

# R Notebook

Code ▾

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```
library(gitcreds)
gitcreds_set()
```

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```
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
```

```
-> Adding new credentials...
-> Removing credentials from cache...
-> Done.
```

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```
git config --global user.email " celiaok44@gmail.com"
git config --global user.name "herroug celia"
```

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```
library(dada2); packageVersion("dada2")
```

```
[1] '1.28.0'
```

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```
library(dplyr)
```

```
library(dplyr)

iris %>%
  filter(Sepal.Length > 6) %>% # filtered using size of Sepal Length
  filter(Species == "versicolor") # and the species
```

| Sepal.Length | Sepal.Width | Petal.Length | Petal.Width | Species    |
|--------------|-------------|--------------|-------------|------------|
| <dbl>        | <dbl>       | <dbl>        | <dbl>       | <fctr>     |
| 7.0          | 3.2         | 4.7          | 1.4         | versicolor |
| 6.4          | 3.2         | 4.5          | 1.5         | versicolor |
| 6.9          | 3.1         | 4.9          | 1.5         | versicolor |
| 6.5          | 2.8         | 4.6          | 1.5         | versicolor |
| 6.3          | 3.3         | 4.7          | 1.6         | versicolor |
| 6.6          | 2.9         | 4.6          | 1.3         | versicolor |
| 6.1          | 2.9         | 4.7          | 1.4         | versicolor |
| 6.7          | 3.1         | 4.4          | 1.4         | versicolor |
| 6.2          | 2.2         | 4.5          | 1.5         | versicolor |
| 6.1          | 2.8         | 4.0          | 1.3         | versicolor |

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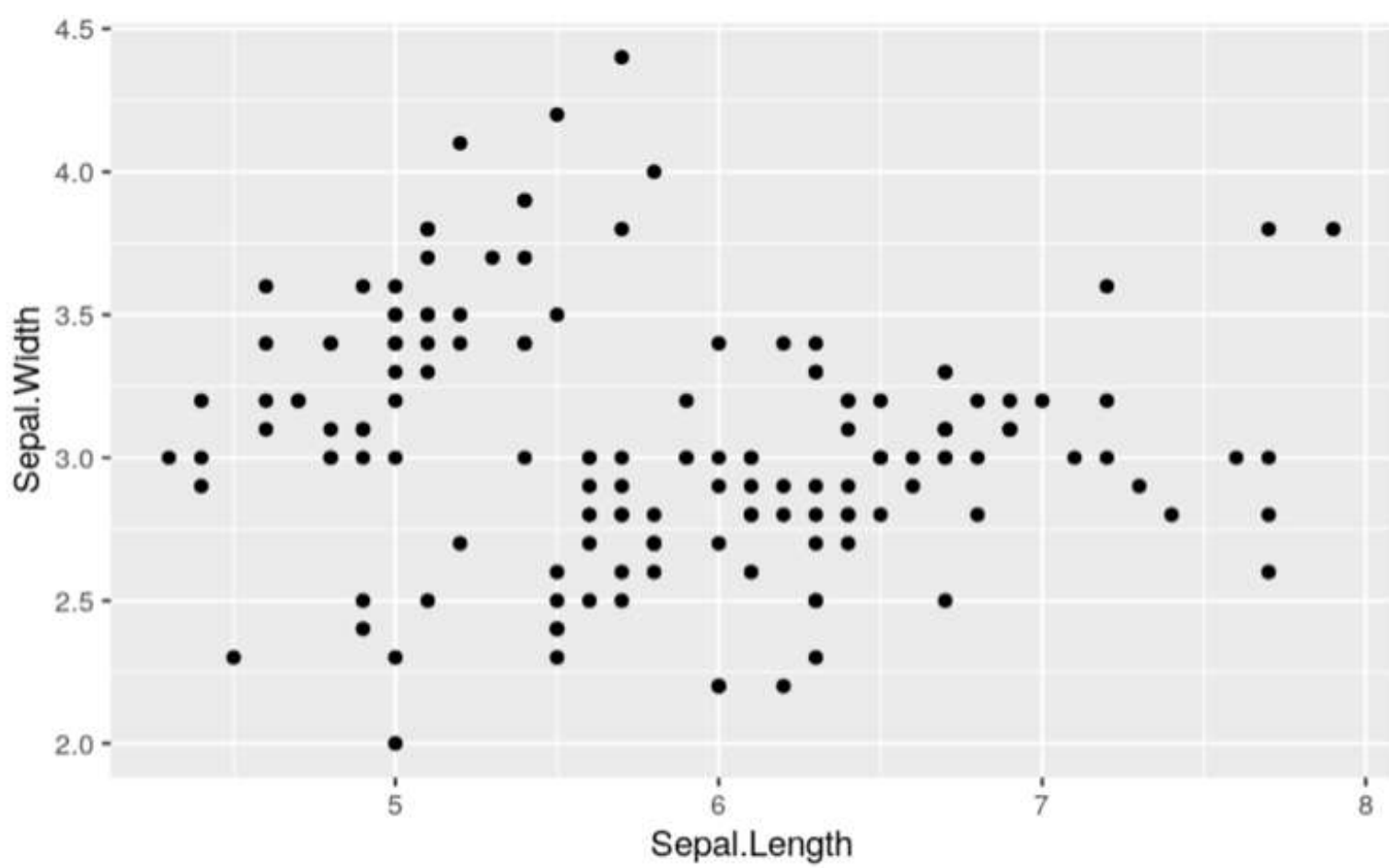
NA

run

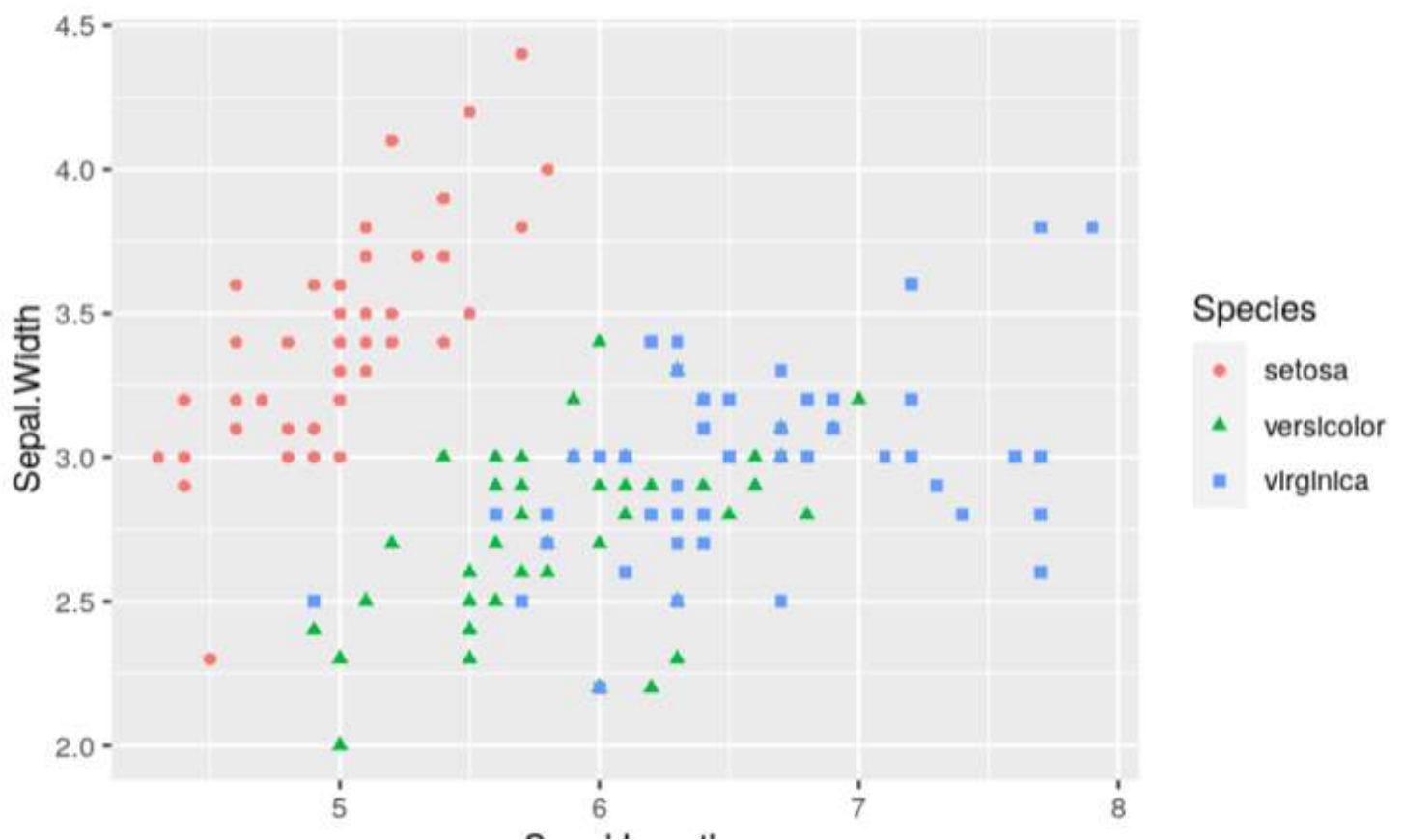
```
iris %>%
  select(Sepal.Length, Species) %>%
  mutate(Sepal.Length2 = Sepal.Length * 2) %>%
  mutate(Sepal.Length2_squared = Sepal.Length2 * Sepal.Length2)
```

| Sepal.Length | Species | Sepal.Length2 | Sepal.Length2_squared |
|--------------|---------|---------------|-----------------------|
| <dbl>        | <fctr>  | <dbl>         | <dbl>                 |
| 5.1          | setosa  | 10.2          | 104.04                |
| 4.9          | setosa  | 9.8           | 96.04                 |
| 4.7          | setosa  | 9.4           | 88.36                 |
| 4.6          | setosa  | 9.2           | 84.64                 |
| 5.0          | setosa  | 10.0          | 100.00                |
| 5.4          | setosa  | 10.8          | 116.64                |
| 4.6          | setosa  | 9.2           | 84.64                 |
| 5.0          | setosa  | 10.0          | 100.00                |
| 4.4          | setosa  | 8.8           | 77.44                 |
| 4.9          | setosa  | 9.8           | 96.04                 |

1-10 of 150 rows



```
# We can colorize and give a shape by month
# scatter plot
ggplot(data = iris, aes(Sepal.Length, Sepal.Width)) +
  geom_point(aes(color = Species, shape = Species)) # shape is the same thing with classical plot on R
```



```
#Boxplot
ggplot(data = iris, aes(Species, Sepal.Length)) + # what I want to represent
  geom_boxplot(aes(color = Species, fill = Species), alpha=0.4) + # how I represent it. Here I changed geometry using boxplot.
  labs(x = "Species", y = "Sepal.Length (mm)") +
  theme_minimal()
```

