

# 1808-351-351m-451-w01b-p-RIntro4-code.R

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```
# Script Name: tea_time_class_1.R
# Purpose: Learning R
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# Latest Changelog Entries: v0.00.01 - tea_time_class1.R - Ethan Started it

# Basics of R
# Math operations
2 + 2

## [1] 4

8 / 4

## [1] 2

2 * 3

## [1] 6

3 ^ 3

## [1] 27

a = 3
b = 3
a + b

## [1] 6

# Assignment operator is usually used instead of =
# It is directional
# = works in most cases too but may cause problems

a <- 3
b <- 6

a <- b
a

## [1] 6

a <- 3
b <- 6

a -> b
a

## [1] 3

# Object classes
# numerics
a <- 5
class(a)
```

```
## [1] "numeric"

# integers
b <- as.integer(42)
class(b)

## [1] "integer"

# logicals
c <- TRUE
class(b)

## [1] "integer"

# characters
d <- "hello world"
class(c)

## [1] "logical"

# factors
e <- as.factor(c("1", "1", "a", "1", "c", "a"))
class(e)

## [1] "factor"
e

## [1] 1 1 a 1 c a
## Levels: 1 a c

# quick note it you want to convert a factor to a numeric
# You have to convert it to a character first, then a numeric

# R uses brackets to reference a data index
# data["row","column"]
# Standard organization for data set has variables as columns and observations as rows
# Keep in mind that R indexing starts from 1, not 0
# Load a test data set
data("iris")

iris[1,2]

## [1] 3.5

# Leaving a row or column input blank puts all values
# First column
iris[,1]

## [1] 5.1 4.9 4.7 4.6 5.0 5.4 4.6 5.0 4.4 4.9 5.4 4.8 4.8 4.3 5.8 5.7 5.4
## [18] 5.1 5.7 5.1 5.4 5.1 4.6 5.1 4.8 5.0 5.0 5.2 5.2 4.7 4.8 5.4 5.2 5.5
## [35] 4.9 5.0 5.5 4.9 4.4 5.1 5.0 4.5 4.4 5.0 5.1 4.8 5.1 4.6 5.3 5.0 7.0
## [52] 6.4 6.9 5.5 6.5 5.7 6.3 4.9 6.6 5.2 5.0 5.9 6.0 6.1 5.6 6.7 5.6 5.8
## [69] 6.2 5.6 5.9 6.1 6.3 6.1 6.4 6.6 6.8 6.7 6.0 5.7 5.5 5.5 5.8 6.0 5.4
## [86] 6.0 6.7 6.3 5.6 5.5 5.5 6.1 5.8 5.0 5.6 5.7 5.7 6.2 5.1 5.7 6.3 5.8
## [103] 7.1 6.3 6.5 7.6 4.9 7.3 6.7 7.2 6.5 6.4 6.8 5.7 5.8 6.4 6.5 7.7 7.7
## [120] 6.0 6.9 5.6 7.7 6.3 6.7 7.2 6.2 6.1 6.4 7.2 7.4 7.9 6.4 6.3 6.1 7.7
## [137] 6.3 6.4 6.0 6.9 6.7 6.9 5.8 6.8 6.7 6.7 6.3 6.5 6.2 5.9
```

```

# First row
iris[1,]

## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1          5.1          3.5          1.4          0.2 setosa

# Data frames have associated column names
colnames(iris)

## [1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width"
## [5] "Species"

# Columns can be called by name using $
# Rstudio features tab completion for things like column names
iris$Sepal.Length

## [1] 5.1 4.9 4.7 4.6 5.0 5.4 4.6 5.0 4.4 4.9 5.4 4.8 4.8 4.3 5.8 5.7 5.4
## [18] 5.1 5.7 5.1 5.4 5.1 4.6 5.1 4.8 5.0 5.0 5.2 5.2 4.7 4.8 5.4 5.2 5.5
## [35] 4.9 5.0 5.5 4.9 4.4 5.1 5.0 4.5 4.4 5.0 5.1 4.8 5.1 4.6 5.3 5.0 7.0
## [52] 6.4 6.9 5.5 6.5 5.7 6.3 4.9 6.6 5.2 5.0 5.9 6.0 6.1 5.6 6.7 5.6 5.8
## [69] 6.2 5.6 5.9 6.1 6.3 6.1 6.4 6.6 6.8 6.7 6.0 5.7 5.5 5.5 5.8 6.0 5.4
## [86] 6.0 6.7 6.3 5.6 5.5 5.5 6.1 5.8 5.0 5.6 5.7 5.7 6.2 5.1 5.7 6.3 5.8
## [103] 7.1 6.3 6.5 7.6 4.9 7.3 6.7 7.2 6.5 6.4 6.8 5.7 5.8 6.4 6.5 7.7 7.7
## [120] 6.0 6.9 5.6 7.7 6.3 6.7 7.2 6.2 6.1 6.4 7.2 7.4 7.9 6.4 6.3 6.1 7.7
## [137] 6.3 6.4 6.0 6.9 6.7 6.9 5.8 6.8 6.7 6.7 6.3 6.5 6.2 5.9

# Functions are processes that take an input and give an output
# Rstudio has tab completion for function inputs
max(iris$Petal.Length)

## [1] 6.9

mean(iris$Sepal.Width)

## [1] 3.057333

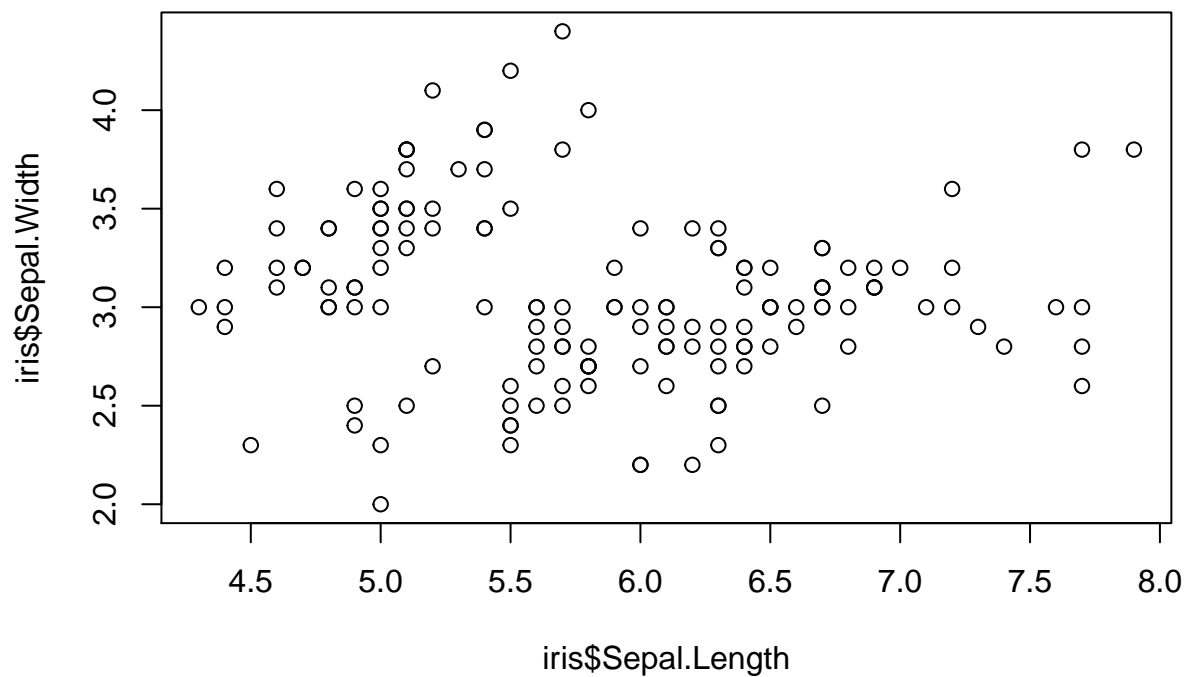
sd(iris$Petal.Width)

## [1] 0.7622377

# Functions can take multiple inputs, they can be named in the call or placed in order
plot(x = iris$Sepal.Length, y = iris$Sepal.Width)

# x and y can be specified with x = ... in any order or the inputs can be given in order
# This plot is the same as the previous
plot(iris$Sepal.Length, iris$Sepal.Width)

```



```
# Matrix operations
```

```
mat <- matrix(data = 1:9, nrow = 3, ncol = 3)
mat
```

```
##      [,1] [,2] [,3]
## [1,]    1    4    7
## [2,]    2    5    8
## [3,]    3    6    9
```

```
# Element multiplication
```

```
mat*mat
```

```
##      [,1] [,2] [,3]
## [1,]    1   16   49
## [2,]    4   25   64
## [3,]    9   36   81
```

```
# Matrix multiplication
```

```
mat %*% mat
```

```
##      [,1] [,2] [,3]
## [1,]   30   66  102
## [2,]   36   81  126
## [3,]   42   96  150
```

```
# t() function is for transposing
```

```
t(mat)
```

```
##      [,1] [,2] [,3]
```

```
## [1,] 1 2 3
## [2,] 4 5 6
## [3,] 7 8 9
```

```
mat %*% t(mat)
```

```
##      [,1] [,2] [,3]
## [1,] 66 78 90
## [2,] 78 93 108
## [3,] 90 108 126
```

```
# Inverse matrix
```

```
mat[2,3] <- 18
solve(mat)
```

```
##      [,1] [,2] [,3]
## [1,] -1.05 0.1 0.61666667
## [2,] 0.60 -0.2 -0.06666667
## [3,] -0.05 0.1 -0.05000000
```

```
solve(mat) %*% mat
```

```
##      [,1] [,2] [,3]
## [1,] 1 0.000000e+00 8.881784e-16
## [2,] 0 1.000000e+00 2.220446e-16
## [3,] 0 -5.551115e-17 1.000000e+00
```

```
# Structures in R
```

```
# for loops
```

```
for (i in 1:5) {
  print(i)
}
```

```
## [1] 1
## [1] 2
## [1] 3
## [1] 4
## [1] 5
```

```
# While loops
```

```
i <- 10
while (i > 5) {
  i <- i - 1
  print(i)
}
```

```
## [1] 9
## [1] 8
## [1] 7
## [1] 6
## [1] 5
```

```
# if statments
```

```
dave <- TRUE
```

```
# if (dave) {} also works
```

```
if (dave == TRUE) {
  print("good morning dave")
}
```

```
## [1] "good morning dave"
```

```
# User defined functions
```

```
math <- function(a,b) {
```

```
  c <- a + b*2
```

```
  # return defines what the output of the function is
```

```
  return(c)
```

```
}
```

```
math(2,6)
```

```
## [1] 14
```