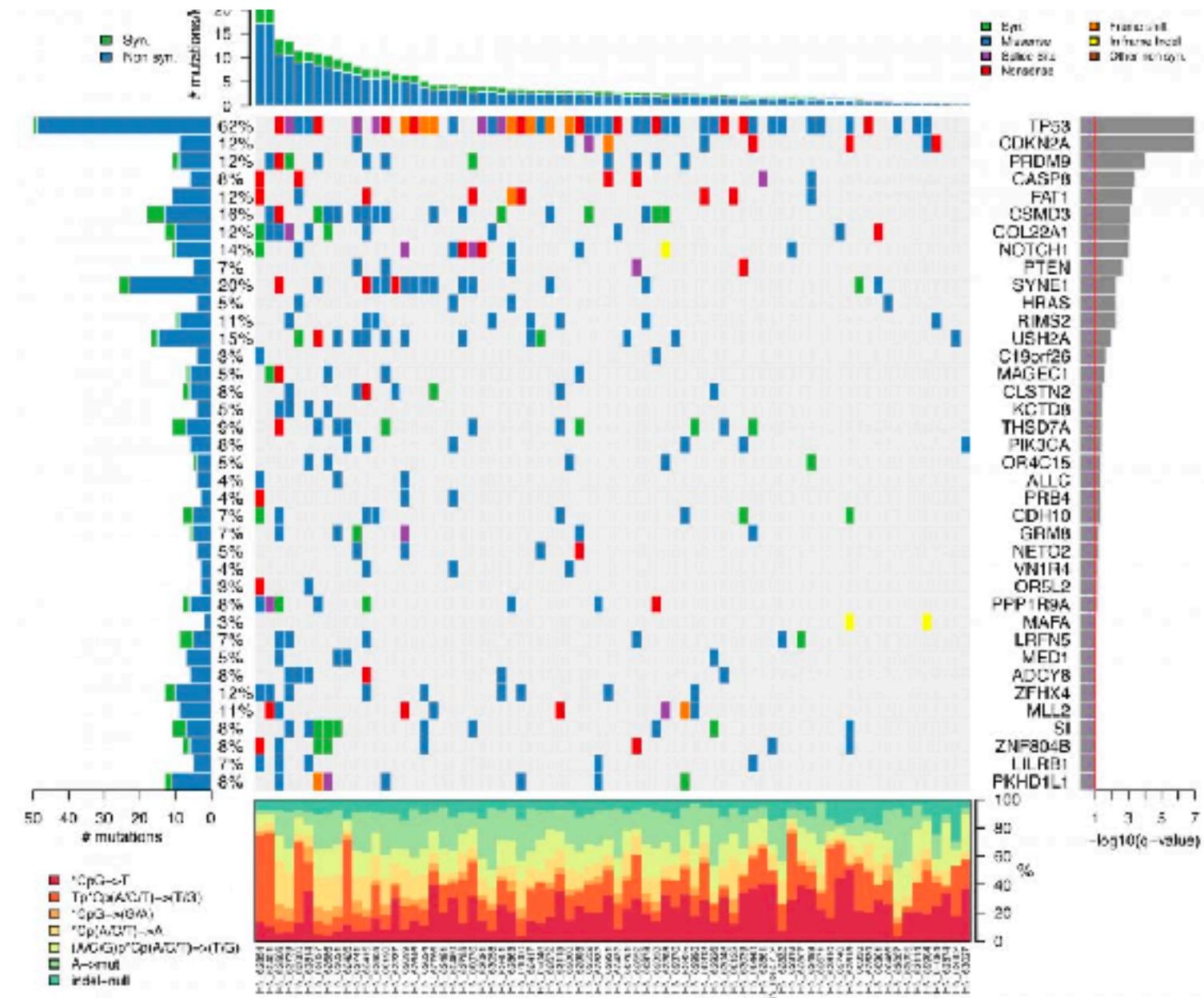


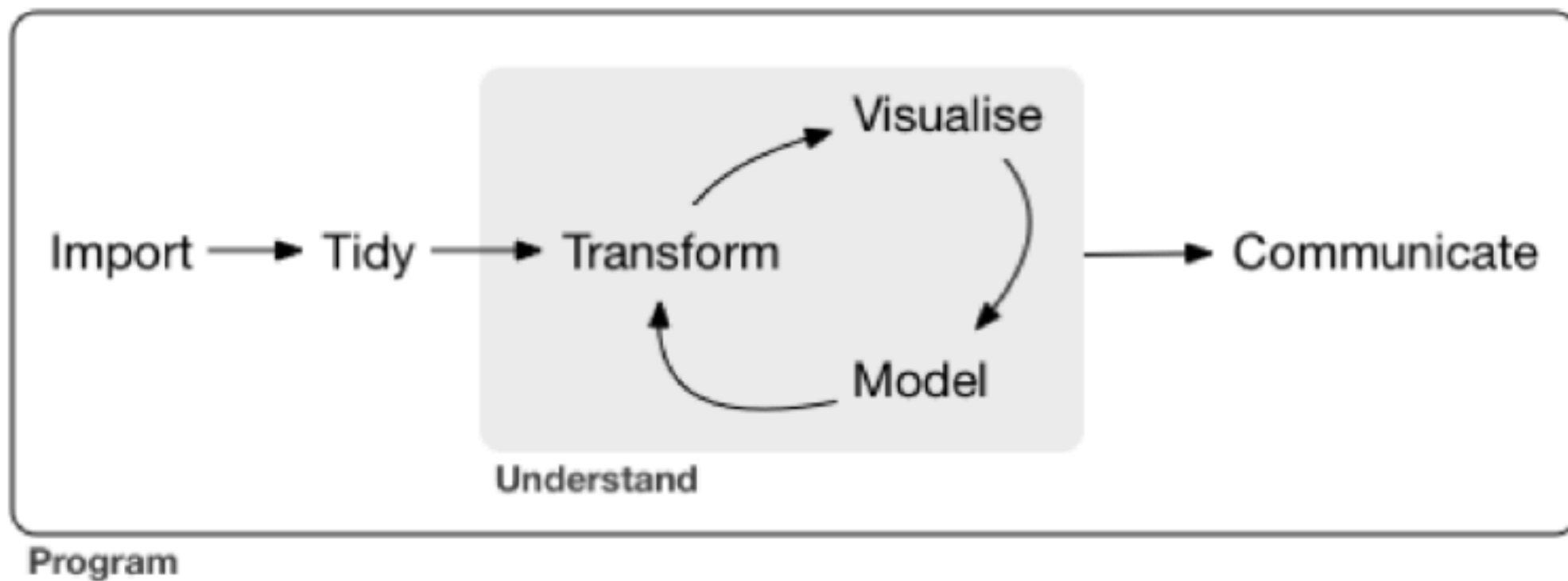
R for data transformation and visualisation

- March 2019
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```
install.packages(c("tidyverse", "ggrepel")); library(tidyverse); library(ggrepel)
```

Job of a Data Scientist

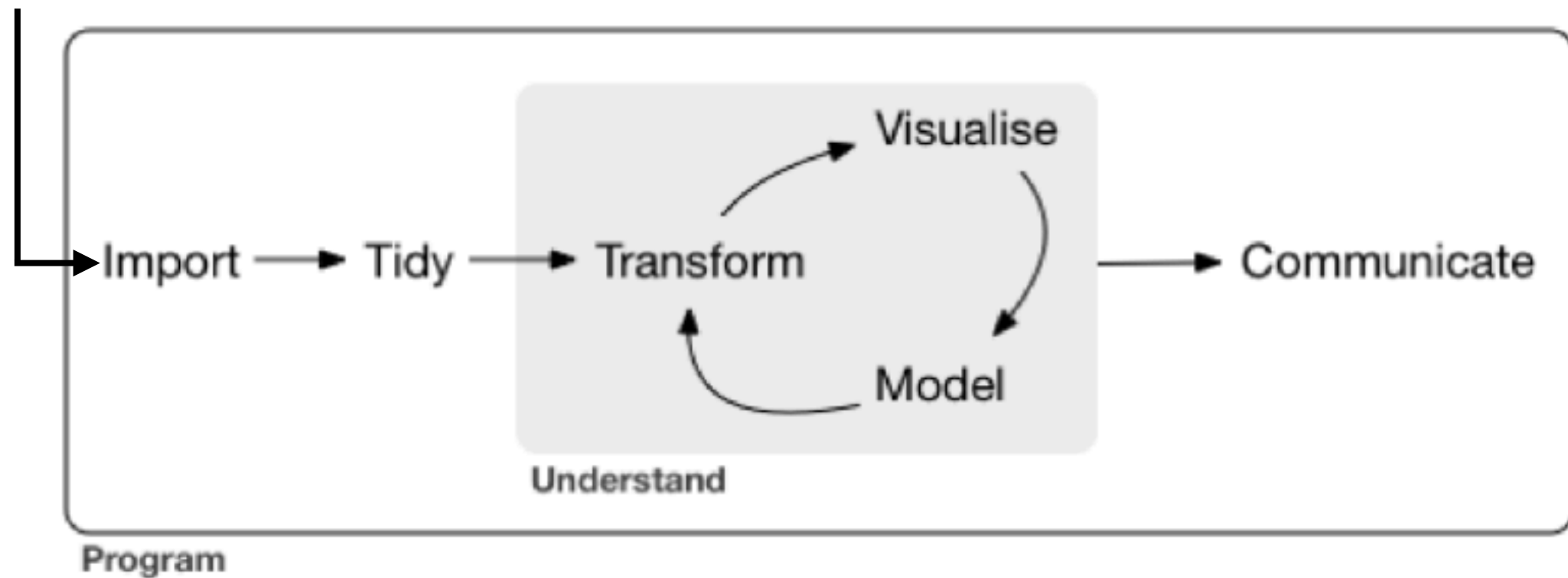


Job of a **Biological** Scientist

Perform experiment

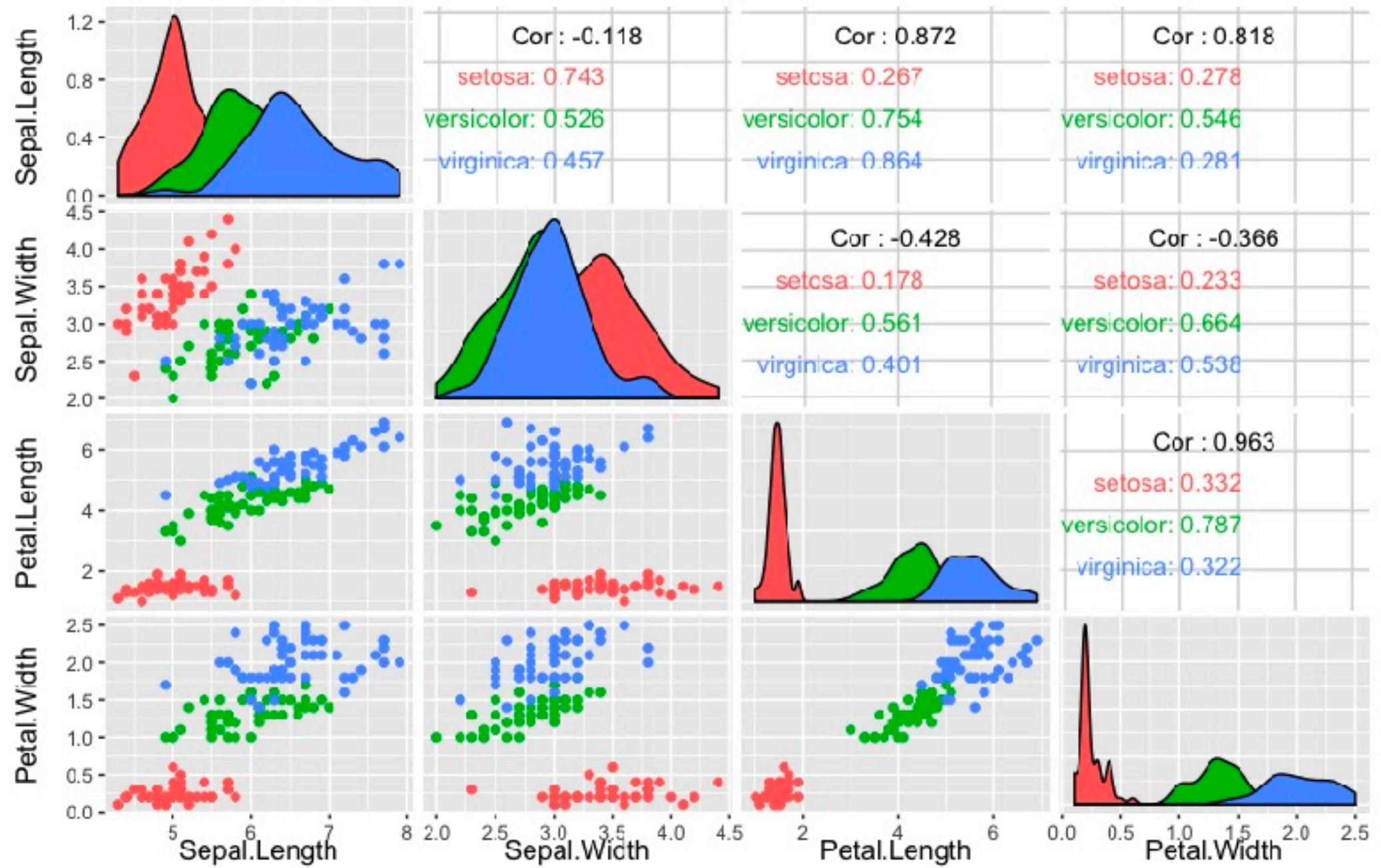


Generate (gigabytes of) data



Pair-wise Correlations

Iris phenotypes



RStudio interface

The screenshot shows the RStudio interface with several panels and annotations:

- Source Editor (Top Left):** Contains R code for reading a CSV file, calculating survival percentages by gender, and plotting the results as a bar plot. A red box highlights this area with the text **saved code**.
- Environment and History (Top Right):** Shows the Global Environment with a data object named 'titanic' (891 obs. of 13 variables). The History panel shows the execution of the code. A red box highlights this area with the text **history and environment**.
- Console (Bottom Left):** Shows the command prompt with a red box and the text **run code & see output**.
- Files Panel (Bottom Right):** Shows the file explorer with a red box and the text **plotting!**.

```
1 # read in data
2 titanic <- read.csv("https://goo.gl/4Gqsnz")
3
4 # calculate percentage survival of each gender of passenger
5 counts <- table(titanic$Sex, titanic$Pclass, titanic$Survived)
6 percentageSurvival <- counts[,2] / (counts[,1] + counts[,2])
7
8 # plot the results as a bar plot
9 par(bg = "blue")
10 barplot(percentageSurvival, beside = T, col = "hotpink4", der
11
12
13
```

Data	
titanic	891 obs. of 13 variables
Values	
counts	'table' int [1:2, 1:3, 1:2] 3 77 6 9...
percentageSurv...	table [1:2, 1:3] 96.8 36.9 92.1 15.7...

Name	Size	Modified
.RData	2.5 KB	Jul 13, 2016, 10:10 PM
.Rhistory	5.1 KB	Aug 14, 2016, 3:07 PM
node_modules		
old-container		
R		

Assigning a variable

opt -

example
`a <- 'apple'`

Running a line of code



cmd return



ctrl return

Pipe



cmd shift M



ctrl shift M

example
`diamonds %>% str()`

practice

ds **opt -** diamonds **cmd shift M** str() **cmd return**

ds <- diamonds %>% str()

Types

Converting between common data types in R. Can always go from a higher value in the table to a lower value.

<code>as.logical</code>	TRUE, FALSE, TRUE	Boolean values (TRUE or FALSE).
<code>as.numeric</code>	1, 0, 1	Integers or floating point numbers.
<code>as.character</code>	'1', '0', '1'	Character strings. Generally preferred to factors.
<code>as.factor</code>	'1', '0', '1', levels: '1', '0'	Character strings with preset levels. Needed for some statistical models.

Maths Functions

<code>log(x)</code>	Natural log.	<code>sum(x)</code>	Sum.
<code>exp(x)</code>	Exponential.	<code>mean(x)</code>	Mean.
<code>max(x)</code>	Largest element.	<code>median(x)</code>	Median.
<code>min(x)</code>	Smallest element.	<code>quantile(x)</code>	Percentage quantiles.
<code>round(x, n)</code>	Round to n decimal places.	<code>rank(x)</code>	Rank of elements.
<code>sig.fig(x, n)</code>	Round to n significant figures.	<code>var(x)</code>	The variance.
<code>cor(x, y)</code>	Correlation.	<code>sd(x)</code>	The standard deviation.

Variable Assignment

```
> a <- 'apple'
> a
[1] 'apple'
```

Vectors

Creating Vectors

<code>c(2, 4, 6)</code>	2 4 6	Join elements into a vector
<code>2:6</code>	2 3 4 5 6	An integer sequence
<code>seq(2, 3, by=0.5)</code>	2.0 2.5 3.0	A complex sequence
<code>rep(1:2, times=3)</code>	1 2 1 2 1 2	Repeat a vector
<code>rep(1:2, each=3)</code>	1 1 1 2 2 2	Repeat elements of a vector

Grammar of Graphics (ggplot2)

data

a `data_frame` containing values for plotting

aesthetic mappings

map to individual columns in data_frame

columns in the `data_frame` that map to features on the plot
e.g. x axis, y axis, point color, point size

geometric objects

set by you

What geometric shapes will be used?

`geom_point()` `geom_bar()` `geom_boxplot()` `geom_text()` `geom_histogram()` etc

coordinates

set by you

control plot layout

`coord_cartesian` (x vs y) ; `coord_map`; `coord_flipped`; `coord_polar`; `coord_equal` etc.

faceting

map to individual columns in data_frame

break data into sub-plots based on a particular grouping e.g. `facet_wrap(~ group)`

ggplots built up in layers (+)

```
mpg %>%  
ggplot(aes(x=displ, y=cty)) +  
  geom_point(col="red") +  
  geom_line(lwd=2) +  
  geom_smooth() +  
  facet_wrap(~cyl)
```

map to individual column in data_frame

No quotes

character set by you

Quotes

number

No quotes