# R Markdown Example

## Sergio Cárcamo

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# Combined R Markdown Example

This is a simple R Markdown document with code chunks and a plot included.

Use Ctrl+Alt+I to create new code chunk

### R Code Chunk with Plot

```
# Generate some random data
set.seed(42)
x <- rnorm(100)

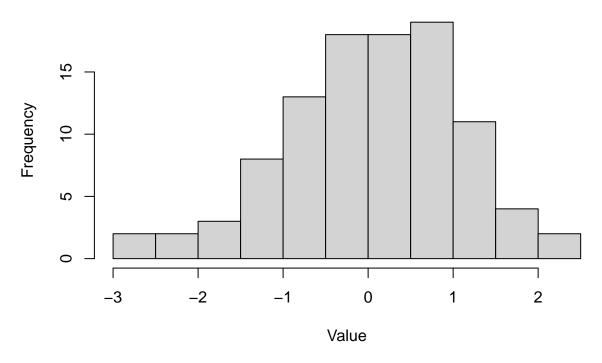
# Summary statistics
summary(x)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## -2.99309 -0.61669 0.08980 0.03251 0.66156 2.28665
```

### Plot histogram

```
hist(x, main = "Histogram of Random Data", xlab = "Value")
```

# **Histogram of Random Data**



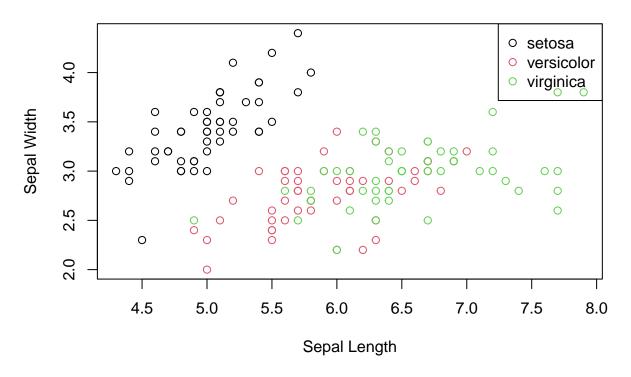
## Loading and Exploring Data

```
# Load a sample dataset
data(iris)
# View the first few rows of the dataset
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
              4.7
                          3.2
                                       1.3
                                                    0.2 setosa
## 3
## 4
              4.6
                          3.1
                                       1.5
                                                    0.2 setosa
## 5
              5.0
                          3.6
                                                    0.2 setosa
                                       1.4
## 6
              5.4
                          3.9
                                       1.7
                                                    0.4 setosa
```

#### Data visualization

```
# Scatter plot of iris dataset
plot(iris$Sepal.Length, iris$Sepal.Width,
    main = "Sepal Length vs. Sepal Width",
    xlab = "Sepal Length", ylab = "Sepal Width",
    col = iris$Species)
legend("topright", legend = levels(iris$Species), col = 1:3, pch = 1)
```

# Sepal Length vs. Sepal Width

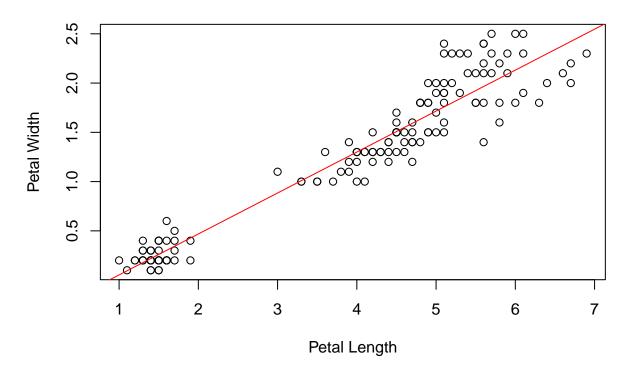


### Linear Regression

```
# Fit a linear regression model
lm_model <- lm(Petal.Width ~ Petal.Length, data = iris)</pre>
# Summary of the model
summary(lm_model)
##
## lm(formula = Petal.Width ~ Petal.Length, data = iris)
##
## Residuals:
##
       Min
                  1Q
                       Median
                                    3Q
                                            Max
## -0.56515 -0.12358 -0.01898 0.13288
                                       0.64272
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
                            0.039762 -9.131 4.7e-16 ***
## (Intercept) -0.363076
## Petal.Length 0.415755
                            0.009582
                                     43.387
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.2065 on 148 degrees of freedom
## Multiple R-squared: 0.9271, Adjusted R-squared: 0.9266
## F-statistic: 1882 on 1 and 148 DF, p-value: < 2.2e-16
# Plot the regression line
plot(iris$Petal.Length, iris$Petal.Width,
```

```
main = "Petal Width vs. Petal Length with Regression Line",
    xlab = "Petal Length", ylab = "Petal Width")
abline(lm_model, col = "red")
```

# Petal Width vs. Petal Length with Regression Line



## Interactive Plot (Using Plotly)

```
# Load the plotly library
library(plotly)
## Loading required package: ggplot2
##
## Attaching package: 'plotly'
## The following object is masked from 'package:ggplot2':
##
       last_plot
##
## The following object is masked from 'package:stats':
##
##
       filter
## The following object is masked from 'package:graphics':
##
##
       layout
# Scatter plot using plotly
plot_ly(iris, x = ~Petal.Length, y = ~Petal.Width, color = ~Species,
        type = 'scatter', mode = 'markers',
        marker = list(size = 10)) %>%
  layout(title = "Interactive Scatter Plot: Petal Width vs. Petal Length",
```

```
xaxis = list(title = "Petal Length"),
yaxis = list(title = "Petal Width"))
```