

7.1. Create a 2x3x2 array with values from 1 to 12.

- Name the rows as "Row1", "Row2", and columns as "Col1", "Col2", "Col3".
- Access the elements in the first row and second column of the second layer.
- Replace all elements greater than 6 with NA.
- Calculate the sum of all elements in the array.

#Code:

```
# a: Create the array with names
my_array <- array(1:12, dim = c(2, 3, 2),
  dimnames = list(c("Row1", "Row2"),
    c("Col1", "Col2", "Col3")))
print(my_array)

# b: Access element in first row, second column, second layer
element <- my_array["Row1", "Col2", 2]
print(element)

# c: Replace values > 6 with NA
my_array[my_array > 6] <- NA
print(my_array)

# d: Calculate sum ignoring NA
total_sum <- sum(my_array, na.rm = TRUE)
print(total_sum)
```

#output:

```
,, 1
  Col1 Col2 Col3
Row1  1  3  5
Row2  2  4  6
,, 2
  Col1 Col2 Col3
Row1  7  9 11
Row2  8 10 12
```

```
[1] 9
```

```
, , 1
```

```
Col1 Col2 Col3
```

```
Row1  1   3   5
```

```
Row2  2   4   6
```

```
, , 2
```

```
Col1 Col2 Col3
```

```
Row1 NA  NA  NA
```

```
Row2 NA  NA  NA
```

```
[1] 21
```

7.2. Create a factor using the vector `c("Red", "Blue", "Green", "Red", "Blue")`.

- a. Display the levels of the factor.**
- b. Generate a factor with 3 levels ("High", "Medium", "Low") repeated 4 times.**
- c. Modify the factor to include a new level "Very High".**
- d. Drop the level "Medium" from the factor and display the updated factor.**

#Code:

```
# Step 1: Create a factor
```

```
colors <- factor(c("Red", "Blue", "Green", "Red", "Blue"))
```

```
# Step a: Display levels
```

```
print(levels(colors))
```

```
# Step b: Create repeated factor
```

```
levels_vector <- factor(rep(c("High", "Medium", "Low"), times = 4))
```

```
print(levels_vector)
```

```
# Step c: Add new level
```

```
levels(levels_vector) <- c(levels(levels_vector), "Very High")
```

```
# Step d: Drop "Medium" and display updated factor
levels_vector <- levels_vector[levels_vector != "Medium"]
levels_vector <- droplevels(levels_vector)
print(levels_vector)
```

#output:

```
[1] "Blue" "Green" "Red"
```

```
[1] High Medium Low High Medium Low High Medium Low High Medium Low
Levels: High Low Medium
```

```
[1] High Low High Low High Low High Low
Levels: High Low
```

8. Write a script to process the text "R Programming is Fun and Challenging".

- a. Extract every second word from the sentence.
- b. Count the number of occurrences of vowels (a, e, i, o, u) in the string.
- c. Replace the word "Challenging" with "Exciting".

Code:

```
# Load required library
library(stringr)
```

```
# Original text
text <- "R Programming is Fun and Challenging"
```

```
# a. Extract every second word
words <- strsplit(text, " ")[[1]]
second_words <- words[seq(2, length(words), by = 2)]
print(second_words)
```

```
# b. Count number of vowels
```

```
lower_text <- tolower(text)
vowel_count <- sum(str_count(lower_text, "[aeiou]"))
print(vowel_count)
```

```
# c. Replace "Challenging" with "Exciting"
new_text <- sub("Challenging", "Exciting", text)
print(new_text)
```

#Output:

```
[1] "Programming" "Fun" "Challenging"
```

```
[1] 9
```

```
[1] "R Programming is Fun and Exciting"
```

9.1. Create a nested list containing:

- a. A data frame with student names, marks, and grades.
- b. A vector with the total marks for each student.
- c. A list of factors indicating the performance category ("Excellent", "Good", "Average").

Code:

```
# a. Data frame with student names, marks, and grades
```

```
students_df <- data.frame(
  Name = c("Alice", "Bob", "Charlie"),
  Marks = c(88, 75, 62),
  Grade = c("A", "B", "C")
)
```

```
# b. Vector with total marks (assuming one subject each for now)
```

```
total_marks <- c(88, 75, 62)
```

```
# c. List of factors for performance category
performance <- list(
  factor(c("Excellent", "Good", "Average"))
)
```

```
# Combine all into a nested list
nested_list <- list(
  StudentData = students_df,
  TotalMarks = total_marks,
  PerformanceCategory = performance
)
```

```
# View the nested list
print(nested_list)
```

#output:

```
$StudentData
```

	Name	Marks	Grade
1	Alice	88	A
2	Bob	75	B
3	Charlie	62	C

```
$TotalMarks
```

```
[1] 88 75 62
```

```
$PerformanceCategory
```

```
$PerformanceCategory[[1]]
```

```
[1] Excellent Good Average
```

```
Levels: Average Excellent Good
```

9.2. Write a script to:

- Access and modify the data frame inside the nested list.
- Add a new entry for a student.
- Extract students with "Excellent" performance.

Code:

Step a: Modify Bob's marks

```
nested_list$StudentData$Marks[nested_list$StudentData$Name == "Bob"] <- 80
```

Step b: Add new student David

```
new_student <- data.frame(Name = "David", Marks = 91, Grade = "A")
```

```
nested_list$StudentData <- rbind(nested_list$StudentData, new_student)
```

```
nested_list$TotalMarks <- c(nested_list$TotalMarks, 91)
```

```
nested_list$PerformanceCategory[[1]] <- factor(  
  c(as.character(nested_list$PerformanceCategory[[1]]), "Excellent"),  
  levels = c("Average", "Good", "Excellent")  
)
```

Step c: Extract students with "Excellent" performance (Grade A)

```
excellent_students <- nested_list$StudentData[nested_list$StudentData$Grade == "A", ]  
print(excellent_students)
```

Output:

	Name	Marks	Grade
1	Alice	88	A
4	David	91	A

Question 10:

Code:

```
# Install missing package if needed
if (!require("corrplot")) install.packages("corrplot")

# Load libraries
library(dplyr)
library(ggplot2)
library(readr)
library(lubridate)
library(corrplot)

# Load dataset
avocado <- read_csv("C:/Users/Suman Bajani/OneDrive/Desktop/avocado.csv")

# View structure and dimensions
str(avocado)
cat("Number of rows:", nrow(avocado), "\n")
cat("Number of columns:", ncol(avocado), "\n")

# Check for duplicates and missing values
duplicates <- avocado[duplicated(avocado), ]
cat("Number of duplicate rows:", nrow(duplicates), "\n")
print(colSums(is.na(avocado)))
missing_summary <- sapply(avocado, function(x) sum(is.na(x)))
cat("Total missing values in dataset:", sum(is.na(avocado)), "\n")

# Clean missing values
avocado_clean <- na.omit(avocado)
```

```
cat("Rows after removing missing values:", nrow(avocado_clean), "\n")
```

```
# Summary statistics
```

```
mean_price <- mean(avocado_clean$AveragePrice)
```

```
median_price <- median(avocado_clean$AveragePrice)
```

```
sd_price <- sd(avocado_clean$AveragePrice)
```

```
cat("Mean:", mean_price, "\n")
```

```
cat("Median:", median_price, "\n")
```

```
cat("Standard Deviation:", sd_price, "\n")
```

```
# Group by type
```

```
avg_price_by_type <- avocado_clean %>%
```

```
  group_by(type) %>%
```

```
  summarise(AveragePrice = mean(AveragePrice))
```

```
print(avg_price_by_type)
```

```
# Standardize name (only if needed)
```

```
names(avocado_clean)[names(avocado_clean) == "Total Volume"] <- "Total.Volume"
```

```
# Scatter plot
```

```
# Scatter plot to visualize Total.Volume vs AveragePrice
```

```
plot <- ggplot(avocado_clean, aes(x = Total.Volume, y = AveragePrice)) +
```

```
  geom_point(alpha = 0.5, color = "darkgreen") +
```

```
  labs(title = "Total Volume vs Average Price",
```

```
        x = "Total Volume", y = "Average Price")
```

```
print(plot)
```

```
# Boxplot
```

```
ggplot(avocado_clean, aes(x = type, y = AveragePrice, fill = type)) +
```

```
  geom_boxplot() +
```

```
  labs(title = "Average Price by Avocado Type", x = "Type", y = "Average Price")
```



```
# Date conversion and line plot
avocado_clean$Date <- as.Date(avocado_clean$Date)
ggplot(avocado_clean, aes(x = Date, y = AveragePrice, color = type)) +
  geom_line() +
  labs(title = "Average Price Over Time by Avocado Type", x = "Date", y = "Average Price")
```

```
# Top 10 regions
top_regions <- avocado_clean %>%
  group_by(region) %>%
  summarise(AvgPrice = mean(AveragePrice)) %>%
  arrange(desc(AvgPrice)) %>%
  slice(1:10)
ggplot(top_regions, aes(x = reorder(region, AvgPrice), y = AvgPrice)) +
  geom_bar(stat = "identity", fill = "steelblue") +
  coord_flip() +
  labs(title = "Top 10 Regions by Average Price", x = "Region", y = "Average Price")
```

```
#Correlation
numeric_data <- select(avocado_clean, where(is.numeric))
cor_matrix <- cor(numeric_data, use = "complete.obs")
print(cor_matrix)
corrplot(cor_matrix, method = "color", type = "upper",
  tl.col = "black", tl.cex = 0.8,
  title = "Correlation Matrix", mar = c(0, 0, 1, 0))
```

#output:

Number of rows: 18249

Number of columns: 14

Number of duplicate rows: 0

Total missing values in dataset: 0

Rows after removing missing values: 18249

Mean: 1.405978

Median: 1.37

Standard Deviation: 0.4026766

A tibble: 2 × 2

type	AveragePrice
<chr>	<dbl>
1 conventional	1.16
2 organic	1.65

