R for Geoscience

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2023-02-26

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Part I

Chapter 1

Base R you have to know

1.1

1.2 R?

1992   Ross Ihaka Robert Gentleman

Figure 1.1: *Ross Ihaka and Robert Gentleman, the creators of R.*

R   R   R   R   The

R Base Package

     B   R Python stata   R

1.3   Vector( )

Vector R

     1 5 5

```
x <- c(1,2,3,4,5)
x
#> [1] 1 2 3 4 5
```

   c()   1 2 3 4 5   <-   x   1 2 3 4 5   x   R
     R   c()

  R   google   :,   R

```
x <- c(1:5)
x
#> [1] 1 2 3 4 5
```

   vector   typeof()

```
typeof(x)
#> [1] "integer"
```

            length ,length()

```
length(x)
#> [1] 5
```

        R seq()

```
seq(1, 9, 0.5)
#>  [1] 1.0 1.5 2.0 2.5 3.0 3.5 4.0 4.5 5.0 5.5 6.0 6.5 7.0 7.5
#> [15] 8.0 8.5 9.0
```

            1 9 0.5   3   ??   Console
??seq
   Help   ::   :: package::function   cli   a
generation
            Description   Usage,seq(...)   seq   (),   A
        vector

```
# Vector of logical values
log_values <- c(TRUE, FALSE, TRUE, FALSE)

log_values
#> [1]  TRUE FALSE  TRUE FALSE
```

   #   R

```
fruits <- c("banana", "apple", "orange", "mango", "lemon", "50")
fruits
#> [1] "banana" "apple"  "orange" "mango"  "lemon"  "50"
```

                   []   brackets,   fruits "banana" "mango"

```
fruits[c(1,4)]
#> [1] "banana" "mango"
```

```
fruits[1:4]
#> [1] "banana" "apple"  "orange" "mango"
```

     "banana

```
fruits[-1]
#> [1] "apple"  "orange" "mango"  "lemon"  "50"
```

                sort,

```
fruits <- c("banana", "apple", "orange", "mango", "lemon")
numbers <- c(13, 3, 5, 7, 20, 2)

sort(fruits)  # Sort a string
#> [1] "apple"  "banana" "lemon"  "mango"  "orange"
sort(numbers) # Sort numbers
#> [1]  2  3  5  7 13 20
```

1.4   Lists( )

R   `list()`

```r
thislist <- list(
  a = c("apple", "banana", "cherry"),
  b = c(1,2,5,6,7,9),
  c = c(TRUE, FALSE, TRUE)
)
# Print the list
thislist
#> $a
#> [1] "apple"  "banana" "cherry"
#>
#> $b
#> [1] 1 2 5 6 7 9
#>
#> $c
#> [1]  TRUE FALSE  TRUE
```

```r
typeof(thislist)
#> [1] "list"
```

```r
length(thislist)
#> [1] 3
```

1.5   Matrices( )

(column) (row)   `matrix()`

,

```r
# Create a matrix
thismatrix <- matrix(c(1,2,3,4,5,6), nrow = 3, ncol = 2)

# Print the matrix
thismatrix
#>      [,1] [,2]
#> [1,]    1    4
#> [2,]    2    5
#> [3,]    3    6
```

NOTE: c()

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)

thismatrix
#>       [,1]      [,2]
#> [1,] "apple"   "cherry"
#> [2,] "banana" "orange"
```

Access Matrix Items You can access the items by using [ ] brackets. The first number "1" in the bracket specifies the row-position, while the second number "2" specifies the column-position:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)

thismatrix[1, 2]
#> [1] "cherry"
```

The whole row can be accessed if you specify a comma after the number in the bracket:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)

thismatrix[2,]
#> [1] "banana" "orange"
```

The whole column can be accessed if you specify a comma before the number in the bracket:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)

thismatrix[,2]
#> [1] "cherry" "orange"
```

Access More Than One Row More than one row can be accessed if you use the c() function:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange","grape", "pineapple", "pear", "melon

thismatrix[c(1,2),]
#>       [,1]      [,2]      [,3]
#> [1,] "apple"   "orange" "pear"
#> [2,] "banana" "grape"   "melon"
```

Access More Than One Column More than one column can be accessed if you use the c() function:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange","grape", "pineapple", "pea

thismatrix[, c(1,2)]
#>      [,1]      [,2]
#> [1,] "apple"   "orange"
#> [2,] "banana"  "grape"
#> [3,] "cherry"  "pineapple"
```

Add Rows and Columns Use the cbind() function to add additional columns in a Matrix:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange","grape", "pineapple", "pea

newmatrix <- cbind(thismatrix, c("strawberry", "blueberry", "raspberry"))

# Print the new matrix
newmatrix
#>      [,1]      [,2]         [,3]     [,4]
#> [1,] "apple"   "orange"     "pear"   "strawberry"
#> [2,] "banana"  "grape"      "melon"  "blueberry"
#> [3,] "cherry"  "pineapple"  "fig"    "raspberry"
```

Use the rbind() function to add additional rows in a Matrix:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange","grape", "pineapple", "pea

newmatrix <- rbind(thismatrix, c("strawberry", "blueberry", "raspberry"))

# Print the new matrix
newmatrix
#>      [,1]          [,2]          [,3]
#> [1,] "apple"       "orange"      "pear"
#> [2,] "banana"      "grape"       "melon"
#> [3,] "cherry"      "pineapple"   "fig"
#> [4,] "strawberry"  "blueberry"   "raspberry"
```

Remove Rows and Columns Use the c() function to remove rows and columns in a Matrix:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange", "mango", "pineapple"), n
```

```
#Remove the first row and the first column
thismatrix <- thismatrix[-c(1), -c(1)]

thismatrix
#> [1] "mango"      "pineapple"
```

Check if an Item Exists To find out if a specified item is present in a matrix, use the %in% operator:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)

"apple" %in% thismatrix
#> [1] TRUE
```

Number of Rows and Columns Use the dim() function to find the number of rows and columns in a Matrix:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)

dim(thismatrix)
#> [1] 2 2
```

Matrix Length Use the length() function to find the dimension of a Matrix:

```
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)

length(thismatrix)
#> [1] 4
```

Combine two Matrices Again, you can use the rbind() or cbind() function to combine two or more matrices together:

```
# Combine matrices
Matrix1 <- matrix(c("apple", "banana", "cherry", "grape"), nrow = 2, ncol = 2)
Matrix2 <- matrix(c("orange", "mango", "pineapple", "watermelon"), nrow = 2, ncol = 2)

# Adding it as a rows
Matrix_Combined <- rbind(Matrix1, Matrix2)
Matrix_Combined
#>      [,1]     [,2]
#> [1,] "apple"  "cherry"
#> [2,] "banana" "grape"
#> [3,] "orange" "pineapple"
```

```
#> [4,] "mango"   "watermelon"

# Adding it as a columns
Matrix_Combined <- cbind(Matrix1, Matrix2)
Matrix_Combined
#>      [,1]     [,2]     [,3]     [,4]
#> [1,] "apple"  "cherry" "orange" "pineapple"
#> [2,] "banana" "grape"  "mango"  "watermelon"
```

1.6 Data Frame( )

data.frame()

```
# Create a data frame
Data_Frame <- data.frame (
  Training = c("Strength", "Stamina", "Other"),
  Pulse = c(100, 150, 120),
  Duration = c(60, 30, 45)
)

# Print the data frame
Data_Frame
#>   Training Pulse Duration
#> 1 Strength   100       60
#> 2  Stamina   150       30
#> 3    Other   120       45
```

Use the `summary()` function to summarize the data from a Data Frame:

```
summary(Data_Frame)
#>    Training             Pulse          Duration
#>  Length:3          Min.   :100.0   Min.   :30.0
#>  Class :character  1st Qu.:110.0   1st Qu.:37.5
#>  Mode  :character  Median :120.0   Median :45.0
#>                    Mean   :123.3   Mean   :45.0
#>                    3rd Qu.:135.0   3rd Qu.:52.5
#>                    Max.   :150.0   Max.   :60.0
```

[]   [[ ]] $

```
Data_Frame[1]
#>   Training
#> 1 Strength
#> 2  Stamina
#> 3    Other

Data_Frame[["Training"]]
#> [1] "Strength" "Stamina"  "Other"

Data_Frame$Training
#> [1] "Strength" "Stamina"  "Other"
```

rbind()

```
# Add a new row
New_row_DF <- rbind(Data_Frame, c("Strength", 110, 110))

# Print the new row
New_row_DF
#>   Training Pulse Duration
#> 1 Strength   100       60
#> 2  Stamina   150       30
#> 3    Other   120       45
#> 4 Strength   110      110
```

cbind()

```
# Add a new column
New_col_DF <- cbind(New_row_DF, Steps = c(1000, 6000, 2000,5000))

# Print the new column
New_col_DF
#>   Training Pulse Duration Steps
#> 1 Strength   100       60  1000
#> 2  Stamina   150       30  6000
#> 3    Other   120       45  2000
#> 4 Strength   110      110  5000
```

rbind()   R

```
Data_Frame1 <- data.frame (
  Training = c("Strength", "Stamina", "Other"),
  Pulse = c(100, 150, 120),
  Duration = c(60, 30, 45)
```

```
)

Data_Frame2 <- data.frame (
  Training = c("Stamina", "Stamina", "Strength"),
  Pulse = c(140, 150, 160),
  Duration = c(30, 30, 20)
)

New_Data_Frame <- rbind(Data_Frame1, Data_Frame2)
New_Data_Frame
#>    Training Pulse Duration
#> 1 Strength   100       60
#> 2  Stamina   150       30
#> 3    Other   120       45
#> 4  Stamina   140       30
#> 5  Stamina   150       30
#> 6 Strength   160       20
```

cbind()   R

```
Data_Frame3 <- data.frame (
  Training = c("Strength", "Stamina", "Other"),
  Pulse = c(100, 150, 120),
  Duration = c(60, 30, 45)
)

Data_Frame4 <- data.frame (
  Steps = c(3000, 6000, 2000),
  Calories = c(300, 400, 300)
)

New_Data_Frame1 <- cbind(Data_Frame3, Data_Frame4)
New_Data_Frame1
#>    Training Pulse Duration Steps Calories
#> 1 Strength   100       60  3000      300
#> 2  Stamina   150       30  6000      400
#> 3    Other   120       45  2000      300
```