

Practical 1

Introduction to R

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EXERCICE 1

```
# Generate the numbers 1, 2, . . . , 12,  
# and store the result in the vector x
```

```
x <- 1:12  
print(x)
```

```
## [1] 1 2 3 4 5 6 7 8 9 10 11 12
```

EXERCICE 2

```
# Generate four repetitions of the sequence of numbers (6, 2, 4).
```

```
y <- rep(c(6, 2, 4), 4)  
print(y)
```

```
## [1] 6 2 4 6 2 4 6 2 4 6 2 4
```

EXERCICE 3

```
# Generate the sequence consisting of six 9s, then five 2s, and finally four  
# 5s. Store the numbers in a 5 by 3 matrix (populating it columnwise).
```

```
sequence <- c(rep(9, 6), rep(2, 5), rep(5, 4))  
matrix_sequence <- matrix(sequence, nrow = 5, ncol = 3, byrow = FALSE)  
print(matrix_sequence)
```

```
##      [,1] [,2] [,3]  
## [1,]    9    9    2  
## [2,]    9    2    5  
## [3,]    9    2    5  
## [4,]    9    2    5  
## [5,]    9    2    5
```

EXERCICE 4

```
# Generate a vector consisting of 20 numbers generated randomly from  
# a normal distribution. Use the value 100 as seed (in order to be able
```

```
# to replicate the experiments).
```

```
set.seed(100)
random_vector <- rnorm(20)
print(random_vector)
```

```
## [1] -0.50219235  0.13153117 -0.07891709  0.88678481  0.11697127  0.31863009
## [7] -0.58179068  0.71453271 -0.82525943 -0.35986213  0.08988614  0.09627446
## [13] -0.20163395  0.73984050  0.12337950 -0.02931671 -0.38885425  0.51085626
## [19] -0.91381419  2.31029682
```

```
# Then, calculate the following statistics about the generated vector:
# mean, median, variance and the standard deviation.
```

```
mean_value <- mean(random_vector)
median_value <- median(random_vector)
variance_value <- var(random_vector)
standard_deviation_value <- sd(random_vector)
```

```
print(paste("Mean: ", mean_value))
```

```
## [1] "Mean:  0.107867147646684"
```

```
print(paste("Median: ", median_value))
```

```
## [1] "Median:  0.0930803020313303"
```

```
print(paste("Variance: ", variance_value))
```

```
## [1] "Variance:  0.516335002736962"
```

```
print(paste("Standard deviation: ", standard_deviation_value))
```

```
## [1] "Standard deviation:  0.718564543194946"
```

```
# Repeat the generation of the vector and the statistics with and without
# changing the seed and observe what happens.
```

```
set.seed(100)
random_vector <- rnorm(20)
print(random_vector)
```

```
## [1] -0.50219235  0.13153117 -0.07891709  0.88678481  0.11697127  0.31863009
## [7] -0.58179068  0.71453271 -0.82525943 -0.35986213  0.08988614  0.09627446
## [13] -0.20163395  0.73984050  0.12337950 -0.02931671 -0.38885425  0.51085626
## [19] -0.91381419  2.31029682
```

```
mean_value <- mean(random_vector)
median_value <- median(random_vector)
variance_value <- var(random_vector)
standard_deviation_value <- sd(random_vector)
```

```
print(paste("Mean: ", mean_value))
```

```
## [1] "Mean:  0.107867147646684"
```

```
print(paste("Median: ", median_value))
```

```
## [1] "Median:  0.0930803020313303"
```

```

print(paste("Variance: ", variance_value))

## [1] "Variance: 0.516335002736962"
print(paste("Standard deviation: ", standard_deviation_value))

## [1] "Standard deviation: 0.718564543194946"
# Random vector with seed 200
set.seed(200)
random_vector <- rnorm(20)
print(random_vector)

## [1] 0.08475635 0.22646034 0.43255650 0.55806524 0.05975527 -0.11464087
## [7] -1.02057835 -0.29705130 0.16815003 1.41987233 -0.09952507 -0.81829697
## [13] -0.46930224 0.57504497 -1.87174513 -0.63183110 -0.04243820 1.44210693
## [19] -0.92089342 -0.01560860

mean_value <- mean(random_vector)
median_value <- median(random_vector)
variance_value <- var(random_vector)
standard_deviation_value <- sd(random_vector)

print(paste("Mean: ", mean_value))

## [1] "Mean: -0.0667571647297675"
print(paste("Median: ", median_value))

## [1] "Median: -0.0290233979238126"
print(paste("Variance: ", variance_value))

## [1] "Variance: 0.616924355239516"
print(paste("Standard deviation: ", standard_deviation_value))

## [1] "Standard deviation: 0.785445322883468"
# Random vector with seed 300
set.seed(300)
random_vector <- rnorm(20)
print(random_vector)

## [1] 1.37379088 0.86210687 0.47348910 0.70126281 -0.08505527 1.56870212
## [7] 0.81739197 0.39476860 1.21269855 0.35508066 2.21627421 -0.09054039
## [13] -1.31652811 0.06653479 0.51217262 0.05002974 1.46739718 -1.28952052
## [19] -0.25769151 0.29305131

mean_value <- mean(random_vector)
median_value <- median(random_vector)
variance_value <- var(random_vector)
standard_deviation_value <- sd(random_vector)

print(paste("Mean: ", mean_value))

## [1] "Mean: 0.466270781584136"
print(paste("Median: ", median_value))

```

```
## [1] "Median: 0.434128853552544"
print(paste("Variance: ", variance_value))

## [1] "Variance: 0.77779170713689"
print(paste("Standard deviation: ", standard_deviation_value))

## [1] "Standard deviation: 0.881925000857153"
```

EXERCICE 5

```
# Read the data into an R object named students (data is in a
# space-delimited text file and there is no header row).

students <- read.table("./dataset/data1.txt", header = FALSE)

# Add the following titles for columns:
# height, shoesize, gender, population

colnames(students) <- c("height", "shoesize", "gender", "population")

# Check that R reads the file correctly.

print(students)
```

```
##   height shoesize gender population
## 1    181      44   male    kuopio
## 2    160      38 female    kuopio
## 3    174      42 female    kuopio
## 4    170      43   male    kuopio
## 5    172      43   male    kuopio
## 6    165      39 female    kuopio
## 7    161      38 female    kuopio
## 8    167      38 female tampere
## 9    164      39 female tampere
## 10   166      38 female tampere
## 11   162      37 female tampere
## 12   158      36 female tampere
## 13   175      42   male tampere
## 14   181      44   male tampere
## 15   180      43   male tampere
## 16   177      43   male tampere
## 17   173      41   male tampere
```

```
# Print the header names only.
```

```
print(colnames(students))
```

```
## [1] "height"      "shoesize"    "gender"      "population"
```

```
# Print the column height.
```

```
print(students$height)
```

```
## [1] 181 160 174 170 172 165 161 167 164 166 162 158 175 181 180 177 173
```

```
# What is the gender distribution (how many observations are in  
# each group) and the distribution of sampling sites (column population)?
```

```
gender_distribution <- table(students$gender)  
population_distribution <- table(students$population)  
print(gender_distribution)
```

```
##  
## female    male  
##         9     8
```

```
print(population_distribution)
```

```
##  
## kuopio tampere  
##         7     10
```

```
# Show the distributions in the above item at the same time by  
# using a contingency table.
```

```
contingency_table <- table(students$gender, students$population)  
print(contingency_table)
```

```
##  
##           kuopio tampere  
## female      4          5  
## male        3          5
```

```
# Make two subsets of your dataset by splitting it according to gender.  
# Use data frame operations first and then do the same using  
# the function subset.
```

```
male_students <- students[students$gender == "Male", ]  
female_students <- students[students$gender == "Female", ]
```

```
# Subsets by gender using subset
```

```
male_students_subset <- subset(students, gender == "Male")  
female_students_subset <- subset(students, gender == "Female")
```

```
# Make two subsets containing individuals below and above the  
# median height. Use data frame operations first and then do the  
# same using the function subset.
```

```
median_height <- median(students$height)  
below_median_height <- students[students$height < median_height, ]  
above_median_height <- students[students$height > median_height, ]
```

```
# Subsets by height using subset
```

```
below_median_subset <- subset(students, height < median_height)  
above_median_subset <- subset(students, height > median_height)
```

```
# Change height from centimetres to metres for all rows in the  
# data frame. Do this using in three different ways: with basic  
# primitives, a loop using for and the function apply.
```

```
# Using basic primitives
```

```

students$height <- students$height / 100

# Using a loop
for (i in seq_len(nrow(students))) {
  students$height[i] <- students$height[i] / 100
}

# Using apply
students$height <- sapply(students$height, function(x) x / 100)

# Plot height against shoesize, using blue circles
# for males and magenta crosses for females.
# Add a legend.

# Correct the conversion of height to meters
students$height <- students$height * 10000

plot(students$height, students$shoesize,
     col = ifelse(students$gender == "Male", "blue", "magenta"),
     pch = ifelse(students$gender == "Male", 16, 4),
     xlab = "Height (m)", ylab = "Shoe Size")
legend("topright", legend = c("Male", "Female"),
     col = c("blue", "magenta"),
     pch = c(16, 4))

```

