithms from https://github.com/timholly/AxisAlgorithms.jl.git
INFO: Cloning cache of Calculus from https://github.com/johnmyleswhite/Calculus.jl.git
INFO: Cloning cache of CommonSubexpressions from https://github.com/rdeits/CommonSubexpressions.jl.git
INFO: Cloning cache of Compose from https://github.com/GiovineItalia/Compose.jl.git
INFO: Cloning cache of CoupledFields from https://github.com/Mattriks/CoupledFields.jl.git
INFO: Cloning cache of DiffResults from https://github.com/JuliaDiff/DiffResults.jl.git
INFO: Cloning cache of DiffRules from https://github.com/JuliaDiff/DiffRules.jl.git
```

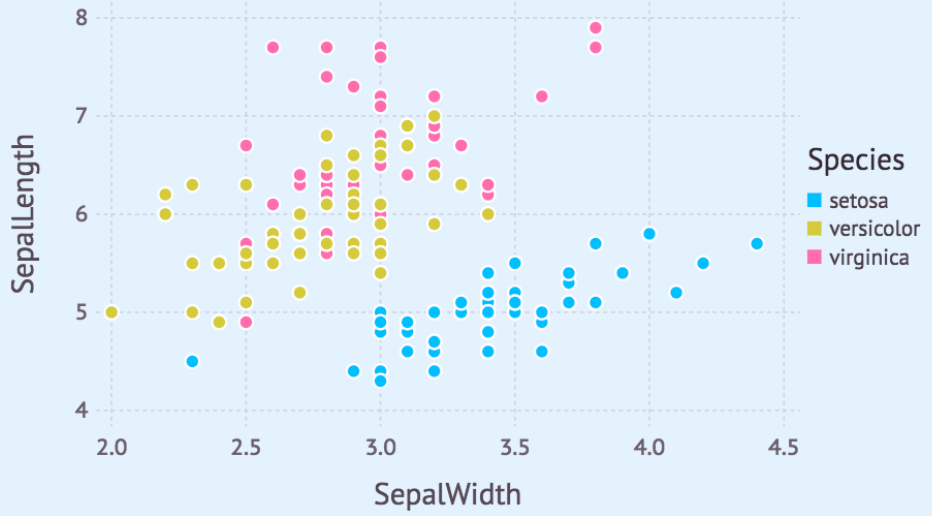





```
In [2]: #Pkg.add("RDatasets")
#Pkg.add("Dataframe")
#Pkg.add("Gadfly")
using RDatasets, DataFrames, Gadfly
set_default_plot_size(5inch, 5inch/golden)
describe(dataset("datasets", "iris"))
plot(dataset("datasets", "iris"), x="SepalWidth", y="SepalLength", color="Species")
```

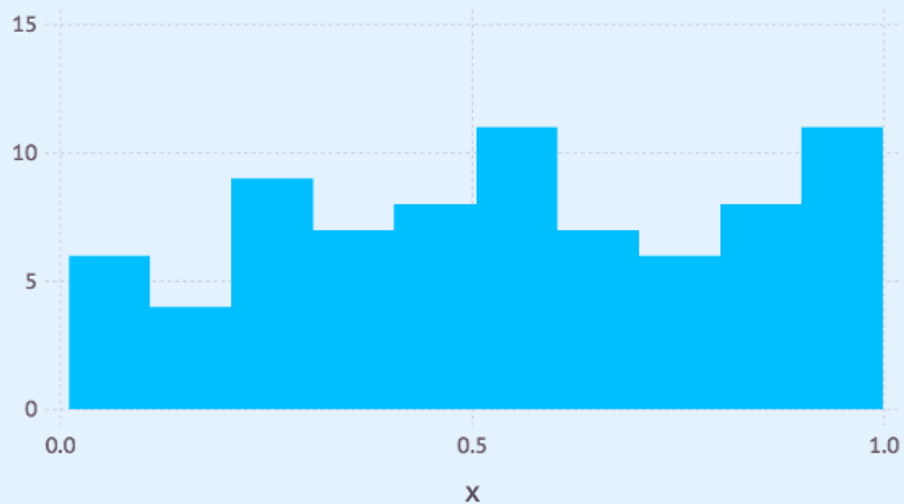
```
SepalLength
Summary Stats:
Mean:          5.843333
Minimum:       4.300000
1st Quartile:  5.100000
Median:        5.800000
3rd Quartile:  6.400000
Maximum:       7.900000
Length:        150
Type:          Float64
Number Missing: 0
% Missing:     0.000000
```

Out[2]:

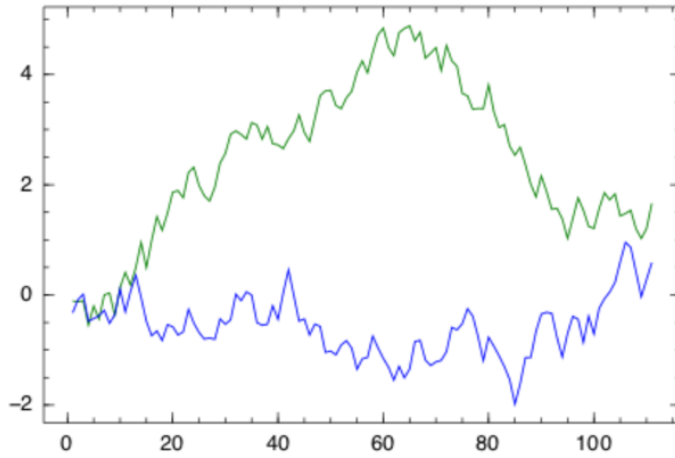


```
In [3]: using Gadfly
        srand(111)
        plot(x=rand(77), Geom.histogram(bincount=10))
```

Out[3]:



```
In [1]: using Winston
        srand(111)
        pl = plot(cumsum(rand(111) .- 0.5), "g", cumsum(rand(111) .- 0.5), "b")
        display(pl)
```



Chapter 5: Working with Widgets

```
In [16]: import ipywidgets as widgets
         from IPython.display import display
```

```
In [17]: my_button = widgets.Button(description='Click My Button')
         display(my_button)
```

Click My Button

```
In [18]: def my_button_clicked(b):
         print("You clicked on My Button")
```

```
In [19]: my_button.on_click(my_button_clicked)
```

```
In [20]: import ipywidgets as widgets
         from IPython.display import display
```

```
In [21]: my_button = widgets.Button(description='Click My Button')
         display(my_button)
```

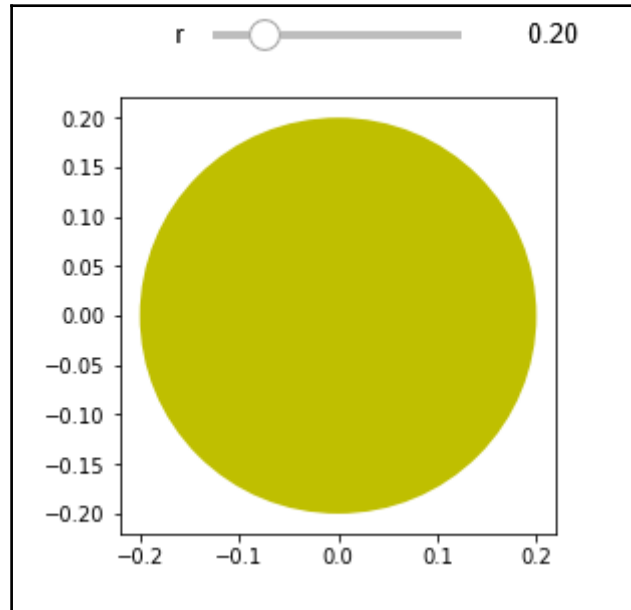
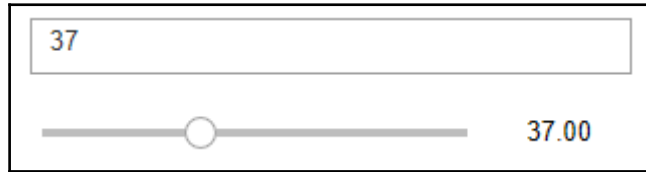
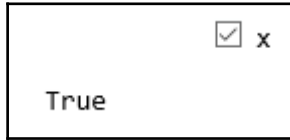
Click My Button

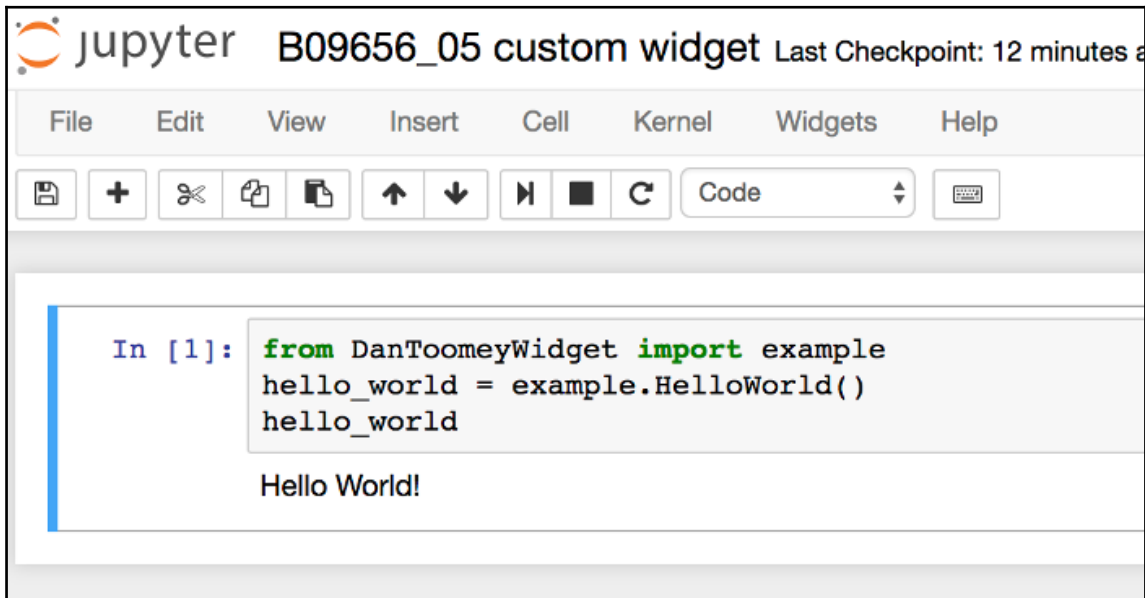
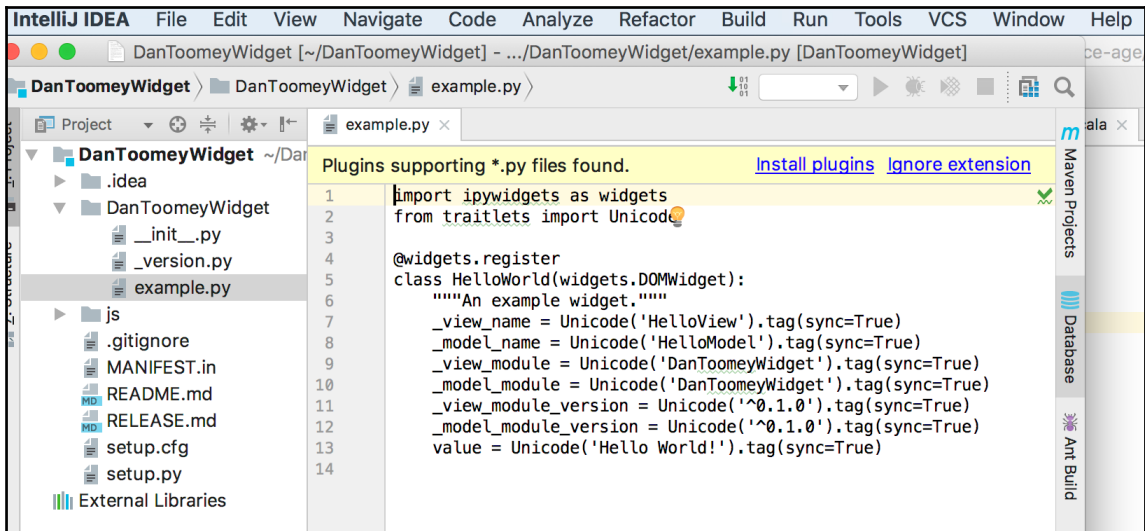
You clicked on My Button

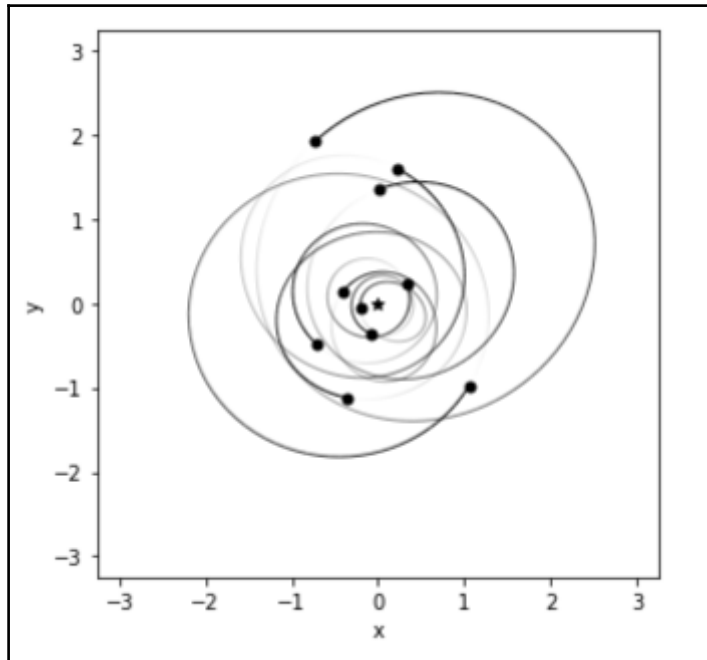
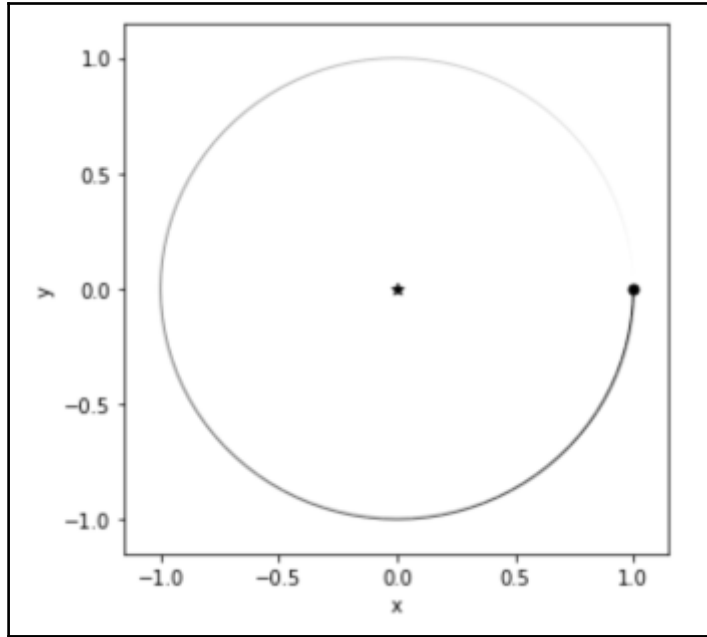
```
In [22]: def my_button_clicked(b):
         print("You clicked on My Button")
```

```
In [23]: my_button.on_click(my_button_clicked)
```

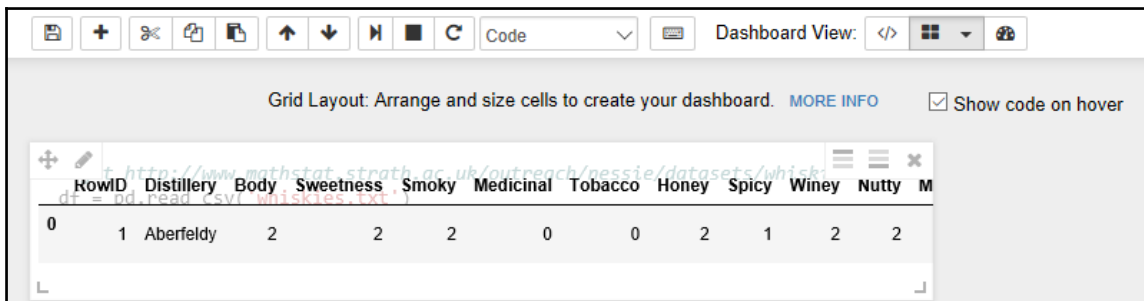
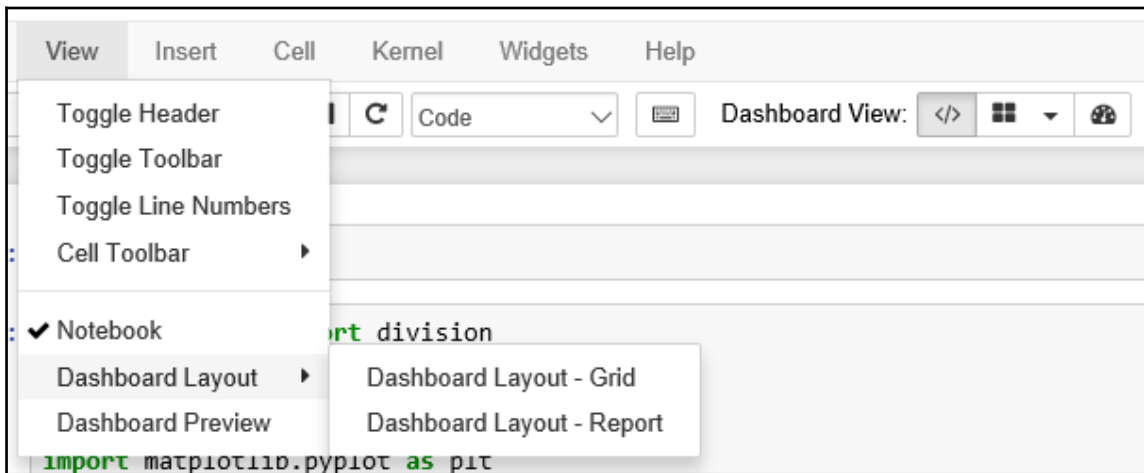
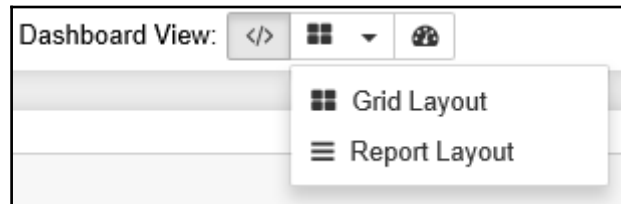
Balloon color: red
 green
 blue

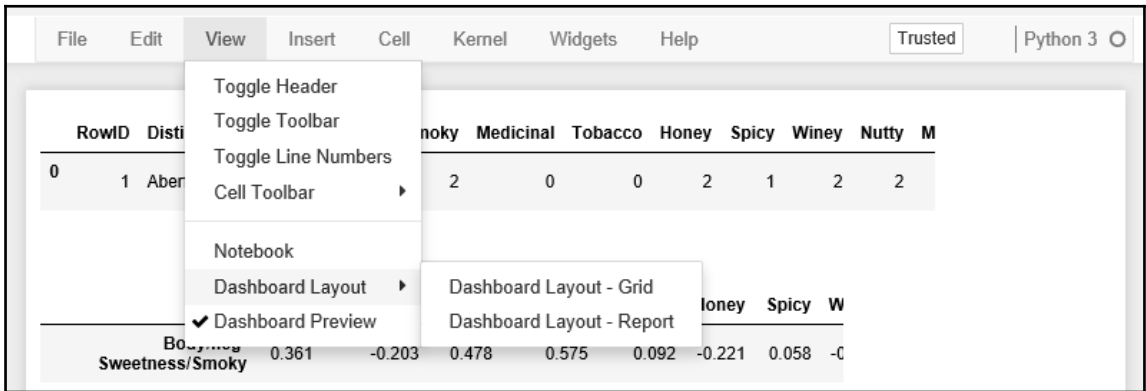




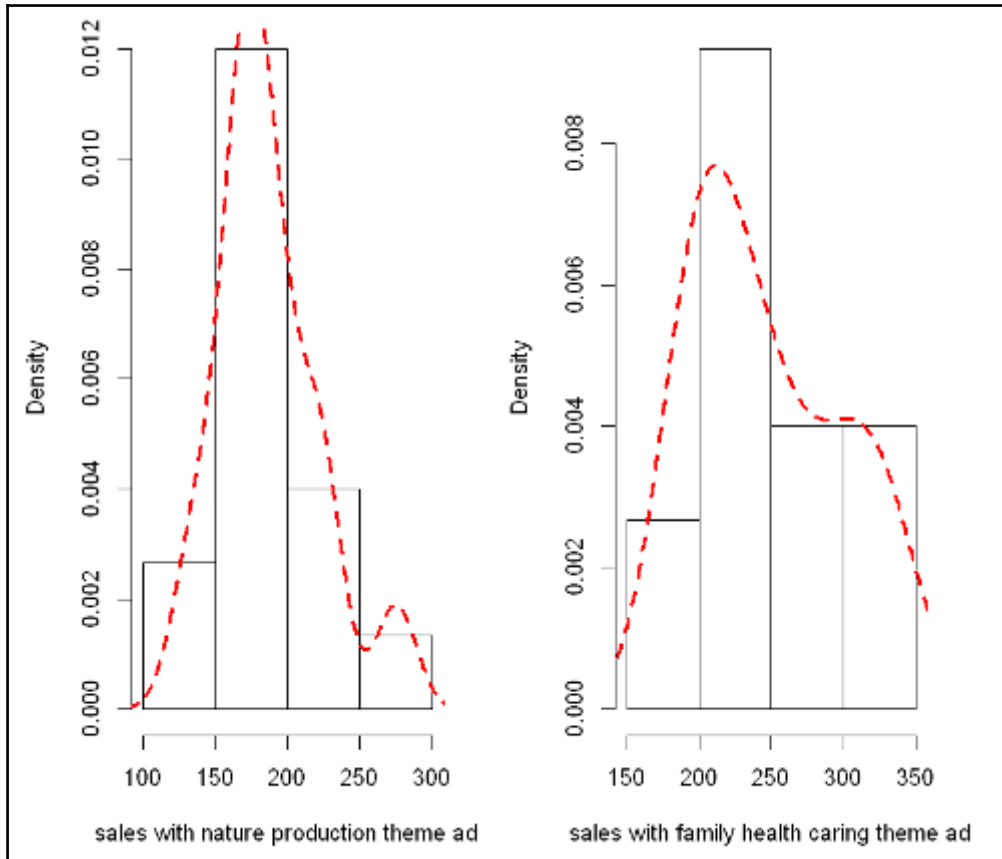


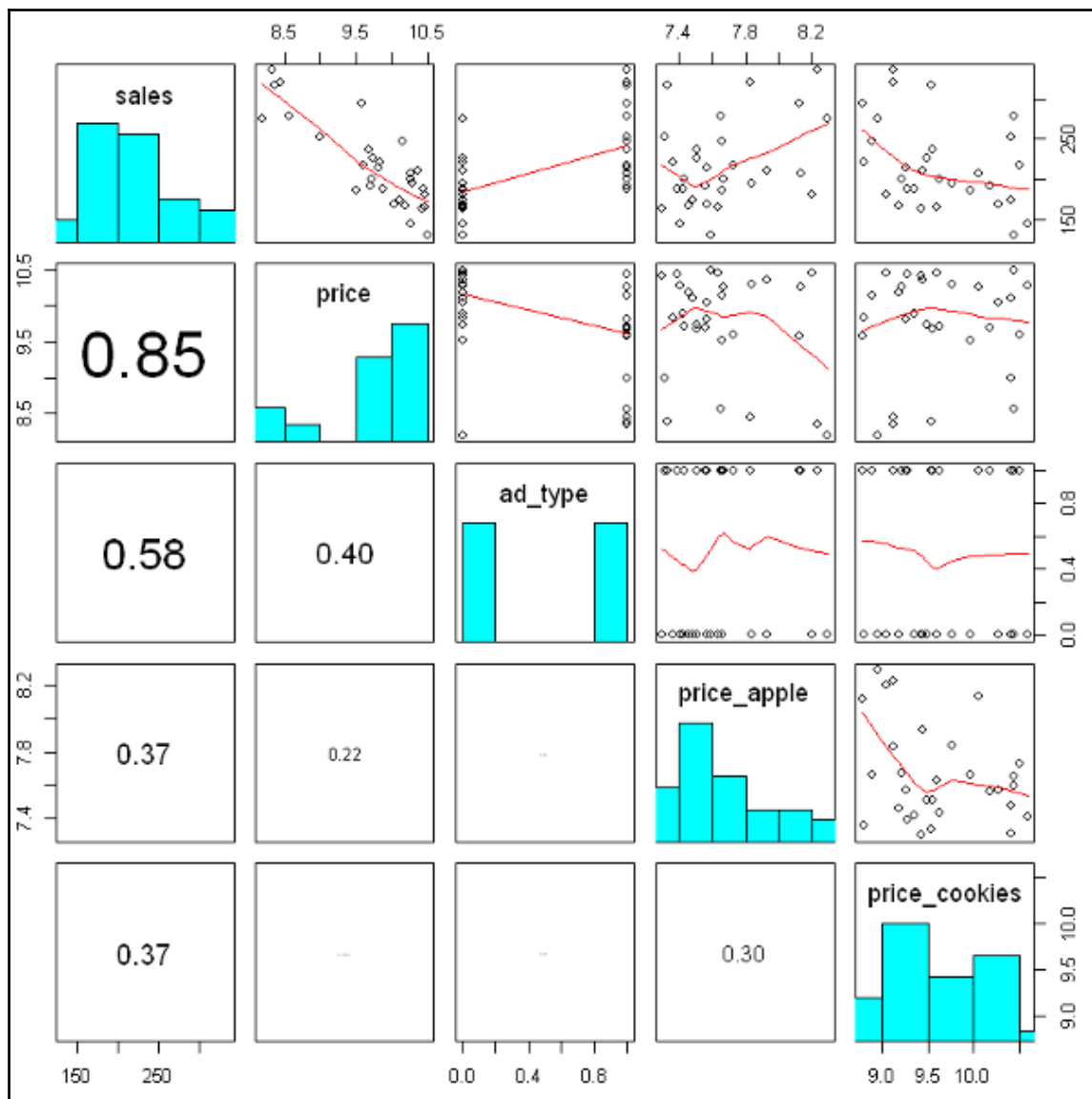
Chapter 6: Jupyter Dashboards





sales	price	ad_type	price_apple	price_cookies
222	9.83	0	7.36	8.80
201	9.72	1	7.43	9.62
247	10.15	1	7.66	8.90
169	10.04	0	7.57	10.26
317	8.38	1	7.33	9.54
227	9.74	0	7.51	9.49





```

Call:
lm(formula = sales ~ price + ad_type + price_apple + price_cookies,
    data = df)

Residuals:
    Min       1Q   Median       3Q      Max
-36.290 -10.488   0.884  10.483  29.471

Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept)   774.813    145.349   5.331 1.59e-05 ***
price         -51.239     5.321  -9.630 6.83e-10 ***
ad_type        29.742     7.249   4.103 0.000380 ***
price_apple    22.089    12.512   1.765 0.089710 .
price_cookies -25.277     6.296  -4.015 0.000477 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 18.2 on 25 degrees of freedom
Multiple R-squared:  0.8974,    Adjusted R-squared:  0.881
F-statistic: 54.67 on 4 and 25 DF,  p-value: 5.318e-12

```

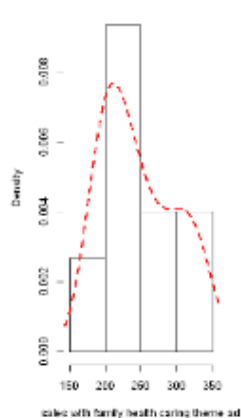
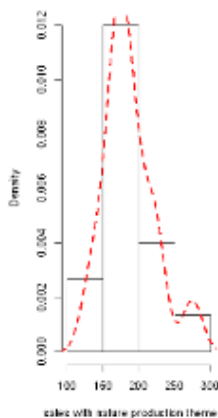
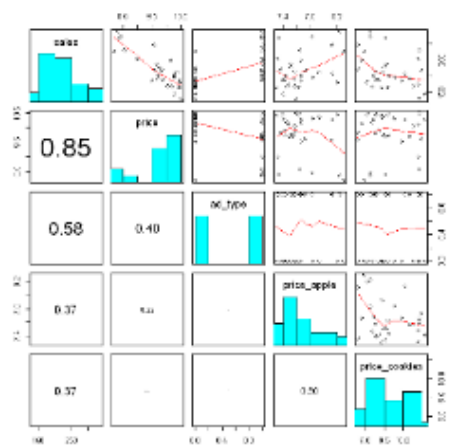
	sales	price	ad_type	price_apple	price_cookie
222	9.83	0	7.36	8.8	
201	9.72	1	7.43	9.6	
247	10.15	1	7.66	8.9	
169	10.04	0	7.57	10.2	
317	8.38	1	7.33	9.5	
227	9.74	0	7.51	9.4	

```
Call:
lm(formula = sales ~ price + ad_type + price_apple + price_cookies,
    data = df)
```

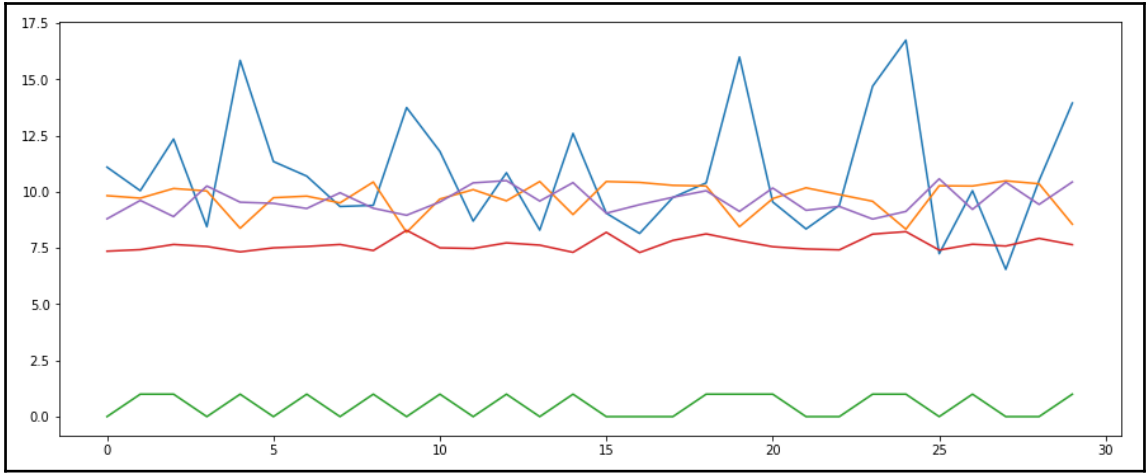
```
Residuals:
    Min       1Q   Median       3Q      Max
-36.298 -10.488  0.884  10.483  29.471
```

```
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  774.813   145.349   5.331 1.59e-05 ***
price        -51.239    5.321  -9.630 6.83e-10 ***
ad_type       29.742    7.249   4.103 0.000380 ***
price_apple   22.089   12.512   1.765 0.089710 .
price_cookies -25.277    6.296  -4.015 0.000477 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Residual standard error: 18.2 on 25 degrees of freedom
Multiple R-squared:  0.8974,    Adjusted R-squared:  0.881
F-statistic: 54.67 on 4 and 25 DF,  p-value: 5.318e-12
```

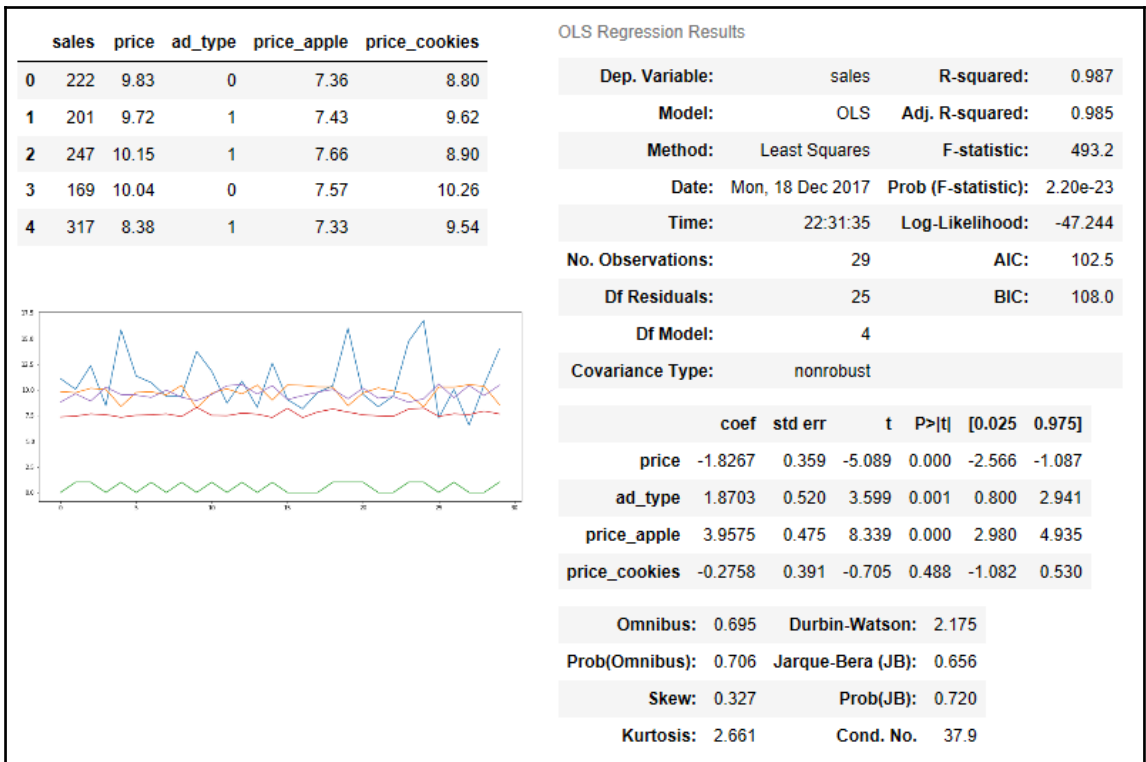


	sales	price	ad_type	price_apple	price_cookies
0	222	9.83	0	7.36	8.80
1	201	9.72	1	7.43	9.62
2	247	10.15	1	7.66	8.90
3	169	10.04	0	7.57	10.26
4	317	8.38	1	7.33	9.54



OLS Regression Results

Dep. Variable:	sales	R-squared:	0.987			
Model:	OLS	Adj. R-squared:	0.985			
Method:	Least Squares	F-statistic:	493.2			
Date:	Mon, 18 Dec 2017	Prob (F-statistic):	2.20e-23			
Time:	22:31:35	Log-Likelihood:	-47.244			
No. Observations:	29	AIC:	102.5			
Df Residuals:	25	BIC:	108.0			
Df Model:	4					
Covariance Type:	nonrobust					
	coef	std err	t	P> t 	[0.025	0.975]
price	-1.8267	0.359	-5.089	0.000	-2.566	-1.087
ad_type	1.8703	0.520	3.599	0.001	0.800	2.941
price_apple	3.9575	0.475	8.339	0.000	2.980	4.935
price_cookies	-0.2758	0.391	-0.705	0.488	-1.082	0.530
Omnibus:	0.695	Durbin-Watson:	2.175			
Prob(Omnibus):	0.706	Jarque-Bera (JB):	0.656			
Skew:	0.327	Prob(JB):	0.720			
Kurtosis:	2.661	Cond. No.	37.9			

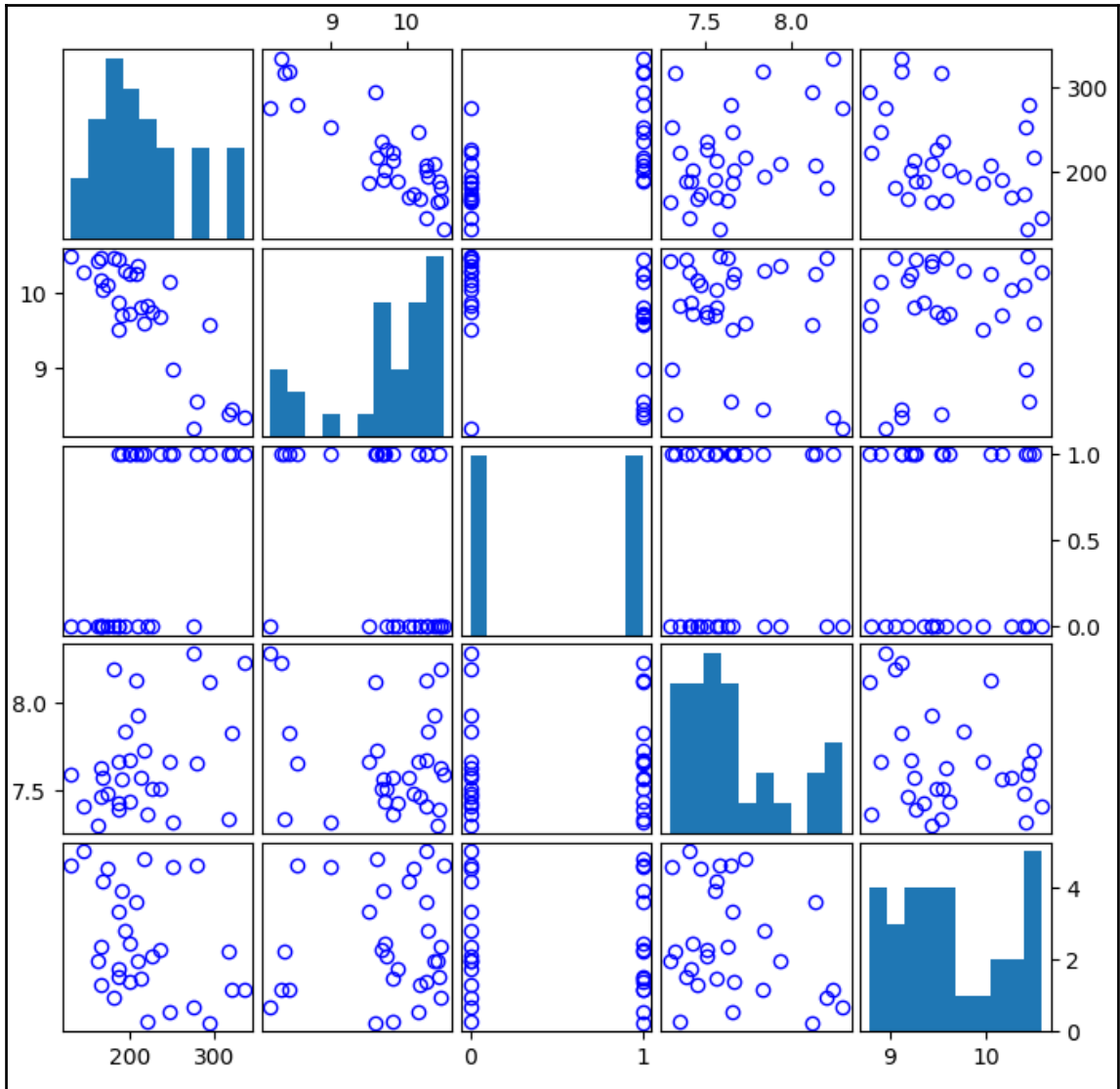


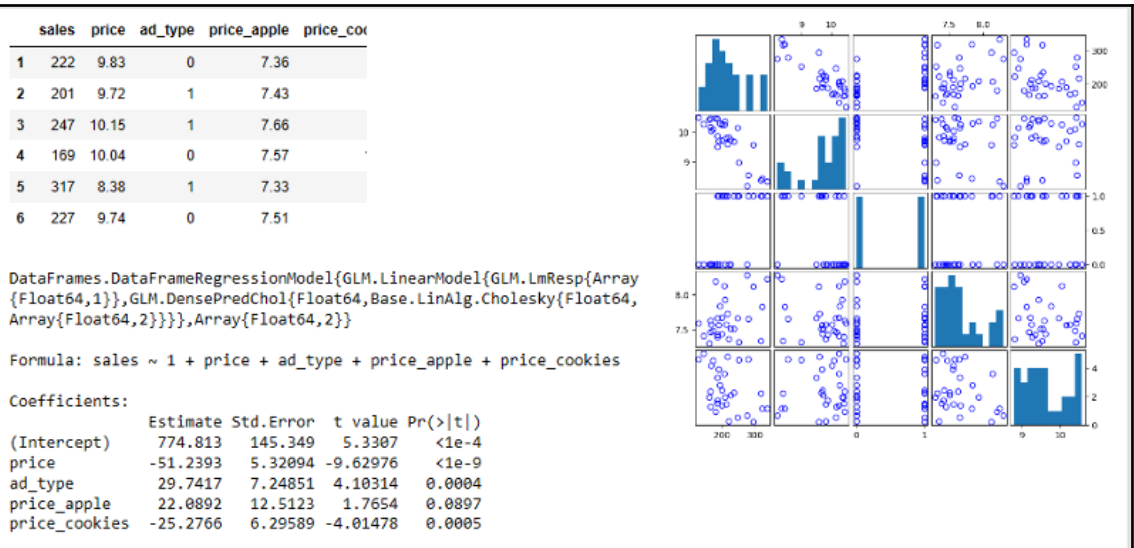
	sales	price	ad_type	price_apple	price_cookies
1	222	9.83	0	7.36	8.8
2	201	9.72	1	7.43	9.62
3	247	10.15	1	7.66	8.9
4	169	10.04	0	7.57	10.26
5	317	8.38	1	7.33	9.54
6	227	9.74	0	7.51	9.49

Formula: sales ~ 1 + price + ad_type + price_apple + price_cookies

Coefficients:

	Estimate	Std.Error	t value	Pr(> t)
(Intercept)	774.813	145.349	5.3307	<1e-4
price	-51.2393	5.32094	-9.62976	<1e-9
ad_type	29.7417	7.24851	4.10314	0.0004
price_apple	22.0892	12.5123	1.7654	0.0897
price_cookies	-25.2766	6.29589	-4.01478	0.0005





```
[
  {
    "name": "lion",
    "avg_weight": "400"
  },
  {
    "name": "tiger",
    "avg_weight": "400"
  },
  {
    "name": "human",
    "avg_weight": "150"
  },
  {
    "name": "elephant",
    "avg_weight": "2000"
  }
]
2000
```

```
Raw data is [ 98, 98.6, 98.4, 98.8, 200, 120, 98.5 ]
Standard Deviation is 35.07
Mean is 116.04
Median is 98.6
Median Abs Deviation is 0.200000000000000284
The outliers of the data set are [ 4, 5, 6 ]
The data set without outliers is [ 98, 98.6, 98.4, 98.8 ]
```

```
[
  {
    "name": "lion",
    "avg_weight": "400"
  },
  {
    "name": "tiger",
    "avg_weight": "400"
  },
  {
    "name": "human",
    "avg_weight": "150"
  },
  {
    "name": "elephant",
    "avg_weight": "2000"
  }
]
2000
```


Raw data is [98, 98.6, 98.4, 98.8, 200, 120, 98.5]
Standard Deviation is 35.07
Mean is 116.04
Median is 98.6
Median Abs Deviation is 0.200000000000000284
The outliers of the data set are [4, 5, 6]
The data set without outliers is [98, 98.6, 98.4, 98.8]

Chapter 7: Sharing Your Code

The screenshot shows the GitHub interface for a repository named 'notebooks' by user 'danieltoomey'. At the top, there's a navigation bar with 'Code', 'Issues 0', 'Pull requests 0', 'Wiki', 'Pulse', 'Graphs', and 'Settings'. Below this, a message states 'No description or website provided. — Edit'. A summary bar indicates '2 commits', '1 branch', and '0 releases'. The main area features a 'Branch: master' dropdown, a 'New pull request' button, and buttons for 'Create new file', 'Upload files', and 'Finish'. A commit history section shows a commit by 'danieltoomey' with files 'README.md' (first commit) and 'Stats Analysis.ipynb' (Add files via upload). Below the commit history, a 'README.md' file is listed. The repository name 'notebooks' is displayed in a large font at the bottom.

Branch: master ▾

notebooks / Stats Analysis.ipynb

 **danieltoomey** Add files via upload

1 contributor

85 lines (84 sloc) | 2.18 KB

```
In [17]: const stats = require("stats-analysis");

var arr = [98, 98.6, 98.4, 98.8, 200, 120, 98.5];

//standard deviation
var my_stddev = stats.stdev(arr).toFixed(2);

//mean
var my_mean = stats.mean(arr).toFixed(2);

//median
var my_median = stats.median(arr);

//median absolute deviation
var my_mad = stats.MAD(arr);

// Outlier detection. Returns indexes of outliers
var my_outliers = stats.indexOfOutliers(arr);
```

```
data(iris)
head(iris)
```

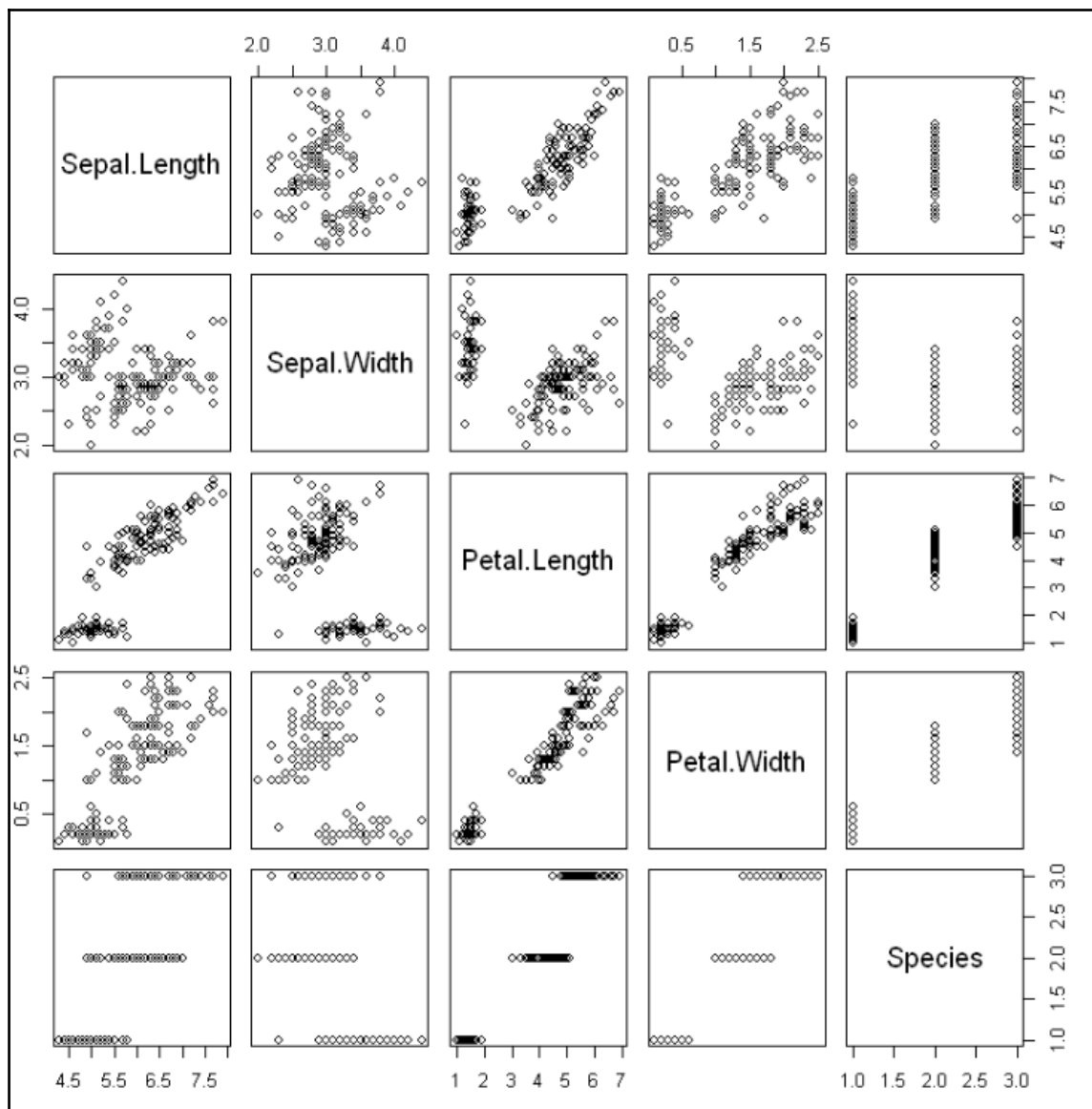
Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa

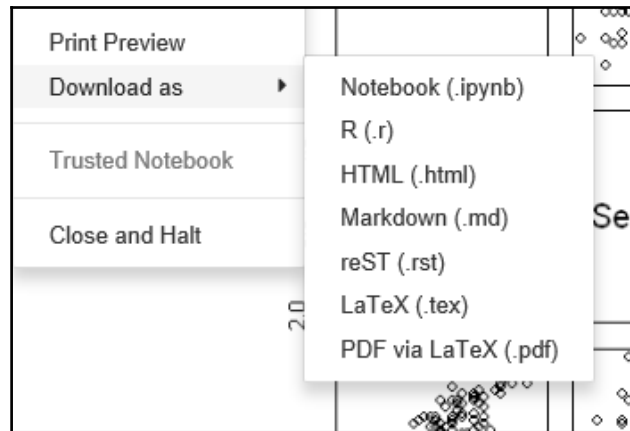
```
summary(iris)
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Min. :4.300	Min. :2.000	Min. :1.000	Min. :0.100
1st Qu.:5.100	1st Qu.:2.800	1st Qu.:1.600	1st Qu.:0.300
Median :5.800	Median :3.000	Median :4.350	Median :1.300
Mean :5.843	Mean :3.057	Mean :3.758	Mean :1.199
3rd Qu.:6.400	3rd Qu.:3.300	3rd Qu.:5.100	3rd Qu.:1.800
Max. :7.900	Max. :4.400	Max. :6.900	Max. :2.500

Species

setosa	:50
versicolor	:50
virginica	:50





C:\Users\Dan\Downloads\B09656_07+r+iris+for+conversions.r
File Edit Search View Encoding Language Settings Tools
B09656_07+r+iris+for+conversions.r

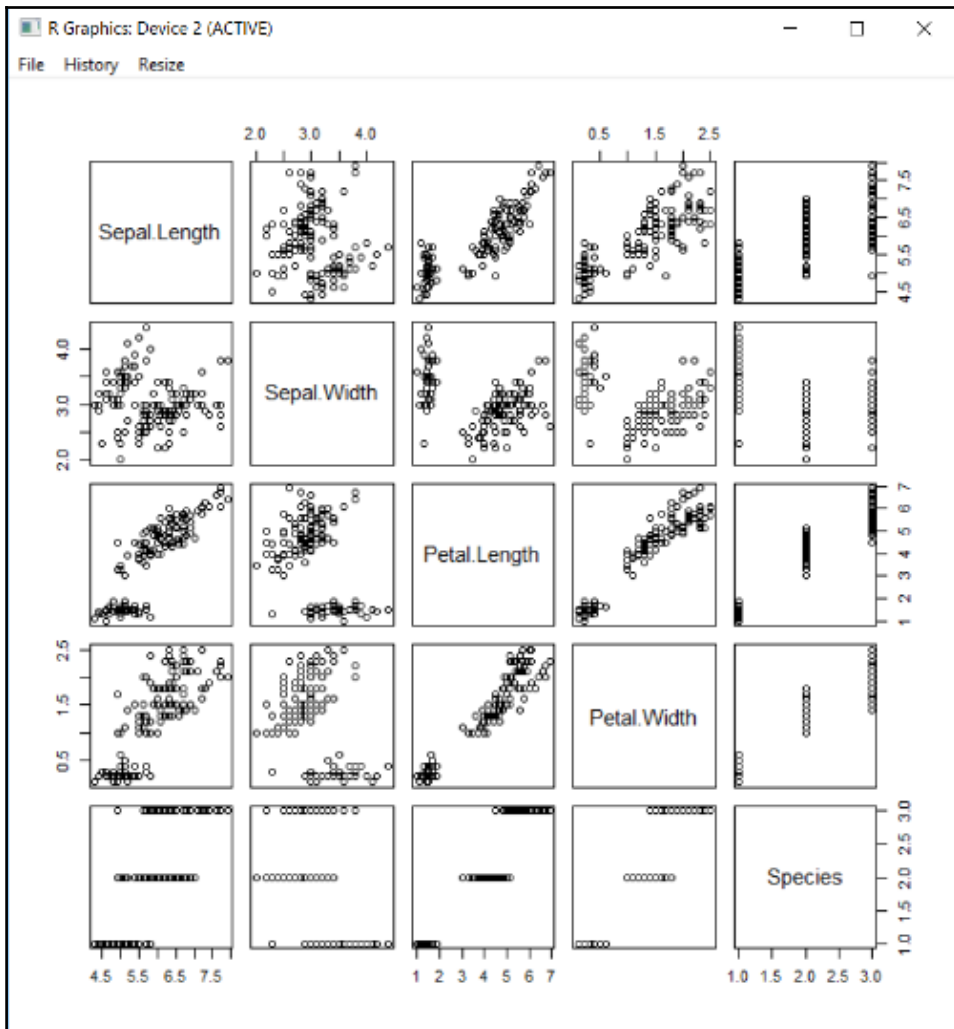
```
1  
2 data(iris)  
3 head(iris)  
4  
5 summary(iris)  
6  
7 plot(iris)  
8
```

The screenshot shows the RStudio interface with a script editor and a console. The script editor contains the following R code:

```
1  
2 data(iris)  
3 head(iris)  
4  
5 summary(iris)  
6  
7 plot(iris)  
8
```

The console shows the output of the commands:

```
> head(iris)  
  Sepal.Length Sepal.width Petal.Length Petal.width Species  
1           5.1           3.5           1.4           0.2 setosa  
2           4.9           3.0           1.4           0.2 setosa  
3           4.7           3.2           1.3           0.2 setosa  
4           4.6           3.1           1.5           0.2 setosa  
5           5.0           3.6           1.4           0.2 setosa  
6           5.4           3.9           1.7           0.4 setosa  
>  
> summary(iris)  
  Sepal.Length      Sepal.width      Petal.Length  
Min.   :4.300      Min.   :2.000      Min.   :1.000  
1st Qu.:5.100      1st Qu.:2.800      1st Qu.:1.600  
Median :5.800      Median :3.000      Median :4.350  
Mean   :5.843      Mean   :3.057      Mean   :3.758  
3rd Qu.:6.400      3rd Qu.:3.300      3rd Qu.:5.100  
Max.   :7.900      Max.   :4.400      Max.   :6.900  
  Petal.width      Species  
Min.   :0.100      setosa   :50  
1st Qu.:0.300      versicolor:50  
Median :1.300      virginica :50  
Mean   :1.199  
3rd Qu.:1.800  
Max.   :2.500  
>  
> plot(iris)  
>
```



B09656_07 r iris for convi x

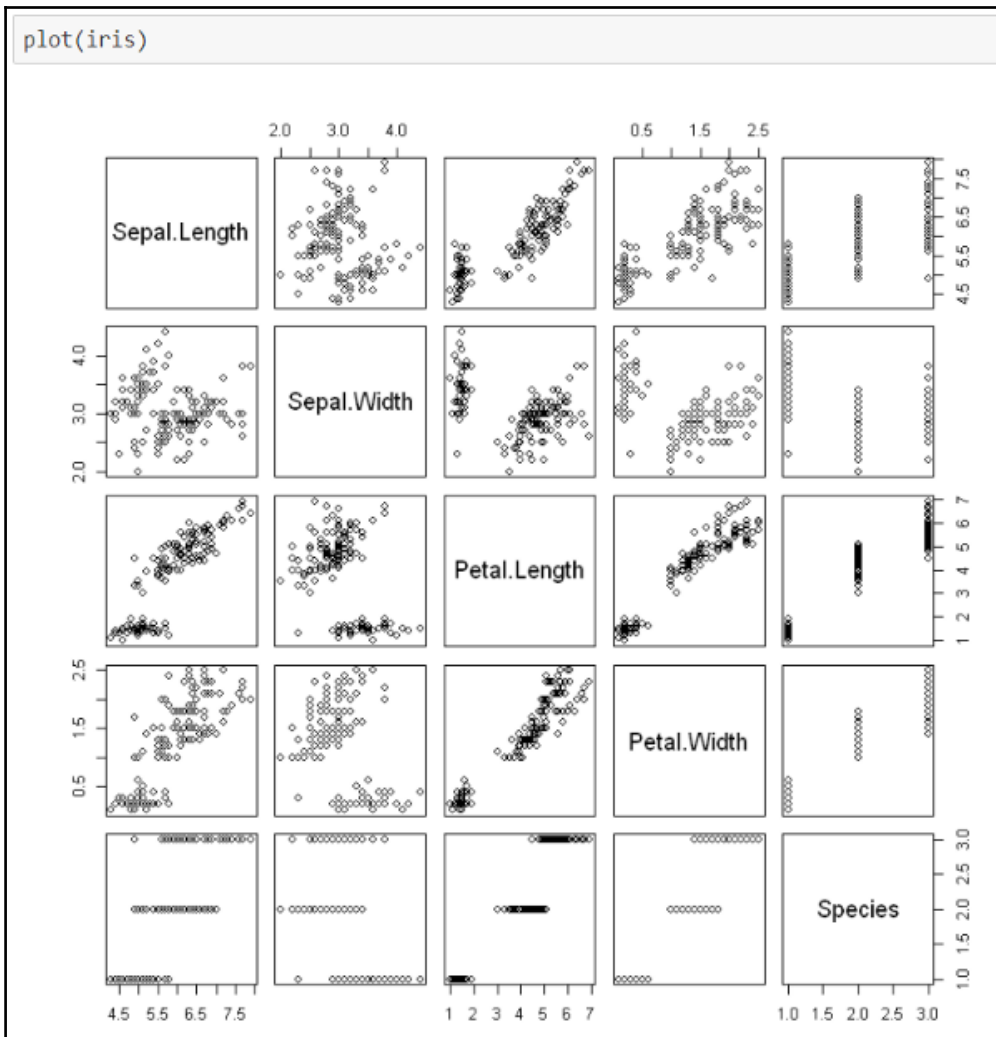
file:///C:/Users/Dan/Downloads/B09656_07+r+iris+for+conversions.html

In [4]: `data(iris)`
`head(iris)`

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa

In [5]: `summary(iris)`

```
Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
Min.   :4.300      Min.   :2.000      Min.   :1.000      Min.   :0.100
1st Qu.:5.100      1st Qu.:2.800      1st Qu.:1.600      1st Qu.:0.300
Median :5.800      Median :3.000      Median :4.350      Median :1.300
Mean   :5.843      Mean   :3.057      Mean   :3.758      Mean   :1.199
3rd Qu.:6.400      3rd Qu.:3.300      3rd Qu.:5.100      3rd Qu.:1.800
Max.   :7.900      Max.   :4.400      Max.   :6.900      Max.   :2.500
Species
setosa   :50
versicolor:50
virginica :50
```




```

11841 <div class="output_text output_subarea ">
11842 <pre> Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
11843 Min.      :4.300    Min.      :2.000    Min.      :1.000    Min.      :0.100
11844 1st Qu.:5.100    1st Qu.:2.800    1st Qu.:1.600    1st Qu.:0.300
11845 Median :5.800    Median :3.000    Median :4.350    Median :1.300
11846 Mean   :5.843    Mean   :3.057    Mean   :3.758    Mean   :1.199
11847 3rd Qu.:6.400    3rd Qu.:3.300    3rd Qu.:5.100    3rd Qu.:1.800
11848 Max.   :7.900    Max.   :4.400    Max.   :6.900    Max.   :2.500
11849      Species
11850      setosa      :50
11851      versicolor:50
11852      virginica  :50
11853
11854
11855 </pre>
11856
11857 <div class="output_png output_subarea ">
11858 <img src="data:image/png;base64,iVBORw0KGgoAAAANSUHEUgAAA0gAAANICAMAAADKOT
11859 jIyampqnp6eysrK9vb3Hx8fQ0NDZ2dnh4eHp6enw8PD///QFLu4AAAACXBIWXMAABJ0AAAS
11860 dAHeZh94AAAgaELEQVR4nO1dizaJOax0n9PbPdP8/99uEsCnLF8CBKl6uxkuS4XtSmVB0mYC
11861 AGAY5mgCAHAFQEgAIAAICQAEACEBgAAgJAAQAIQEAAKakABAABASAAgAQgIAAUBIACAAcAkA
11862 BAAhYAAICQAEACEBAACgJAAQAAQEgAIAEICAFAFASAAgAAgJAAQAIQGAACAKABAahaQAAoCQ

```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa


```
summary(iris)
```


Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Min. :4.300	Min. :2.000	Min. :1.000	Min. :0.100
1st Qu.:5.100	1st Qu.:2.800	1st Qu.:1.600	1st Qu.:0.300

```

1  ```R
2  data(iris)
3  head(iris)
4  ```
5  <table>
6  <thead><tr><th scope=col>Sepal.Length</th><th scope=col>Sepal.Width</th><th scope=col>Petal.Length</th><th scope=col>Petal.Width</th><th scope=col>Species</th></tr>
7  <tbody>
8  <tr><td>5.1 </td><td>3.5 </td><td>1.4 </td><td>0.2 </td><td>setosa</td></tr>
9  <tr><td>4.9 </td><td>3.0 </td><td>1.4 </td><td>0.2 </td><td>setosa</td></tr>
10 <tr><td>4.7 </td><td>3.2 </td><td>1.3 </td><td>0.2 </td><td>setosa</td></tr>
11 <tr><td>4.6 </td><td>3.1 </td><td>1.5 </td><td>0.2 </td><td>setosa</td></tr>
12 <tr><td>5.0 </td><td>3.6 </td><td>1.4 </td><td>0.2 </td><td>setosa</td></tr>
13 <tr><td>5.4 </td><td>3.9 </td><td>1.7 </td><td>0.4 </td><td>setosa</td></tr>
14 </tbody>
15 </table>
16
17 ```R
18 summary(iris)
19 ```
20      Sepal.Length   Sepal.Width   Petal.Length   Petal.Width
21  Min.   :4.300   Min.   :2.000   Min.   :1.000   Min.   :0.100
22  1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300
23  Median :5.800   Median :3.000   Median :4.350   Median :1.300
24  Mean   :5.843   Mean   :3.057   Mean   :3.758   Mean   :1.199
25  3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800
26  Max.   :7.900   Max.   :4.400   Max.   :6.900   Max.   :2.500
27
28      Species
29  setosa   :50
30  versicolor:50
31  virginica :50
32
33 ```R
34 plot(iris)
35 ```
36 ! [png] (output_2_0.png)

```

```
data(iris)
head(iris)

Sepal.Length Sepal.Width Petal.Length Petal.Width Species
5.1          3.5         1.4         0.2         setosa
4.9          3.0         1.4         0.2         setosa
4.7          3.2         1.3         0.2         setosa
4.6          3.1         1.5         0.2         setosa
5.0          3.6         1.4         0.2         setosa
5.4          3.9         1.7         0.4         setosa

summary(iris)

Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
Min.   :4.300    Min.   :2.000    Min.   :1.000    Min.   :0.100
1st Qu.:5.100    1st Qu.:2.800    1st Qu.:1.600    1st Qu.:0.300
Median :5.800    Median :3.000    Median :4.350    Median :1.300
Mean   :5.843    Mean   :3.057    Mean   :3.758    Mean   :1.199
3rd Qu.:6.400    3rd Qu.:3.300    3rd Qu.:5.100    3rd Qu.:1.800
Max.   :7.900    Max.   :4.400    Max.   :6.900    Max.   :2.500

Species
setosa      :50
versicolor:50
```

```

.. code:: r

    data(iris)
    head(iris)

.. raw:: html

    <table>
      <thead><tr><th scope=col>Sepal.Length</th><th scope=col>Sepal.Width</th><th
scope=col>Petal.Length</th><th scope=col>Petal.Width</th><th scope=col>Species</th>
</tr></thead>
      <tbody>
        <tr><td>5.1 </td><td>3.5 </td><td>1.4 </td><td>0.2
</td><td>setosa</td></tr>
        <tr><td>4.9 </td><td>3.0 </td><td>1.4 </td><td>0.2
</td><td>setosa</td></tr>
        <tr><td>4.7 </td><td>3.2 </td><td>1.3 </td><td>0.2
</td><td>setosa</td></tr>
        <tr><td>4.6 </td><td>3.1 </td><td>1.5 </td><td>0.2
</td><td>setosa</td></tr>
        <tr><td>5.0 </td><td>3.6 </td><td>1.4 </td><td>0.2
</td><td>setosa</td></tr>
        <tr><td>5.4 </td><td>3.9 </td><td>1.7 </td><td>0.4
</td><td>setosa</td></tr>
      </tbody>
    </table>

.. code:: r

    summary(iris)

```

B09656_07 r iris for conversions

January 1, 2018

```
In [4]: data(iris)
        head(iris)
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa

```
In [5]: summary(iris)
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Min. :4.300	Min. :2.000	Min. :1.000	Min. :0.100
1st Qu.:5.100	1st Qu.:2.800	1st Qu.:1.600	1st Qu.:0.300
Median :5.800	Median :3.000	Median :4.350	Median :1.300
Mean :5.843	Mean :3.057	Mean :3.758	Mean :1.199
3rd Qu.:6.400	3rd Qu.:3.300	3rd Qu.:5.100	3rd Qu.:1.800
Max. :7.900	Max. :4.400	Max. :6.900	Max. :2.500

Species

setosa	:50
versicolor	:50
virginica	:50

```

281
282
283 ▾ \begin{Verbatim}[commandchars=\\\{\}]
284 {\color{incolor}In [{\color{incolor}4}]:} data\PY{p}{()iris\PY{p}{}}
285 \PY{k+kp}{head}\PY{p}{()iris\PY{p}{}}
286 \end{Verbatim}
287
288
289 ▾ \begin{tabular}{r|lllll}
290 Sepal.Length & Sepal.Width & Petal.Length & Petal.Width & Species\\
291 \hline
292 5.1 & 3.5 & 1.4 & 0.2 & setosa\\
293 4.9 & 3.0 & 1.4 & 0.2 & setosa\\
294 4.7 & 3.2 & 1.3 & 0.2 & setosa\\
295 4.6 & 3.1 & 1.5 & 0.2 & setosa\\
296 5.0 & 3.6 & 1.4 & 0.2 & setosa\\
297 5.4 & 3.9 & 1.7 & 0.4 & setosa\\
298 \end{tabular}
299
300
301
302 ▾ \begin{Verbatim}[commandchars=\\\{\}]
303 {\color{incolor}In [{\color{incolor}5}]:} \PY{k+kp}{summary}\PY{p}{()iris\PY{p}{}}
304 \end{Verbatim}
305
306
307
308 ▾ \begin{verbatim}
309 Sepal.Length Sepal.Width Petal.Length Petal.Width
310 Min. :4.300 Min. :2.000 Min. :1.000 Min. :0.100
311 1st Qu.:5.100 1st Qu.:2.800 1st Qu.:1.600 1st Qu.:0.300
312 Median :5.800 Median :3.000 Median :4.350 Median :1.300
313 Mean :5.843 Mean :3.057 Mean :3.758 Mean :1.199
314 3rd Qu.:6.400 3rd Qu.:3.300 3rd Qu.:5.100 3rd Qu.:1.800
315 Max. :7.900 Max. :4.400 Max. :6.900 Max. :2.500
316 Species
317 setosa :50
318 versicolor:50
319 virginica :50
320
321
322 \end{verbatim}
323
324
325
326 ▾ \begin{Verbatim}[commandchars=\\\{\}]
327 {\color{incolor}In [{\color{incolor}6}]:} plot\PY{p}{()iris\PY{p}{}}
328 \end{Verbatim}
329

```

500 : Internal Server Error

The error was:

```
nbconvert failed: xelatex not found on PATH, if you have not installed xelatex you may need to do so. Find further instructions at https://nbconvert.readthedocs.io/en/latest/install.html#installing-tex.
```

B09656_07 r iris for conversions

January 1, 2018

```
In [4]: data(iris)
        head(iris)
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa

```
In [5]: summary(iris)
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Min. :4.300	Min. :2.000	Min. :1.000	Min. :0.100
1st Qu.:5.100	1st Qu.:2.800	1st Qu.:1.600	1st Qu.:0.300
Median :5.800	Median :3.000	Median :4.350	Median :1.300
Mean :5.843	Mean :3.057	Mean :3.758	Mean :1.199
3rd Qu.:6.400	3rd Qu.:3.300	3rd Qu.:5.100	3rd Qu.:1.800
Max. :7.900	Max. :4.400	Max. :6.900	Max. :2.500

Species

setosa	:50
versicolor	:50
virginica	:50

Chapter 8: Multiuser Jupyter

```
In [1]: from ipywidgets import interact  
  
def myfunction(x):  
    return x  
  
interact(myfunction, x="Hello World")
```

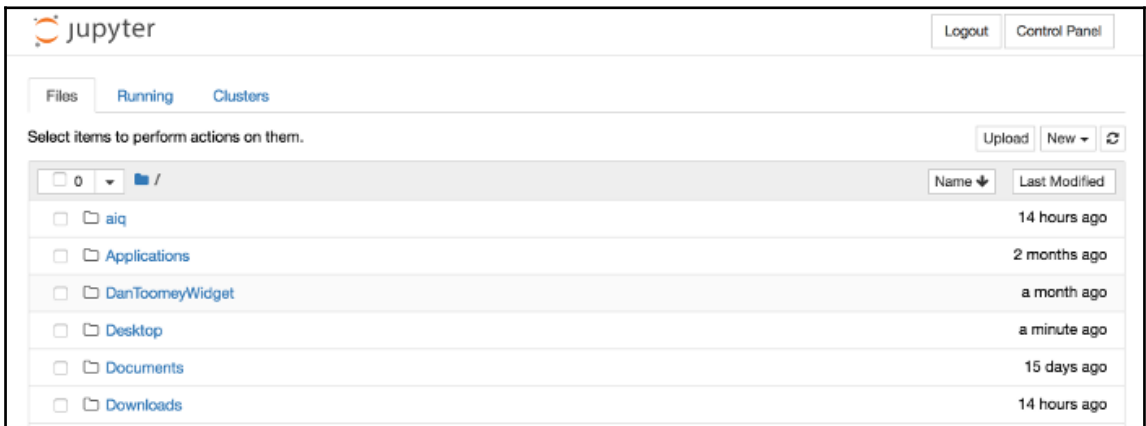
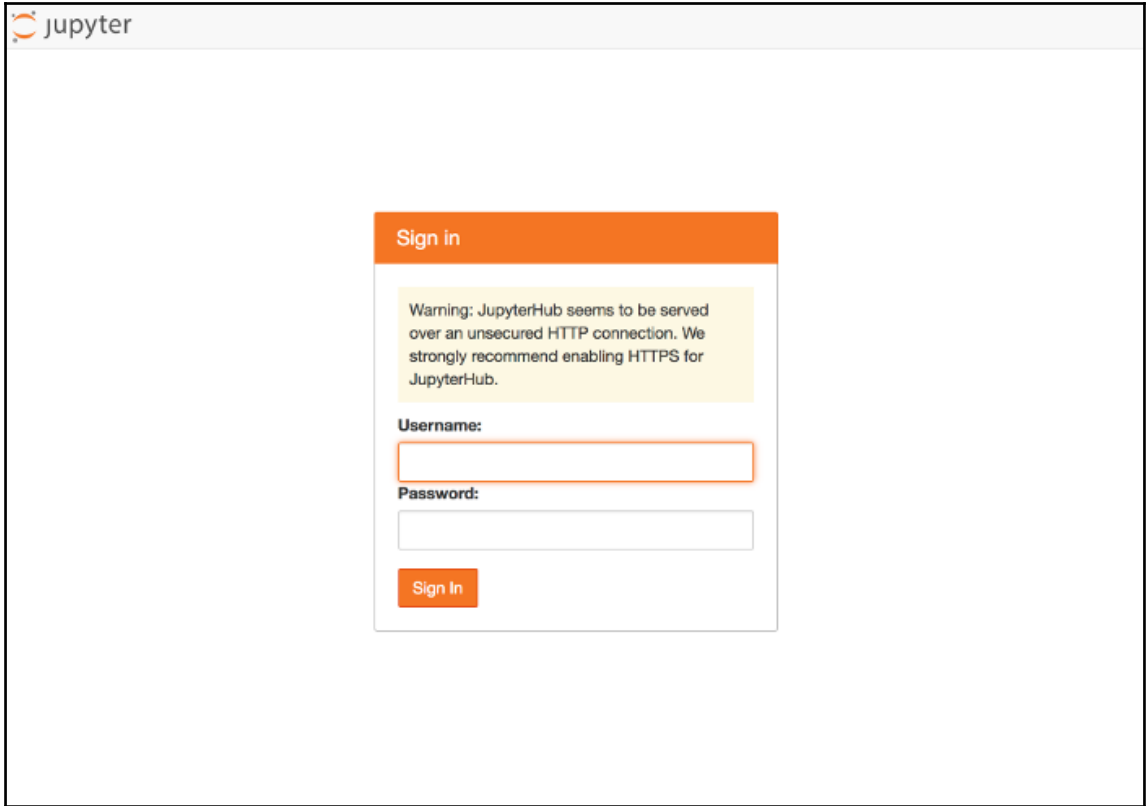
x

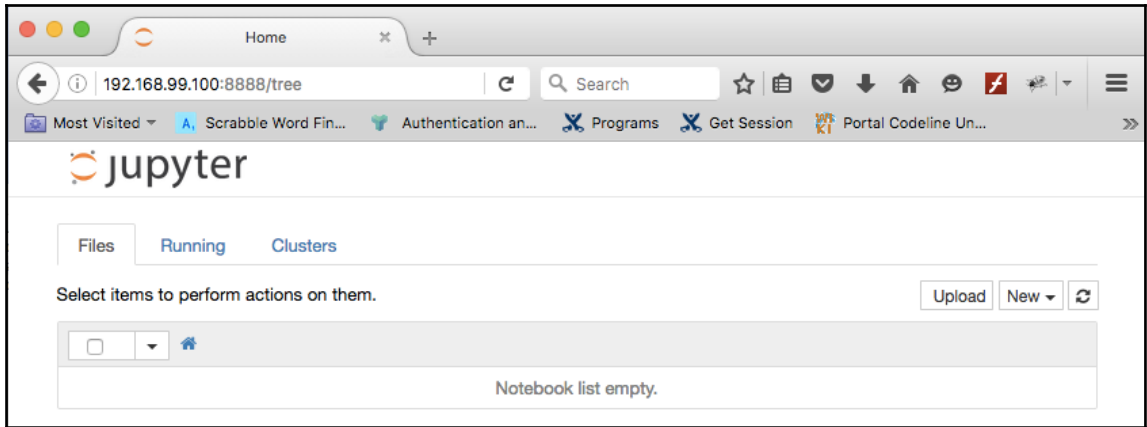
'Hello World'

```
In [1]: from ipywidgets import interact  
  
def myfunction(x):  
    return x  
  
interact(myfunction, x="Hello World")
```

x

'Hello Dan'





Chapter 9: Interacting with Big Data

```
In [9]: import pyspark
        if not 'sc' in globals():
            sc = pyspark.SparkContext()

        text_file = sc.textFile('B09656_09_word_count.ipynb')
        counts = text_file.flatMap(lambda line: line.split(" ")) \
            .map(lambda word: (word, 1)) \
            .reduceByKey(lambda a, b: a + b)

        for x in counts.collect():
            print (x)

(' ', 291)
('"cells":', 1)
([' ', 4)
('"code"', 1)
('8,', 1)
('"outputs":', 1)
('"NameError"', 1)
('"evaluate":', 1)
('"name', 1)
```

```

In [1]: import pyspark
        if not 'sc' in globals():
            sc = pyspark.SparkContext()

        text_file = sc.textFile('B09656_09_word_count.ipynb')
        counts = text_file.flatMap(lambda line: line.split(" ")) \
            .map(lambda word: (word, 1)) \
            .reduceByKey(lambda a, b: a + b) \
            .sortByKey()

        for x in counts.collect():
            print (x)

(' ', 929)
('\"', 5)
('\"(\\"', 1)
('\"(\'(most', 1)
('\"(\'(word,\"', 1)
('\"(\'(x)\"\", 1)
('\"(\'+\", 1)
('\"(\'.map(lambda', 1)
('\"(\'.reduceByKey(lambda', 1)
('\"(\'1)\"', 1)

```

```
In [3]: import pyspark
if not 'sc' in globals():
    sc = pyspark.SparkContext()

textFile = sc.textFile('access_log')
print(textFile.count(), "access records")

gets = textFile.filter(lambda line: "GET" in line)
print(gets.count(), "GETs")

posts = textFile.filter(lambda line: "POST" in line)
print(posts.count(), "POSTs")

other = textFile.subtract(gets).subtract(posts)
print(other.count(), "Other")

for x in other.collect():
    print(x)

1546 access records
1525 GETs
14 POSTs
7 Other
h194n2f1s308o1033.telia.com - - [09/Mar/2004:13:49:05 -0800] "-" 408 -
64.246.94.141 - - [10/Mar/2004:16:31:19 -0800] "HEAD /twiki/bin/view/Main/SpamAssassinDeletin
g HTTP/1.1" 200 0
206-15-133-154.dialup.ziplink.net - - [11/Mar/2004:16:33:23 -0800] "HEAD /twiki/bin/view/Mai
n/SpamAssassinDeleting HTTP/1.1" 200 0
```

```
In [1]: import pyspark
if not 'sc' in globals():
    sc = pyspark.SparkContext()

#check if a number is prime
def isprime(n):
    # must be positive
    n = abs(int(n))

    # 2 or more
    if n < 2:
        return False

    # 2 is the only even prime number
    if n == 2:
        return True
    if not n & 1:
        return False

    # range starts with 3 and only needs to go up the square root of n
    # for all odd numbers
    for x in range(3, int(n**0.5)+1, 2):
        if n % x == 0:
            return False
    return True

nums = sc.parallelize(range(1000000))

# Compute the number of primes in the RDD
print(nums.filter(isprime).count())

78498
```

```

In [1]: import pyspark
        if not 'sc' in globals():
            sc = pyspark.SparkContext()

        sentences = sc.textFile('B09656_09_article.txt') \
            .glom() \
            .map(lambda x: " ".join(x)) \
            .flatMap(lambda x: x.split("."))
        print(sentences.count(), "sentences")

        bigrams = sentences.map(lambda x:x.split()) \
            .flatMap(lambda x: [(x[i],x[i+1]),1] for i in range(0,len(x)-1)])
        print(bigrams.count(), "bigrams")

        frequent_bigrams = bigrams.reduceByKey(lambda x,y:x+y) \
            .map(lambda x:(x[1],x[0])) \
            .sortByKey(False)
        frequent_bigrams.take(10)

140 sentences
2448 bigrams

Out[1]: [(11, ('of', 'the')),
         (8, ('in', 'the')),
         (6, ('to', 'the')),
         (6, ('for', 'the')),
         (6, ('of', 'college')),
         (5, ('more', 'than')),
         (5, ('I'm', 'cynical')),
         (5, ('cynical', 'about')),
         (5, ('The', 'vast')),
         (5, ('vocational', 'education'))]

```

```

syears = sorted(years.items(), key=operator.itemgetter(1), reverse=True)
soccupations = sorted(occupations.items(), key=operator.itemgetter(1), reverse=True)
squests = sorted(quests.items(), key=operator.itemgetter(1), reverse=True)

print(syears[:5])
print(soccupations[:5])
print(squests[:5])

got error
[('2000', 169), ('1999', 166), ('2001', 16)]
[('actor', 122), ('actress', 73), ('comedian', 19), ('film actress', 7), ('NA', 7)]
[('Bob Dole', 5), ('Pamela Anderson', 3), ('Adam Sandler', 3), ('Richard Belzer', 3), ('Tracy Ullman', 2)]

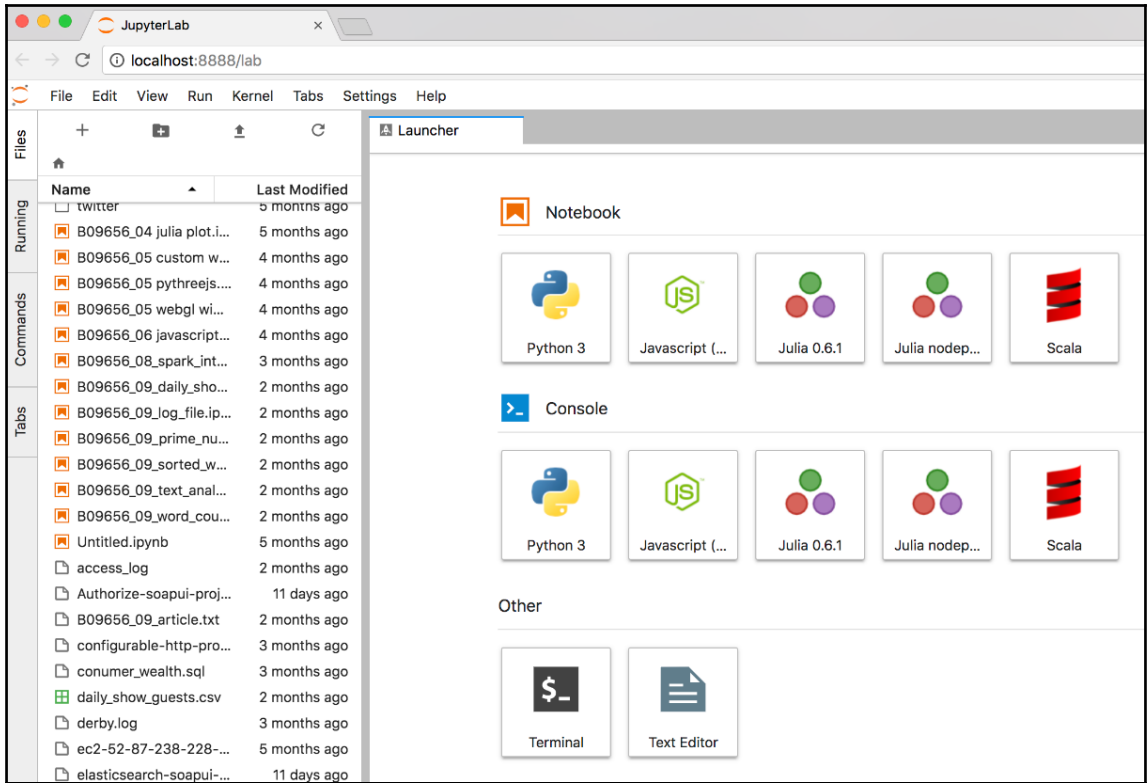
```

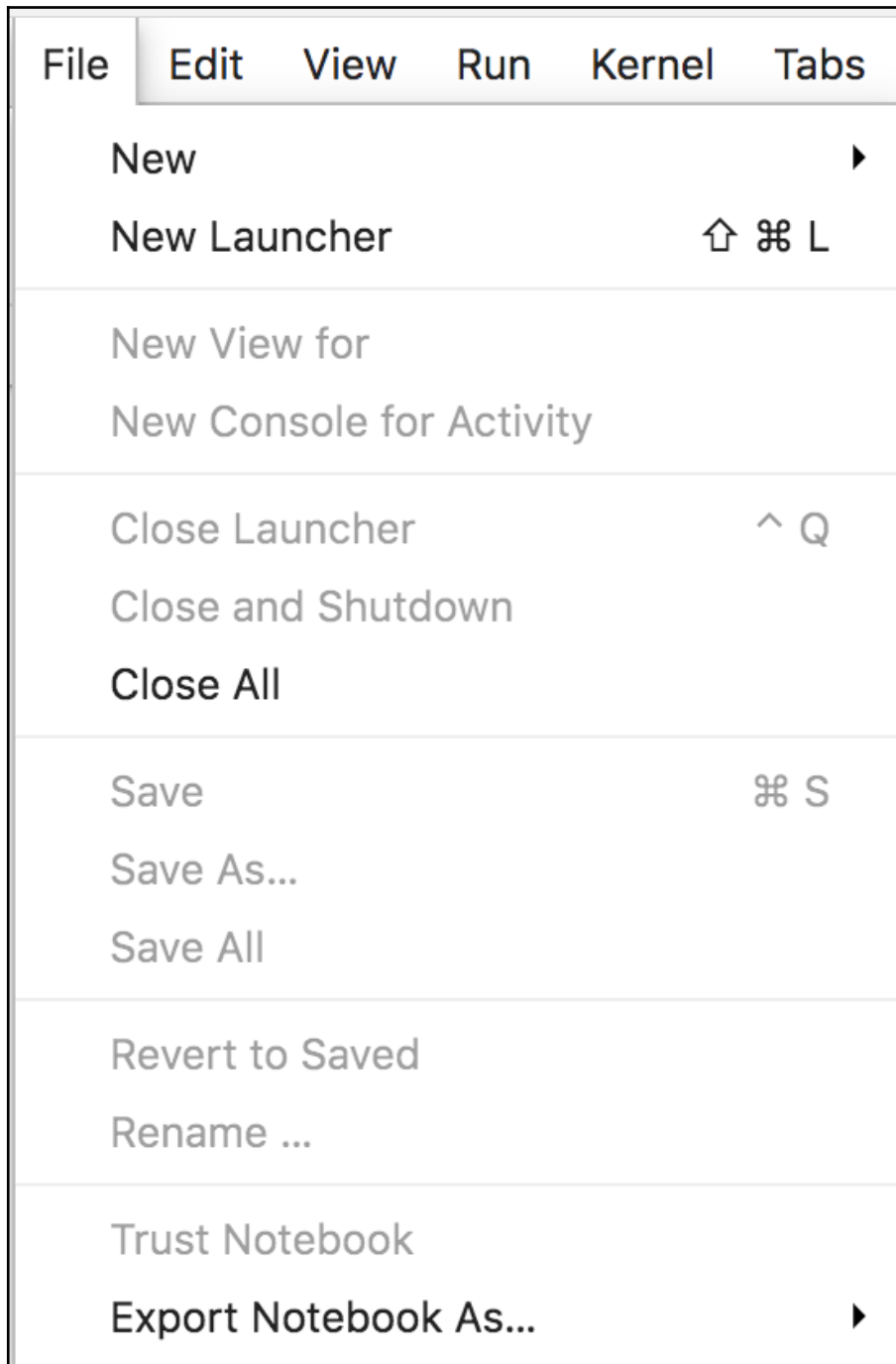
Chapter 11: Jupyter Labs

```

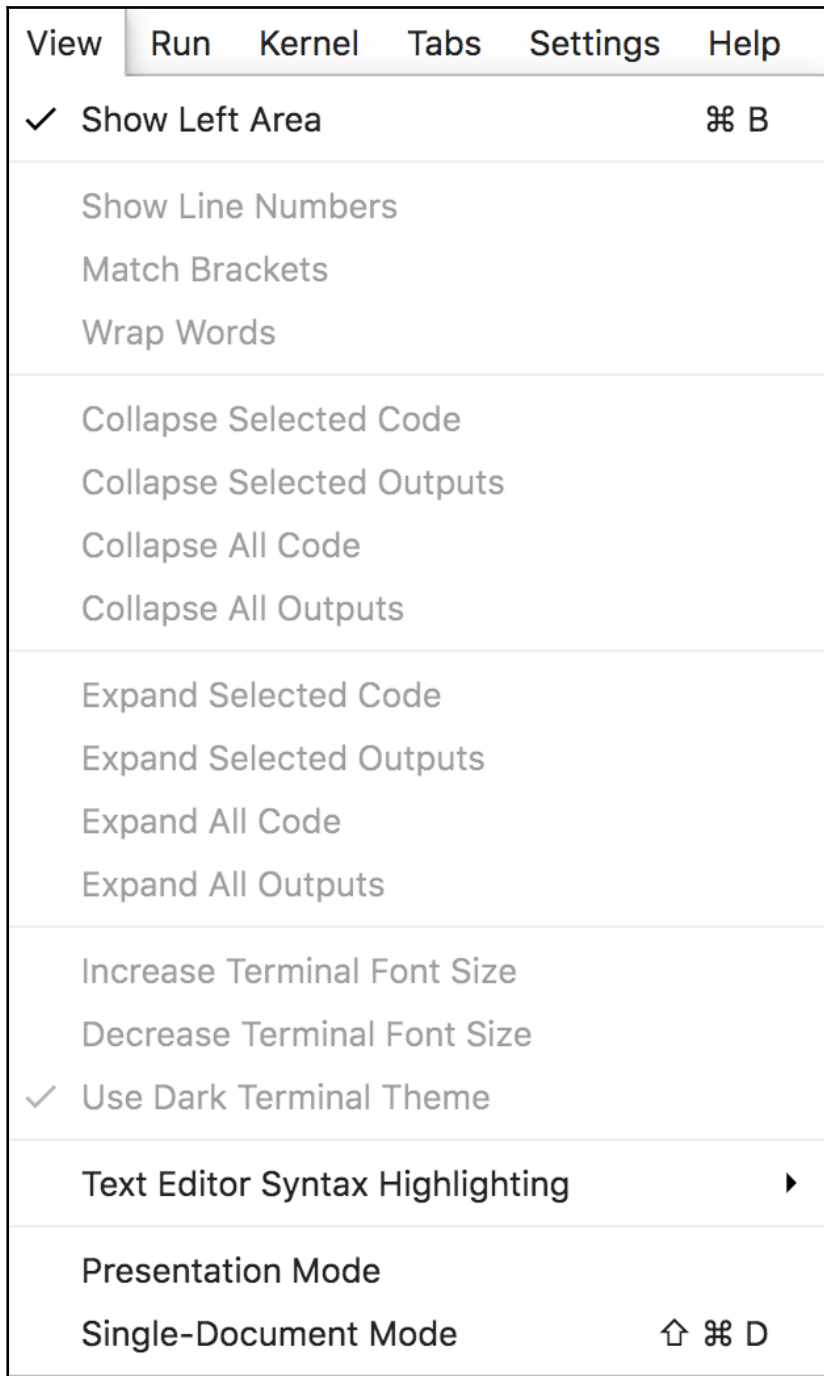
ToomeyD — jupyter-lab - python — 122x24
[I 09:14:32.187 LabApp] JupyterLab application directory is /anaconda3/share/jupyter/lab
[I 09:14:32.196 LabApp] Serving notebooks from local directory: /Users/ToomeyD
[I 09:14:32.196 LabApp] 0 active kernels
[I 09:14:32.196 LabApp] The Jupyter Notebook is running at:
[I 09:14:32.196 LabApp] http://localhost:8888/?token=1c41d43a4ee3046e430c4b042149313a4505282dde5ea820
[I 09:14:32.196 LabApp] Use Control-C to stop this server and shut down all kernels (twice to skip confirmation).
[C 09:14:32.199 LabApp]

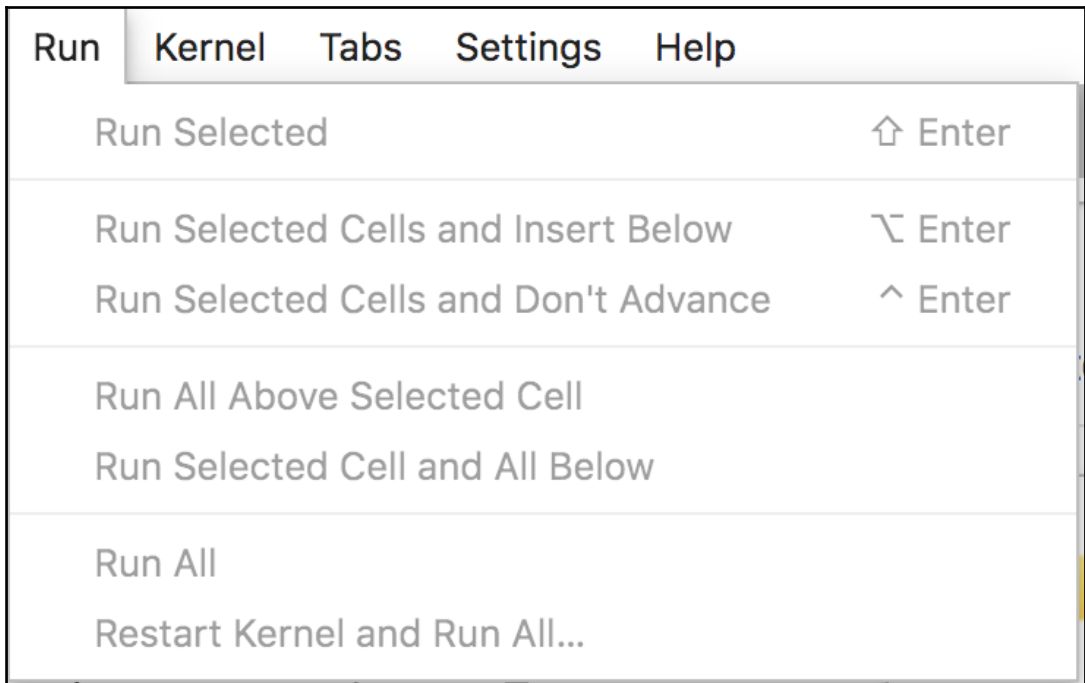
Copy/paste this URL into your browser when you connect for the first time,
to login with a token:
  http://localhost:8888/?token=1c41d43a4ee3046e430c4b042149313a4505282dde5ea820
[I 09:14:32.370 LabApp] Accepting one-time-token-authenticated connection from ::1
[I 09:14:33.963 LabApp] Build is up to date
[I 09:40:29.415 LabApp] Kernel started: 230c9560-e29a-4d13-b838-062f2c774320
Starting kernel event loops.
[I 09:40:33.590 LabApp] Adapting to protocol v5.0 for kernel 230c9560-e29a-4d13-b838-062f2c774320
[I 09:40:33.590 LabApp] Adapting to protocol v5.0 for kernel 230c9560-e29a-4d13-b838-062f2c774320
[I 09:40:33.604 LabApp] Adapting to protocol v5.0 for kernel 230c9560-e29a-4d13-b838-062f2c774320
[I 09:41:16.038 LabApp] Kernel started: 7e0ebc59-ce4e-4ec0-bf30-56d18c22e73b
[I 09:41:17.547 LabApp] Adapting to protocol v5.1 for kernel 7e0ebc59-ce4e-4ec0-bf30-56d18c22e73b
[I 09:41:17.547 LabApp] Adapting to protocol v5.1 for kernel 7e0ebc59-ce4e-4ec0-bf30-56d18c22e73b
[I 09:41:48.342 LabApp] Starting buffering for 7e0ebc59-ce4e-4ec0-bf30-56d18c22e73b:3b803240d6983a9565607ca9297337ba
[I 09:41:48.342 LabApp] Starting buffering for 230c9560-e29a-4d13-b838-062f2c774320:bdc938daa9c2881b293ef6cce5ed4569
```



Edit	View	Run	Kernel	Tabs	Settings
Undo					^ Z
Redo					^ ⬆ Z
Undo Cell Operation					Z
Redo Cell Operation					⬆ Z
Cut Cells					X
Copy Cells					C
Paste Cells Below					V
Paste Cells Above					
Paste Cells and Replace					
Delete Cells					D, D
Select All Cells					⌘ A
Deselect All Cells					
Move Cells Up					
Move Cells Down					
Split Cell					^ ⬆ -
Merge Selected Cells					⬆ M
Clear					
Clear All					
Find...					
Find and Replace...					





Kernel	Tabs	Settings	Help
Interrupt Kernel			I, I
Restart Kernel...			0, 0
Restart Kernel and Clear...			
Restart Kernel and Run All...			
Shutdown Kernel			
Shutdown All Kernels...			
Change Kernel...			

