# Faculty of Science and Technology

# **Project Cover Page**

Assignment Title:	Introduction t	Introduction to Data Science Midterm Project							
Assignment No:	N/A		Date of Submission:	14 December 2024					
Course Title:	INTRODUCTIO	N TO DATA SCIEN	ICE .						
Course Code:	CSC4180		Section:	В					
Semester:	Fall 2024-25		Course Teacher:	TOHEDUL ISLAM					

#### Declaration and Statement of Authorship:

- 1. I/we hold a copy of this Assignment/Case-Study, which can be produced if the original is lost/damaged.
- 2. This Assignment/Case-Study is my/our original work and no part of it has been copied from any other student's work or from any other source except where due acknowledgement is made.
- 3. No part of this Assignment/Case-Study has been written for me/us by any other person except where such collaborationhas been authorized by the concerned teacher and is clearly acknowledged in the assignment.
- 4. I/we have not previously submitted or currently submitting this work for any other course/unit.
- 5. This work may be reproduced, communicated, compared and archived for the purpose of detecting plagiarism.
- 6. I/we give permission for a copy of my/our marked work to be retained by the Faculty for review and comparison, including review by external examiners.
- 7. I/we understand thatPlagiarism is the presentation of the work, idea or creation of another person as though it is your own. It is a formofcheatingandisaveryseriousacademicoffencethatmayleadtoexpulsionfromtheUniversity. Plagiarized material can be drawn from, and presented in, written, graphic and visual form, including electronic data, and oral presentations. Plagiarism occurs when the origin of them arterial used is not appropriately cited.
- 8. I/we also understand that enabling plagiarism is the act of assisting or allowing another person to plagiarize or to copy my/our work.
- \* Student(s) must complete all details except the faculty use part.
- \*\* Please submit all assignments to your course teacher or the office of the concerned teacher.

Group Name/No.: 08

No	Name	ID	Program	Signature
1	DEVDOOT PARIAL	21-45061-2	BSc [CSE]	
2	ABDULLAH AL SHAHRIAR	21-44760-1	BSc [CSE]	
3	MD. SABBIR HOSSAIN KHAN	21-45256-2	BSc [CSE]	
4	GALIB HASAN ALVEE	21-44549-1	BSc [CSE]	

Faculty use only		
FACULTYCOMMENTS		
	Marks Obtained	
	Total Marks	

### **Dataset Description:**

This dataset is called "Depression student dataset" as mentioned in kaggle. This dataset is of depressed students based on various factors. Those factors are demographic, academic, and lifestyle factors. It includes attributes such as gender, age, academic pressure, study satisfaction, sleep duration, dietary habits etc. It shows a connection between various factors and the mental wellbeing of students. Here the class variable is "depression". So, it can help us to detect depression status. Also, it can help identify mental health risks and suggest strategies to improve wellbeing based on those factors after analyzing the dataset.

### **Attributes:**

**Gender:** Specifies the gender of the students. There are two genders in the dataset and they are male and female. It is a categorical attribute

Age: This indicates the age of the students. This is a numeric attribute

**Academic Pressure:** Levels of academic pressure experienced by the students. There are 5 levels, starting from 1 to maximum 5. This is a numeric attribute

**Study Satisfaction:** Satisfaction levels of the students with their studies. Here in the dataset, it is indicated in a range of 1 to 5. This is a numeric attribute

**Sleep Duration**: This indicates the hours of sleep students get per night. They are in 4 categories and those are "Less than 5 hours", "5-6 hours", "7-8 hours", "More than 8 hours". It is a categorical attribute.

**Dietary Habits**: Classification of dietary habits. They are "Unhealthy", "Moderate", "Healthy". It is a categorical attribute

**Have you ever had suicidal thoughts?:** Indicates whether students have had suicidal thoughts. The values are in Yes and No. It is a categorical attribute

**Study Hours:** This attribute indicates the time spent studying daily by the students. Value ranging from 0 to 12 in hours. This is a numeric attribute.

**Financial Stress:** Level of financial stress experienced. Ranging from 1 to 5. This is a numerical attribute.

**Family History of Mental Illness:** Indicates if there's a family history of mental illness of the students. The value is in Yes and No. This is a categorical attribute

**Depression:** Indicates whether the student is depressed or not and the value is in Yes and No. This a categorical column

### **Purpose:**

The main goal of this dataset is to study the relationship between mental health and various demographic, academic, and lifestyle factors among students. By identifying trends and patterns, the dataset aims to help detect depression and assess mental health risks. Insights derived from this analysis can guide the development of preventive measures and interventions to promote student well-being.

### **Project Overview**

This project explores the "Depression Student Dataset", analyzing how demographic, academic, and lifestyle factors impact student mental health. Key objectives include examining mental health patterns (e.g., links between sleep, diet, academic workload, and depression), identifying significant risk factors like family history of mental illness or suicidal thoughts, and proposing actionable interventions to improve mental well-being. The dataset supports descriptive and predictive analyses, using attributes such as gender, age, study hours, academic pressure, and dietary habits. Applications include tailoring preventive measures, providing academic support, and promoting healthy lifestyles to address mental health challenges and enhance student outcomes.

### **Data Preprocessing:**

# 1. Importing Dataset:

# **Coding Part:**

```
install.packages("openxlsx")
library(openxlsx)
mydata <- read.xlsx("C:/Users/sazin/Downloads/Midterm_Dataset.xlsx")
mydata</pre>
```

•	Gender	Age ÷	Academic_Pressure	Study_Satisfaction	Sleep_Duration ÷	Dietary_Habits	+ Have_you_ever_had_suicidal_thoughts.?	Study_Hours +	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	28		2 4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28		4 !	5 5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25		1	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23		1 4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31		1 !	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19		4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34		4	2 NA	Moderate	Yes	6	2	No	Yes
8	Female	20		4	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	NA		1 4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33		4	B Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31		5	5-6 hours	Healthy	Yes	NA	4	No	Yes
12	Male	24		2	1 7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23		5 !	5 NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25		1	1 5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21		5	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28		5	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23		5	2 More than 8 hours	Moderate	No	NA	4	No	NA
18	Female	23		1	NA NA	Healthy	Yes	0	3	No	No
19	Female	20		5	More than 8 hours	Unhealthy	Yes	2	5	No	Yes
20	Male	29		4	More than 8 hours	Unhealthy	Yes	1	3	No	Yes
21	Male	31		2	8 More than 8 hours	Unhealthy	No	3	3	Yes	No
22	Male	24		3 4	More than 8 hours	Healthy	Yes	1	3	No	No
23	NA	31		2	More than 8 hours	Unhealthy	Noo	10	1	No	No
24	Female	33		3	7-8 hours	Moderate	No	11	5	Yes	No
25	Female	33		2	7-8 hours	Moderate	Yes	12	5	Yes	NA
26	Male	31		2	2 7-8 hours	Healthy	No	2	4	Yes	No
27	Male	30		3 4	7-8 hours	Moderate	Yes	0	2	Yes	No
28	Male	21		5	3 7-8 hours	Unhealthy	No	6	4	Yes	Yes
29	Female	29		3 !	Less than 5 hours	Moderate	Yes	4	3	Yes	No
30	Female	34		3 4	Less than 5 hours	Unhealthy	Yes	12	1	Yes	No
31	Female	20		3	2 More than 8 hours	Healthy	No	2	2	No	No

**Explanation Part:** The openxlsx package was used to import data from an Excel file into R. First, the package was installed and loaded. Then, the read.xlsx() function was used to read the file "Midterm\_Dataset.xlsx" from the specified location into a data frame called mydata.

# 2. Structure of the Dataset:

# **Coding Part:**

```
str(mydata)
```

# **Output Part:**

**Explanation Part**: The str(mydata) function provides a detailed structure of the mydata dataset. It displays the type of each column (e.g., numeric, factor, character), the number of observations (rows), and the first few values of each column. This helps in understanding the data types and gives an overview of the dataset's organization.

### 3. Descriptive Statistics using summary Function:

### **Coding Part:**

```
summary(mydata)
```

### **Output Part:**

```
> summary(mydata)
   Gender
                               Academic_Pressure Study_Satisfaction Sleep_Duration
                                                                               Dietary_Habits
Length: 201
                Min. : 18.00 Min. : 1.000 Min. :1.000 Length:201
                                                                              Length:201
class :character 1st Qu.: 22.00 1st Qu.: 2.000 1st Qu.:2.000 Class :character Class :character
Mode :character Median : 26.00 Median : 3.000 Median :3.000 Mode :character Mode :character
                Mean : 28.25 Mean : 3.154 Mean :3.189
                3rd Qu.: 30.00 3rd Qu.: 4.000 3rd Qu.:4.000
                Max. :230.00 Max. :20.000 Max. :5.000
                NA's :3
Have_you_ever_had_suicidal_thoughts.? Study_Hours Financial_Stress Family,_History_of_Mental_Illness Depression
Length: 201
                              Min. : 0.000 Min. :1.00 Length:201
                                                                                        Length: 201
Class :character
                                1st Qu.: 3.000 1st Qu.:2.00 Class :character
                                                                                        Class :character
                                Median: 7.000 Median: 3.00 Mode: character
Mode :character
                                                                                    Mode :character
                                Mean : 6.332 Mean :2.93
                                 3rd Qu.:10.000 3rd Qu.:4.00
                                 Max. :12.000 Max. :5.00
                                 NA's :2
>
```

**Explanation Part:** The summary(mydata) function generates a statistical summary of the dataset, providing key metrics for numeric columns, such as the minimum, median, mean, and maximum values, along with quartiles. For categorical columns, it shows the count of each category. This summary helps in quickly understanding the distribution and characteristics of the data.

# 4. Visualizing Missing Values on a Graph:

# **4.1 Detecting Missing Values Column Wise:**

### **Code Part:**

```
colsums(is.na(mydata))
sum(is.na(mydata))
```

### **Output Part:**

**Explanation Part:** The colSums(is.na(mydata)) command calculates the number of missing values (NA) in each column of the dataset mydata. As we can see that there are total 5 attributes (Gender, Age, Sleep Duration, Study Hours, Depression) having missing values with the amount of missing values. sum(is.na(mydata) indicates the total number of missing values.

# 4.2 Detecting Row Indices of Missing Values in Each Column:

### **Code Part:**

```
lapply(mydata, function(col) which(is.na(col)))
```

# **Output Part:**

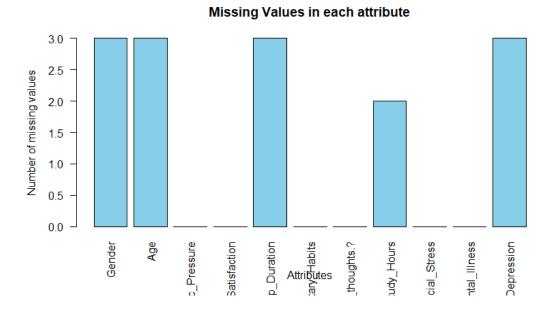
```
> lapply(mydata, function(col) which(is.na(col)))
$Gender
[1] 5 11 23
[1] 9 37 54
$Academic_Pressure
integer (0)
$Study_Satisfaction
integer(0)
$5leep_Duration
[1] 7 13 18
$Dietary_Habits
integer (0)
$`Have_you_ever_had_suicidal_thoughts.?`
integer(0)
$Study_Hours
[1] 11 17
$Financial_Stress
integer(0)
$Family._History_of_Mental_Illness
integer(0)
$Depression
[1] 13 17 25
```

**Explanation Part:** The command lapply (mydata, function(col) which(is.na(col))) identifies the row indices of missing values (NA) in each column of the dataset mydata.

# 4.3 Bar Plot of Missing Value (Graph):

### **Code Part:**

# **Output Part:**



**Explanation Part:** The code generates a bar plot to visualize the number of missing values in each attribute of the dataset. It uses the missing\_counts as parameter, which contains the count of missing values per column, and creates a bar plot with labels for the x-axis (attributes) and y-axis (number of missing values). The bars are colored sky blue, and the x-axis labels are rotated vertically for better readability. This plot helps identify which attributes have missing data.

# 4.4 Missing Values Visualization Gender Attribute:

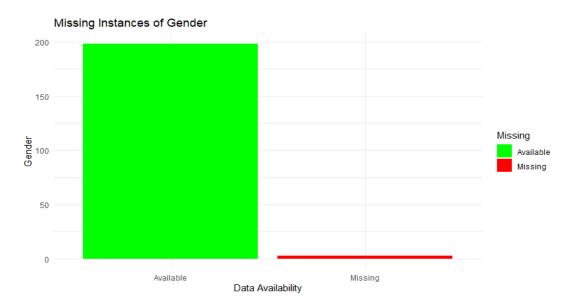
### **Code Part:**

```
install.packages("ggplot2")
library(ggplot2)

missingData <- data.frame(
   Missing = c("Available", "Missing"),
   Gender = c(sum(!is.na(mydata$Gender)), sum(is.na(mydata$Gender)))
)

ggplot(missingData, aes(x = Missing, y = Gender, fill = Missing)) +
   geom_bar(stat = "identity") +
   scale_fill_manual(values = c("Available" = "green", "Missing" = "red")) +
   labs(
      title = "Missing Instances of Gender",
      x = "Data Availability",
      y = "Gender"
   ) +
   theme_minimal()</pre>
```

### **Output Part:**



**Explanation Part:** The code creates a bar plot using ggplot2 to show the availability of data in the "Gender" column of the mydata dataset. It counts the number of available and missing values, then visualizes them with green bars for available data and red bars for missing data. The plot includes a title and axis labels, with a minimalistic theme for clarity. This helps in quickly assessing the completeness of the "Gender" data.

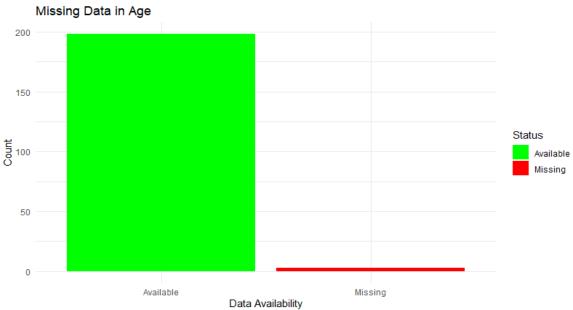
# 4.5 Missing Values Visualization Age Attribute:

### **Code Part:**

```
missingData_Age <- data.frame(
   Status = C("Available", "Missing"),
   Count = c(sum(!is.na(mydata$Age)), sum(is.na(mydata$Age)))
)

ggplot(missingData_Age, aes(x = Status, y = Count, fill = Status)) +
   geom_bar(stat = "identity") +
   scale_fill_manual(values = C("Available" = "green", "Missing" = "red")) +
   labs(
      title = "Missing Data in Age",
      x = "Data Availability",
      y = "Count"
   ) +
   theme_minimal()</pre>
```

### **Output Part:**



**Explanation Part:** The code creates a bar plot using ggplot2 to visualize the availability of data in the "Age" column of the mydata dataset. It counts the available (non-missing) and missing values for the "Age" column, then plots them as bars. Green bars represent available data, and red bars represent missing data. The plot includes a title and axis labels, with a minimalistic theme for clarity. This helps in assessing how complete the "Age" data is in the dataset.

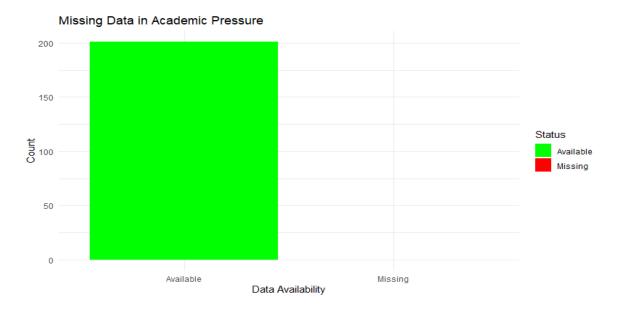
# 4.6 Missing Values Visualization Academic Pressure Attribute:

#### **Code Part:**

```
missingData_AcademicPressure <- data.frame(
    Status = c("Available", "Missing"),
    Count = c(sum(!is.na(mydata$Academic_Pressure)), sum(is.na(mydata$Academic_Pressure)))
)

ggplot(missingData_AcademicPressure, aes(x = Status, y = Count, fill = Status)) +
    geom_bar(stat = "identity") +
    scale_fill_manual(values = c("Available" = "green", "Missing" = "red")) +
    labs(
        title = "Missing Data in Academic Pressure",
        x = "Data Availability",
        y = "Count"
    ) +
    theme_minimal()</pre>
```

### **Output Part:**



**Explanation Part:** The code generates a bar plot using ggplot2 to visualize the availability of data in the "Academic Pressure" column of the mydata dataset. It counts the available (non-missing) and missing values for this column, then plots the results with green bars for available data and red bars for missing data. The plot includes a title and axis labels, with a minimalistic theme applied for simplicity. This visualization helps assess the completeness of the "Academic Pressure" data in the dataset.

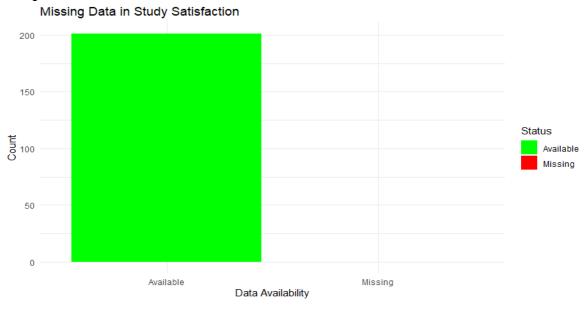
# 4.7 Missing Values Visualization Study Satisfaction Attribute:

#### **Code Part:**

```
missingData_StudySatisfaction <- data.frame(
   Status = c("Available", "Missing"),
   Count = c(sum(!is.na(mydata$Study_Satisfaction)), sum(is.na(mydata$Study_Satisfaction)))
)

ggplot(missingData_StudySatisfaction, aes(x = Status, y = Count, fill = Status)) +
   geom_bar(stat = "identity") +
   scale_fill_manual(values = c("Available" = "green", "Missing" = "red")) +
   labs(
        title = "Missing Data in Study Satisfaction",
        x = "Data Availability",
        y = "Count"
   ) +
   theme_minimal()</pre>
```

### **Output Part:**



**Explanation Part:** The code creates a bar plot using ggplot2 to display the availability of data in the "Study\_Satisfaction" column of the mydata dataset. It calculates the count of available (non-missing) and missing values for this column, then plots them with green bars for available data and red bars for missing data. The plot includes a title, axis labels, and uses a minimalistic theme for a clean and clear visualization. This helps in understanding the completeness of the "Study\_Satisfaction" data in the dataset.

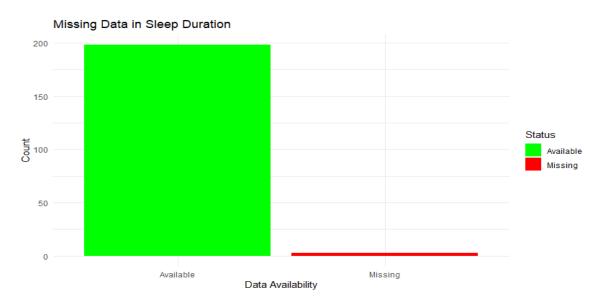
# 4.8 Missing Values Visualization Sleep Duration Attribute:

### **Code Part:**

```
missingData_SleepDuration <- data.frame(
    Status = c("Available", "Missing"),
    Count = c(sum(!is.na(mydata$Sleep_Duration)), sum(is.na(mydata$Sleep_Duration)))
)

ggplot(missingData_SleepDuration, aes(x = Status, y = Count, fill = Status)) +
    geom_bar(stat = "identity") +
    scale_fill_manual(values = c("Available" = "green", "Missing" = "red")) +
    labs(
        title = "Missing Data in Sleep Duration",
        x = "Data Availability",
        y = "Count"
    ) +
    theme_minimal()</pre>
```

### **Output Part:**



**Explanation Part:** The code generates a bar plot using ggplot2 to visualize the availability of data in the "Sleep Duration" column of the mydata dataset. It counts the available (non-missing) and missing values for this column, then creates a plot with green bars representing available data and red bars for missing data. The plot includes a title, axis labels, and uses a minimalistic theme for a clean design. This allows for a clear understanding of the completeness of the "Sleep Duration" data in the dataset.

# **5. Handling Missing Values:**

# **5.1 Detecting Missing values in Dataset:**

٠	Gender ‡	Age ‡	Academic_Pressure ÷	Study_Satisfaction	Sleep_Duration	Dietary_Habits	Have_you_ever_had_suicidal_thoughts.?	Study_Hours	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
7	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
8	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
9	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
LO	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
11	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE
12	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
13	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	TRUE
14	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
15	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
16	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
17	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	TRUE
18	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

# **5.2 Detecting Missing Values Row wise:**

### **Code Part:**

```
missing_rows <- which(rowSums(is.na(mydata)) > 0)
missing_rows
```

# **Output Part:**

```
> missing_rows <- which(rowSums(is.na(mydata)) > 0)
> missing_rows
5  7  9 11 13 17 18 23 25 37 54
5  7  9 11 13 17 18 23 25 37 54
> |
```

**Explanation Part:** The code identifies the rows in the mydata dataset that contain at least one missing value (NA). It does this by checking each row for missing values and then returns the indices of the rows with any missing data. This helps in identifying which rows require attention for data handling.

# **5.3 Discarding Instances of Missing Values:**

# **Code Part:**

mydata\_remove\_missing <- na.omit(mydata)
mydata\_remove\_missing</pre>

# **Initial Part:**

•	Gender ÷	Age ‡	Academic_Pressure ‡	Study_Satisfaction +	Sleep_Duration	Dietary_Habits +	Have_you_ever_had_suicidal_thoughts.?	Study_Hours	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	NA	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	NA	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31	5	4	5-6 hours	Healthy	Yes	NA	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	NA	4	No	NA
18	Female	23	1	3	NA	Healthy	Yes	0	3	No	No

# **Output Part:**

•	Gender ‡	Age ‡	Academic_Pressure	Study_Satisfaction	Sleep_Duration	Dietary_Habits	Have_you_ever_had_suicidal_thoughts.?	Study_Hours ÷	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
19	Female	20	5	5	More than 8 hours	Unhealthy	Yes	2	5	No	Yes
20	Male	29	4	3	More than 8 hours	Unhealthy	Yes	1	3	No	Yes
21	Male	31	2	3	More than 8 hours	Unhealthy	No	3	3	Yes	No
22	Male	24	3	4	More than 8 hours	Healthy	Yes	1	3	No	No
24	Female	33	3	2	7-8 hours	Moderate	No	11	5	Yes	No
26	Male	31	2	2	7-8 hours	Healthy	No	2	4	Yes	No
27	Male	30	3	4	7-8 hours	Moderate	Yes	0	2	Yes	No
28	Male	21	5	3	7-8 hours	Unhealthy	No	6	4	Yes	Yes
29	Female	29	3	5	Less than 5 hours	Moderate	Yes	4	3	Yes	No

**Explanation Part:** The code mydata\_remove\_missing <- na.omit(mydata) removes all rows from the mydata dataset that contain any missing values (NA). The na.omit() function creates a new dataset, mydata\_remove\_missing, which only includes complete rows (i.e., rows without any missing values). This is useful for cleaning the dataset before performing analysis or modeling, ensuring that only complete cases are considered.

# 5.4 Replacing Missing Values with Mean (Average) for Numeric Attribute:

### **Code Part:**

```
mydata[] <- lapply(mydata, function(x) {
   if (is.numeric(x)) {
      x[is.na(x)] <- mean(x, na.rm = TRUE)
      x <- round(x)
   }
   return(x)
}</pre>
```

### **Initial Part:**

٨	Gender ÷	Age ‡	Academic_Pressure	Study_Satisfaction	Sleep_Duration	Dietary_Habits ‡	Have_you_ever_had_suicidal_thoughts.?	Study_Hours ‡	Financial_Stress	FamilyHistory_of_Mental_Illness ÷	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	NA	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	NA	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31	5	4	5-6 hours	Healthy	Yes	NA	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	NA	4	No	NA
18	Female	23	1	3	NA	Healthy	Yes	0	3	No	No

•	Gender	Age ÷	Academic_Pressure +	Study_Satisfaction +	Sleep_Duration	Dietary_Habits	Have_you_ever_had_suicidal_thoughts.?	Study_Hours ÷	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	NA	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	28	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31	5	4	5-6 hours	Healthy	Yes	6	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	6	4	No	NA
18	Female	23	1	3	NA	Healthy	Yes	0	3	No	No
19	Female	20	5	5	More than 8 hours	Unhealthy	Yes	2	5	No	Yes
20	Male	29	4	3	More than 8 hours	Unhealthy	Yes	1	3	No	Yes

**Explanation Part:** The code processes the mydata dataset by replacing missing values (NA) in numeric columns with the mean of the respective column. It first checks if the column is numeric, then fills missing values with the mean of the non-missing values, ignoring any NAs. Afterward, it rounds the values in the column to the nearest integer. This approach helps handle missing data by imputing values with the mean and ensures that the data is rounded for consistency, making it ready for further analysis.

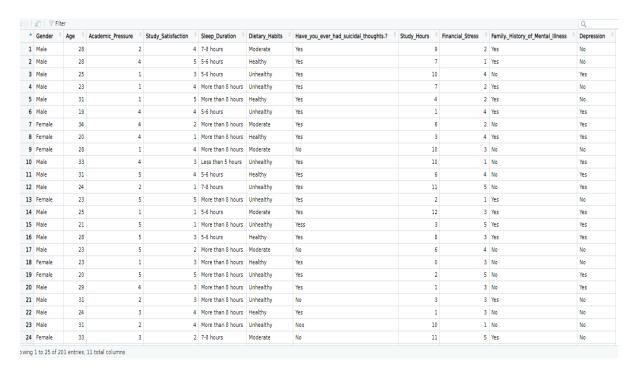
# **5.5** Replacing Missing Values with Mode (Most Frequent) for Categorical Attribute:

```
mydata[] <- lapply(mydata, function(x) {
   if (is.character(x) || is.factor(x)) {
     x[is.na(x)] <- names(which.max(table(x)))
   }
  return(x)
})</pre>
```

### **Initial Part:**

•	Gender ‡	Age ‡	Academic_Pressure	Study_Satisfaction	Sleep_Duration <sup>‡</sup>	Dietary_Habits <sup>‡</sup>	Have_you_ever_had_suicidal_thoughts.?	Study_Hours ‡	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	NA	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	NA	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31	5	4	5-6 hours	Healthy	Yes	NA	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	NA	4	No	NA
18	Female	23	1	3	NA	Healthy	Yes	0	3	No	No

### **Output Part:**



**Explanation Part:** The code addresses missing values (NA) in character or factor columns of the mydata dataset by replacing them with the most frequent value in each respective column. It identifies whether a column is of character or factor type, and for those with missing values, it replaces NA with the most common value (mode). This method ensures that categorical columns retain a meaningful and representative value where data is missing, helping to maintain the integrity of the dataset while handling missing data.

# **5.6** Replacing Missing Values with Forward Fill: Code Part:

```
forward_fill <- mydata
forward_fill[] <- lapply(forward_fill, function(column) {
  zoo::na.locf(column, na.rm = FALSE)
})</pre>
```

# **Initial Part:**

•	Gender ‡	Age ‡	Academic_Pressure	Study_Satisfaction	Sleep_Duration	Dietary_Habits $^{\circ}$	Have_you_ever_had_suicidal_thoughts.?	Study_Hours	Financial_Stress ÷	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	NA	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	NA	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31	5	4	5-6 hours	Healthy	Yes	NA	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	NA	4	No	NA
18	Female	23	1	3	NA	Healthy	Yes	0	3	No	No

# **Output Part:**

^	Gender	Age ‡	Academic_Pressure †	Study_Satisfaction	Sleep_Duration	Dietary_Habits =	Have_you_ever_had_suicidal_thoughts.?	Study_Hours	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	Male	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	5-6 hours	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	20	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	Male	31	5	4	5-6 hours	Healthy	Yes	10	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	7-8 hours	Unhealthy	Yes	2	1	Yes	Yes
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	8	4	No	Yes
18	Female	23	1	3	More than 8 hours	Healthy	Yes	0	3	No	No
19	Female	20	5	5	More than 8 hours	Unhealthy	Yes	2	5	No	Yes
20	Male	29	4	3	More than 8 hours	Unhealthy	Yes	1	3	No	Yes

**Explanation Part:** This code applies forward fill to all columns in the mydata data frame. A copy, forward\_fill, is created, and lapply iterates through each column, using zoo::na.locf to replace NA values with the last observed value(previous value) while preserving NA at the start if no prior value exists

### 5.7 Replacing Missing Values with Backward Fill:

### **Code Part:**

```
backward_fill <- mydata
backward_fill [] <- lapply(backward_fill, function(column) {
  zoo::na.locf(column, na.rm = FALSE, fromLast = TRUE)
})</pre>
```

### **Initial Part:**

*	Gender <sup>‡</sup>	Age ‡	Academic_Pressure ÷	Study_Satisfaction ‡	Sleep_Duration †	Dietary_Habits ‡	Have_you_ever_had_suicidal_thoughts.?	Study_Hours ‡	Financial_Stress <sup>‡</sup>	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	NA	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	NA	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31	5	4	5-6 hours	Healthy	Yes	NA	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	NA	4	No	NA
18	Female	23	1	3	NA	Healthy	Yes	0	3	No	No

•	Gender	Age ÷	Academic_Pressure †	Study_Satisfaction ‡	Sleep_Duration	Dietary_Habits ÷	Have_you_ever_had_suicidal_thoughts.?	Study_Hours +	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	Male	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	More than 8 hours	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	33	1	4	More than 8 hours	Moderate	No	10	3	No	No
10	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	Male	31	5	4	5-6 hours	Healthy	Yes	11	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	5-6 hours	Unhealthy	Yes	2	1	Yes	Yes
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	0	4	No	No
18	Female	23	1	3	More than 8 hours	Healthy	Yes	0	3	No	No
19	Female	20	5	5	More than 8 hours	Unhealthy	Yes	2	5	No	Yes
20	Male	29	4	3	More than 8 hours	Unhealthy	Yes	1	3	No	Yes

**Explanation Part:** This code applies backward fill to all columns in the mydata data frame. A copy, backward\_fill, is created to preserve the original data. Using lapply, it iterates through each column, applying zoo::na.locf with fromLast = TRUE to replace NA values with the next observed value.

# 6. Mean, Median, Variance, SD:

### 6.1 Numeric Mean:

```
numeric_means <- sapply(mydata, function(col) {
   if (is.numeric(col)) {
      mean(col, na.rm = TRUE)
   } else {
      NA
   }
})
numeric_means</pre>
```

```
> numeric_means <- sapply(mydata, function(col) {
   if (is.numeric(col)) {
     mean(col, na.rm = TRUE)
   } else {
   }
+ })
> numeric_means
                              Gender
                                                                                               Academic_Pressure
                                                                        Age
Study_Satisfaction
                                                                 28.233831
                                   NA
                                                                                                         3.154229
3.189055
                       Sleep_Duration
                                                             Dietary_Habits Have_you_ever_had_suicidal_thoughts.?
Study_Hours
6.348259
                     Financial_Stress
                                         Family._History_of_Mental_Illness
                                                                                                      Depression
                            2.930348
```

**Explanation Part:** The code calculates the mean of each numeric column in the mydata dataset, ignoring any missing values (NA). It does this by applying a function to each column, checking if the column is numeric, and then computing the mean of the non-missing values using mean(col, na.rm = TRUE). If the column is not numeric, it returns NA. This provides a summary of the means for all numeric columns in the dataset.

# 6.2 Numeric Median: Code Part:

```
numeric_median <- sapply(mydata, function(col) {
   if (is.numeric(col)) {
      median(col, na.rm = TRUE)
   } else {
      NA
   }
}
numeric_median</pre>
```

```
> numeric_median <- sapply(mydata, function(col) {
   if (is.numeric(col)) {
     median(col, na.rm = TRUE)
   } else {
   }
+ })
> numeric_median
                               Gender
                                                                                               Academic_Pressure
                  Study_Satisfaction
                                                            Sleep_Duration
                                                                                                  Dietary_Habits
Have_you_ever_had_suicidal_thoughts.?
                                                                Study_Hours
                                                                                                Financial_Stress
    Family._History_of_Mental_Illness
                                                                Depression
```

**Explanation Part:** The code calculates the median for each numeric column in the mydata dataset, ignoring any missing values (NA). It applies a function to each column, checks if the column is numeric, and then computes the median of the non-missing values using median (col, na.rm = TRUE). For non-numeric columns, it returns NA. This helps summarize the central tendency of all numeric columns in the dataset.

### **6.3 Numeric Variance:**

```
numeric_var <-sapply(mydata, function(x) {
   if (is.numeric(x)) {
     var(x, na.rm = TRUE)
   } else {
     NA
   }
}
numeric_var</pre>
```

```
> numeric_var <-sapply(mydata, function(x) {</pre>
  if (is.numeric(x)) {
   var(x, na.rm = TRUE)
  } else {
+ NA
+ }
+ })
> numeric_var
                             Gender
                                                                                         Academic_Pressure
                                                            425.850050
                                                                                                4.061095
                                NΔ
                  Study_Satisfaction
                                                         Sleep_Duration
                                                                                           Dietary_Habits
                          1.754080
                                                            Study_Hours
Have_you_ever_had_suicidal_thoughts.?
                                                                                          Financial_Stress
                                                            14.378109
                                                                                              1.995124
    Family._History_of_Mental_Illness
                                                             Depression
> |
```

**Explanation Part:** The code calculates the variance for each numeric column in the mydata dataset, ignoring any missing values (NA). It applies a function to each column, checks if the column is numeric, and then computes the variance using var(x, na.rm = TRUE), which removes NA values during the calculation. For non-numeric columns, it returns NA. This provides a measure of variability for all numeric columns in the dataset.

### **6.4 Numeric Standard Deviation:**

```
numeric_sd <-sapply(mydata, function(x) {
   if (is.numeric(x)) {
      sd(x, na.rm = TRUE)
   } else {
      NA
   }
})
numeric_sd</pre>
```

```
NΑ
> numeric_sd <-sapply(mydata, function(x) {
  if (is.numeric(x)) {
    sd(x, na.rm = TRUE)
   } else {
      NA
   - }
+ })
> numeric_sd
                                                                                              Academic_Pressure
                                                                20.636135
                                                                                                      2.015216
                  Study_Satisfaction
                                                           Sleep_Duration
                                                                                                Dietary_Habits
                            1.324417
Have_you_ever_had_suicidal_thoughts.?
                                                              Study_Hours
                                                                                              Financial_Stress
                                                                3.791848
                                                                                                      1.412489
    Family._History_of_Mental_Illness
                                                               Depression
>
```

**Explanation Part:** The code calculates the standard deviation for each numeric column in the mydata dataset, ignoring any missing values (NA). It applies a function to each column, checks if the column is numeric, and then computes the standard deviation using sd(x, na.rm = TRUE). For non-numeric columns, it returns NA. This helps measure the dispersion or spread of values for all numeric columns in the dataset.

# 7. Handling Duplicate Values:

# 7.1 Detecting Duplicate Values Row Wise:

```
duplicated_rows <- mydata[duplicated(mydata) | duplicated(mydata, fromLast = TRUE), ]
duplicated_rows</pre>
```

```
> duplicated_rows <- mydata[duplicated(mydata) | duplicated(mydata, fromLast = TRUE), ]</pre>
> duplicated_rows
  Gender Age Academic_Pressure Study_Satisfaction
                                                     Sleep_Duration Dietary_Habits
                                                 5 More than 8 hours
                                                                          Unhealthy
46 Male 20
47 Male 20
                              3
                                                 5 More than 8 hours
                                                                          Unhealthy
  Have_you_ever_had_suicidal_thoughts.? Study_Hours Financial_Stress Family._History_of_Mental_Illness Depression
46
                                                  11
                                      NΟ
                                                                                                                No.
                                                                    4
47
                                      No
                                                  11
                                                                                                     No
                                                                                                                No
> |
```

**Explanation Part:** The code identifies and returns all rows in the mydata dataset that are duplicated, either from the beginning or the end of the dataset. It uses the duplicated() function to detect duplicate rows and combines both directions of checking to capture all instances of duplication. This helps in identifying redundant data for further cleaning or analysis.

# 7.2 Removing Duplicated Values:

### **Code Part:**

mydata <-distinct(mydata)</pre>

# **Output Part:**

•	Gender	Age ‡	Academic_Pressure $^{\hat{\circ}}$	Study_Satisfaction $^{\circ}$	Sleep_Duration <sup>‡</sup>	Dietary_Habits	Have_you_ever_had_suicidal_thoughts.?	Study_Hours	Financial_Stress	FamilyHistory_of_Mental_Illness ÷	Depression
10	Female	21	3	3	7-8 hours	Moderate	Yes	8	5	Yes	Yes
1	Male	29	1	1	7-8 hours	Healthy	No	6	1	Yes	No
12	Male	22	1	3	7-8 hours	Unhealthy	No	6	4	No	No
13	Male	21	3	2	7-8 hours	Unhealthy	Yes	1	5	No	Yes
4	Male	31	5	4	5-6 hours	Healthy	No	12	3	No	No
5	Female	24	1	3	5-6 hours	Moderate	No	3	5	No	No
6	Male	20	3	5	More than 8 hours	Unhealthy	No	11	4	No	No
7	Female	31	1	3	7-8 hours	Healthy	No	12	3	Yes	No
8	Male	21	1	5	5-6 hours	Unhealthy	Yes	1	1	No	No
9	Male	24	2	4	5-6 hours	Healthy	No	12	4	Yes	No
0	Male	34	3	4	7-8 hours	Healthy	No	8	3	No	No
1	Female	25	5	4	Less than 5 hours	Healthy	No	7	1	No	No
2	Male	27	2	5	5-6 hours	Healthy	Yes	10	3	No	No
3	Female	28	2	4	5-6 hours	Moderate	No	10	1	Yes	No
4	Male	26	4	4	5-6 hours	Unhealthy	No	9	1	Yes	No
5	Male	23	2	5	Less than 5 hours	Unhealthy	No	8	5	No	No

**Explanation Part:** The code mydata <- distinct(mydata) removes any duplicate rows from the mydata dataset. It keeps only the unique rows, ensuring that the dataset contains no redundancy. This operation helps in cleaning the data by eliminating repeated entries, making it ready for analysis.

# **8. Imbalanced to Balanced Dataset:**

# 8.1 Detecting Imbalance Data:

### **Code Part:**

```
result <- lapply(mydata, function(x) {
   if (is.factor(x) | is.character(x)) {
     table(x)
   }
})
result</pre>
```

### **Output Part:**

```
> mydata <-distinct(mydata)
> result <- lapply(mydata, function(x) {
+    if (is.factor(x) | is.character(x)) {
+       table(x)
+    }
+ })</pre>
 $Gender
Female Male
88 112
$Academic_Pressure
$Study_Satisfaction
NULL
$5leep_Duration
                         7-8 hours Less than 5 hours More than 8 hours
52 48 51
$Dietary_Habits
x
Healthy Moderate Unhealthy
66 66 68
$`Have_you_ever_had_suicidal_thoughts.?`
  No Noo Yes Yess
92 1 106 1
$Study_Hours
$Financial_Stress
$Family._History_of_Mental_Illness
$Depression
No Yes
116 84
> |
```

**Explanation Part:** The code generates frequency tables for each factor or character column in the mydata dataset. It applies a function to each column, checking if the column is a factor or character type, and then counts the occurrences of each unique value using table(x). The result is a list of frequency tables, which helps to analyze the distribution of categorical data in the dataset and understand how often each value appears in the respective columns.

# 8.2 Under Sampling:

### **Code Part:**

```
mydata$Gender <- as.factor(mydata$Gender)
undersampled_data <- ovun.sample(Gender ~ ., data = mydata, method = "under", seed = 123)$data
table(undersampled_data$Gender)</pre>
```

### **Output Part:**

**Explanation Part:** The code converts the Gender column in the mydata dataset into a factor and then applies undersampling to balance the distribution of genders. The ovun.sample() function reduces the size of the majority class, ensuring that both classes in the Gender column have a similar number of instances.

# 8.3 Over Sampling:

### **Code Part:**

```
oversampled_data <- ovun.sample(Gender ~ ., data = mydata, method = "over", seed = 123)$data
table(oversampled_data$Gender)</pre>
```

### **Output Part:**

```
> oversampled_data <- ovun.sample(Gender ~ ., data = mydata, method = "over", seed = 123)$data
> table(oversampled_data$Gender)

Male Female
    112    108
> |
```

**Explanation Part:** The code applies oversampling to balance the Gender column in the mydata dataset by increasing the instances of the minority class. This ensures both classes have an equal number of records, helping to address class imbalance. The table() function then displays the distribution of genders after oversampling.

# 9. Outlier/Invalid data:

# 9.1 Detecting Invalid data:

```
errors <- lapply(mydata, function(col) {
  unique(col)
})
errors</pre>
```

```
> errors <- lapply(mydata, function(col) {
  unique(col)
+ })
> errors
$Gender
[1] "Male" "Female"
[1] 28 25 23 31 19 34 20 33 24 21 29 30 32 26 22 27 18 230 226
$Academic_Pressure
[1] 2 4 1 5 3 20 15
$Study_Satisfaction
[1] 4 5 3 2 1
$51eep_Duration
                      "5-6 hours" "More than 8 hours" "Less than 5 hours"
[1] "7-8 hours"
$Dietary_Habits
[1] "Moderate" "Healthy" "Unhealthy"
$`Have_you_ever_had_suicidal_thoughts.?`
[1] "Yes" "No" "Yess" "Noo"
$Studv_Hours
[1] 9 7 10 4 1 6 3 11 2 12 8 0 5
$Financial_Stress
[1] 2 1 4 3 5
$Family._History_of_Mental_Illness
[1] "Yes" "No"
$Depression
[1] "No" "Yes"
> |
```

**Explanation Part:** The code retrieves the unique values from each of those columns. This helps in understanding the distinct categories/values present in the attributes, which is needed to detect invalid data

# 9.2 Handling Invalid Data:

```
mydata$Have_you_ever_had_suicidal_thoughts.. <- gsub("Yess", "Yes", mydata$Have_you_ever_had_suicidal_thoughts..)
mydata$Have_you_ever_had_suicidal_thoughts.. <- gsub("Noo", "No", mydata$Have_you_ever_had_suicidal_thoughts..)
unique(mydata$Have_you_ever_had_suicidal_thoughts..)</pre>
```

```
> mydata$Have_you_ever_had_suicidal_thoughts.. <- gsub("Yess", "Yes", mydata$Have_you_ever_had_suicidal_thoughts..)
> mydata$Have_you_ever_had_suicidal_thoughts.. <- gsub("Noo", "No", mydata$Have_you_ever_had_suicidal_thoughts..)
> unique(mydata$Have_you_ever_had_suicidal_thoughts..)
[1] "Yes" "No"
> |
```

**Explanation** Part: The code corrects inconsistencies in the Have\_you\_ever\_had\_suicidal\_thoughts.. column by replacing incorrect values ("Yess" with "Yes" and "Noo" with "No") using the gsub() function. It then displays the unique values in the column to ensure the data is consistent and ready for analysis.

### 9.3 Identify Outlier:

### **Code Part:**

# **Output Part:**

Name	Type	Value
outliers_iqr	list [5]	List of length 5
Age	integer [2]	115 121
Academic_Pressure	integer [2]	88 94
Study_Satisfaction	integer [0]	
Study_Hours	integer [0]	
Financial_Stress	integer [0]	

**Explanation Part:** The code detects outliers in the numeric columns of the mydata dataset using the Interquartile Range (IQR) method. It calculates the first and third quartiles (Q1 and Q3) for each column, computes the IQR, and identifies values that fall outside the range of 1.5 times the IQR from the quartiles. The result is a list of indices marking the outliers in each numeric column.

### 9.4 Outlier Visualization in Boxplot:

### For Age Attribute:

#### Code Part:

boxplot(mydata\$Age, main = "Boxplot for Age", col = "skyblue")

### **Output Part:**

# 

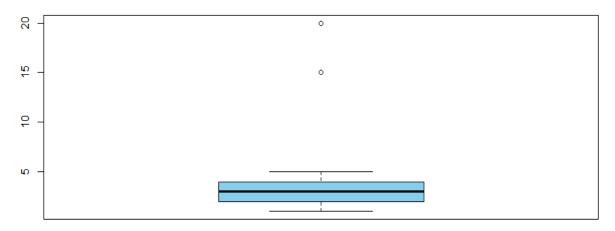
**Explanation Part**: The code creates a boxplot for the Age column in the mydata dataset. The boxplot() function visualizes the distribution of Age, showing the median, quartiles, and potential outliers. The main argument adds a title ("Boxplot for Age"), and the col argument colors the boxplot in "skyblue." This helps to understand the spread and potential outliers in the Age data.

# **Academic Pressure:**

### **Code Part:**

# **Output Part:**

#### **Boxplot for Academic Presssure**



**Explanation Part**: The code generates a boxplot for the Academic\_Pressure column in the mydata dataset. The boxplot() function displays the distribution of Academic\_Pressure, including the median, quartiles, and any potential outliers. The main argument adds the title "Boxplot for Academic Pressure," and the col argument colors the boxplot in "skyblue." This provides a visual representation of how Academic\_Pressure is distributed and helps identify any extreme values.

### 9.5 Replacing Outliers with Median:

#### **Code Part:**

### **Output Part:**

```
> mydata[sapply(mydata, is.numeric)] <- lapply(mydata[sapply(mydata, is.numeric)], function(x) {
+    Q1 <- quantile(x, 0.25, na.rm = TRUE)
+    Q3 <- quantile(x, 0.75, na.rm = TRUE)
+    IQR <- Q3 - Q1
+    x[x < (Q1 - 1.5 * IQR) | x > (Q3 + 1.5 * IQR)] <- median(x, na.rm = TRUE)
+    return(x)
+ }
+ return(x)
+ })
> table(mydata$Age)

18    19    20    21    22    23    24    25    26    26.5    27    28    29    30    31    32    33    34
    8    10    16    10    10    6    14    13    13    2    11    16    15    10    12    7    16    11
> table(mydata$Academic_Pressure)

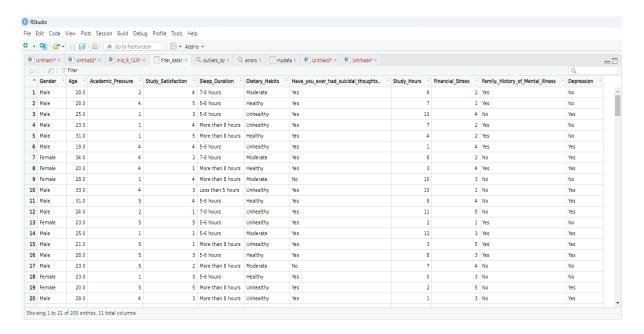
1    2    3    4    5
36    43    44    37    40
>
```

**Explanation:** The code processes the numeric columns in the mydata dataset by identifying and handling outliers. For each numeric column, it calculates the first and third quartiles (Q1 and Q3) and computes the Interquartile Range (IQR). Values that fall outside 1.5 times the IQR from the quartiles are considered outliers and are replaced with the median of the respective column. This approach ensures that extreme values do not skew the analysis, making the dataset more reliable and robust for further processing.

# 10. Filter Data:

### 10.1 Filter Data 1:

```
filter_data1 <- filter(mydata, Age >= 18 & Age <= 60)
```

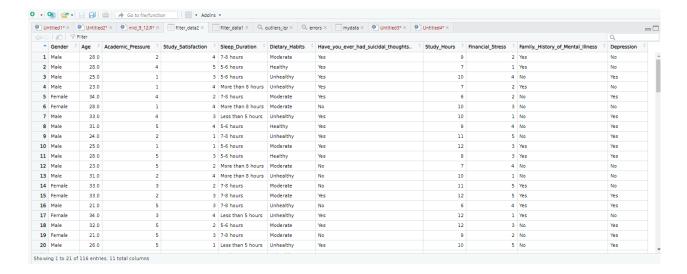


**Explanation:** The code filters the mydata dataset to include only the rows where the Age column values are between 18 and 60, inclusive. The filter () function from the dplyr package is used to select the subset of data that meets the condition Age  $\geq$  18 & Age  $\leq$  60. This helps narrow down the dataset to individuals within the specified age range for further analysis.

### 10.2 Filter Data 2:

### **Code Part:**

filter\_data2 <- filter(mydata, Age >= 18 & Age <= 60 & Study\_Hours > 5)



**Explanation:** The code filters the mydata dataset to include only the rows where the Age is between 18 and 60, and the Study Hours is greater than 5. The filter () function from the dplyr package is used with the condition Age >= 18 & Age <= 60 & study Hours > 5. This allows you to focus on individuals within the specified age range who also study more than 5 hours, refining the dataset for specific analysis.

### 10.3 Filter Data 3:

### **Code Part:**

filter\_data3 <- filter(mydata, Age >= 18 & Age <= 60 & Study\_Hours < 5)

	2 Y	Filter									Q,
٠	Gender ‡	Age ‡	Academic_Pressure <sup>‡</sup>	Study_Satisfaction	Sleep_Duration ÷	Dietary_Habits	<sup>‡</sup> Have_you_ever_had_suicidal_thoughts	Study_Hours	Financial_Stress	FamilyHistory_of_Mental_Illness	Depression
1	Male	31.0	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
2	Male	19.0	4	4	5-6 hours	Unhealthy	Yes		4	Yes	Yes
	Female	20.0	4	1	More than 8 hours	Healthy	Yes		3 4	Yes	Yes
	Female	23.0	5	5	5-6 hours	Unhealthy	Yes		2 1	Yes	No
5	Male	21.0	5	1	More than 8 hours	Unhealthy	Yes		5	Yes	Yes
6	Female	23.0	1	3	5-6 hours	Healthy	Yes		3	No	No
7	Female	20.0	5	5	More than 8 hours	Unhealthy	Yes		2 5	No	Yes
8	Male	29.0	4	3	More than 8 hours	Unhealthy	Yes		1 3	No	Yes
9	Male	31.0	2	3	More than 8 hours	Unhealthy	No		3	Yes	No
0	Male	24.0	3	4	More than 8 hours	Healthy	Yes		1 3	No	No
1	Male	31.0	2	2	7-8 hours	Healthy	No		2 4	Yes	No
12	Male	30.0	3	4	7-8 hours	Moderate	Yes		) 2	Yes	No
3	Female	29.0	3	5	Less than 5 hours	Moderate	Yes	4	3	Yes	No
14	Female	20.0	3	2	More than 8 hours	Healthy	No		2 2	No	No
15	Female	33.0	2	5	Less than 5 hours	Moderate	Yes		3 3	Yes	No
16	Male	26.0	5	4	7-8 hours	Unhealthy	No		3 2	Yes	No
17	Female	28.0	1	1	Less than 5 hours	Healthy	No		2 2	No	No
18	Male	26.0	4	5	Less than 5 hours	Unhealthy	No	4	1	Yes	No
19	Male	21.0	3	2	7-8 hours	Unhealthy	Yes		1 5	No	Yes
20	Female	24.0	1	3	5-6 hours	Moderate	No		5	No	No

**Explanation:** The code filters the mydata dataset to include only the rows where the Age is between 18 and 60, and the study Hours is less than 5. The filter () function from the dplyr package is applied with the condition Age  $\geq$  18 & Age  $\leq$  60 & study Hours  $\leq$  5. This creates a subset of the dataset focusing on individuals within the specified age range who study less than 5 hours, which can be used for further analysis.

# 11. Convert Attribute

# 11.1 Categorical to Numeric:

# **For Sleep Duration:**

```
> mydata$sleep_Duration <- factor(mydata$sleep_Duration,
+ levels = c("Less than 5 hours", "5-6 hours", "7-8 hours", "More than 8 hours"),labels = c(4,5.5,7.5,9))
> table(mydata$sleep_Duration)
4 5.5 7.5 9
48 47 51 55
```

_	Gender ÷	Age ÷	Academic_Pressure	Study_Satisfaction	Sleep_Duration	Dietary_Habits	Have_you_ever_had_suicidal_thoughts.?
1	Male	28	2	4	7.5	Moderate	Yes
2	Male	28	4	5	5.5	Healthy	Yes
3	Male	25	1	3	5.5	Unhealthy	Yes
4	Male	23	1	4	9	Unhealthy	Yes
5	Male	31	1	5	9	Healthy	Yes
6	Male	19	4	4	5.5	Unhealthy	Yes
7	Female	34	4	2	9	Moderate	Yes
8	Female	20	4	1	9	Healthy	Yes
9	Female	28	1	4	9	Moderate	No
10	Male	33	4	3	4	Unhealthy	Yes
11	Male	31	5	4	5.5	Healthy	Yes
12	Male	24	2	1	7.5	Unhealthy	Yes
13	Female	23	5	5	9	Unhealthy	Yes
14	Male	25	1	1	5.5	Moderate	Yes
15	Male	21	5	1	9	Unhealthy	Yess
16	Male	28	5	3	5.5	Healthy	Yes
17	Male	23	5	2	9	Moderate	No
18	Female	23	1	3	9	Healthy	Yes

**Explanation:** This code converts the Sleep Duration column in mydata into a factor with specific levels mapped to numeric labels (4, 5.5, 7.5, 9). The table () function then summarizes the counts of each factor level, showing how many entries correspond to each sleep category.

### For Gender:

```
\label{lem:mydata} $$\operatorname{Gender} \leftarrow \operatorname{factor}(\operatorname{mydata}\operatorname{Gender}, \ \operatorname{levels} = \operatorname{c}(\operatorname{"Male"}, \ \operatorname{"Female"}), \ \operatorname{labels} = \operatorname{c}(1,2))$$ $$ table(\operatorname{mydata}\operatorname{Gender})$$
```

```
> mydata$Gender <- factor(mydata$Gender, levels = c("Male", "Female"), labels = c(1,2))
> table(mydata$Gender)

1  2
112  88
> |
```

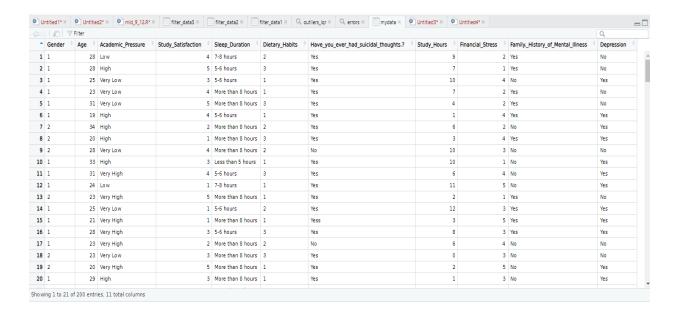
### **Explanation:**

The code converts the gender columns in the mydata dataset. For Gender, "Male" is labeled as 1 and "Female" as 2. The table () function is then used to display the frequency of each category in gender columns.

# 11.2 Numeric to Categorical:

### **Code Part:**

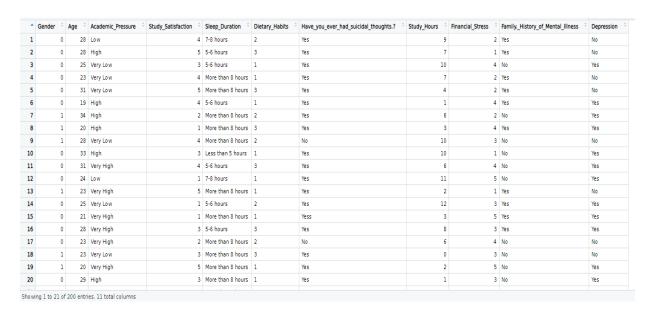
# **Output Part:**



**Explanation:** The code converts the Academic\_Pressure column in the mydata dataset from numeric values to descriptive labels. The numeric values 1 to 5 are mapped to "Very Low", "Low", "Moderate", "High", and "Very High". The table () function then shows the frequency of each category.

# 12. Normalization:

```
minMax <- function(x) {
   (x - min(x)) / (max(x) - min(x))
}
mydata$Gender <- as.numeric(as.factor(mydata$Gender))
mydata$Gender <- minMax(mydata$Gender)</pre>
```



**Explanation:** The code defines a minMax() function to normalize data between 0 and 1. It then converts the Gender column from categorical values ("Male" and "Female") into numeric codes. After that, the minMax() function is applied to the Gender column to scale the values between 0 and 1.

# 13. Conclusion:

In this dataset, initially there were missing values, outliers, invalid data and many more issues. So, the initial dataset is given below:

٨	Gender <sup>‡</sup>	Age ‡	Academic_Pressure ‡	Study_Satisfaction $^{\hat{\circ}}$	Sleep_Duration	Dietary_Habits	Have_you_ever_had_suicidal_thoughts.?	Study_Hours	Financial_Stress	FamilyHistory_of_Mental_Illness †	Depression
1	Male	28	2	4	7-8 hours	Moderate	Yes	9	2	Yes	No
2	Male	28	4	5	5-6 hours	Healthy	Yes	7	1	Yes	No
3	Male	25	1	3	5-6 hours	Unhealthy	Yes	10	4	No	Yes
4	Male	23	1	4	More than 8 hours	Unhealthy	Yes	7	2	Yes	No
5	NA	31	1	5	More than 8 hours	Healthy	Yes	4	2	Yes	No
6	Male	19	4	4	5-6 hours	Unhealthy	Yes	1	4	Yes	Yes
7	Female	34	4	2	NA	Moderate	Yes	6	2	No	Yes
8	Female	20	4	1	More than 8 hours	Healthy	Yes	3	4	Yes	Yes
9	Female	NA	1	4	More than 8 hours	Moderate	No	10	3	No	No
LO	Male	33	4	3	Less than 5 hours	Unhealthy	Yes	10	1	No	Yes
11	NA	31	5	4	5-6 hours	Healthy	Yes	NA	4	No	Yes
12	Male	24	2	1	7-8 hours	Unhealthy	Yes	11	5	No	Yes
13	Female	23	5	5	NA	Unhealthy	Yes	2	1	Yes	NA
14	Male	25	1	1	5-6 hours	Moderate	Yes	12	3	Yes	Yes
15	Male	21	5	1	More than 8 hours	Unhealthy	Yess	3	5	Yes	Yes
16	Male	28	5	3	5-6 hours	Healthy	Yes	8	3	Yes	Yes
17	Male	23	5	2	More than 8 hours	Moderate	No	NA	4	No	NA
8	Female	23	1	3	NA	Healthy	Yes	0	3	No	No

# And this is after applying data preparation steps, our dataset looks like this:

^ Gen	nder ‡	Age	Academic_Pressure	Study_Satisfaction	Sleep_Duration	Dietary_Habits ‡	Have_you_ever_had_suicidal_thoughts	Study_Hours	Financial_Stress ‡	FamilyHistory_of_Mental_Illness +	Depression
1	1	28	Low	4	7.5	Moderate	Yes	9	2	Yes	No
2	1	28	High	5	5.5	Healthy	Yes	7	1	Yes	No
3	1	25	Very Low	3	5.5	Unhealthy	Yes	10	4	No	Yes
4	1	23	Very Low	4	9	Unhealthy	Yes	7	2	Yes	No
5	1	31	Very Low	5	9	Healthy	Yes	4	2	Yes	No
6	1	19	High	4	5.5	Unhealthy	Yes	1	4	Yes	Yes
7	0	34	High	2	9	Moderate	Yes	6	2	No	Yes
8	0	20	High	1	9	Healthy	Yes	3	4	Yes	Yes
9	0	28	Very Low	4	9	Moderate	No	10	3	No	No
10	1	33	High	3	4	Unhealthy	Yes	10	1	No	Yes
1	1	31	Very High	4	5.5	Healthy	Yes	6	4	No	Yes
12	1	24	Low	1	7.5	Unhealthy	Yes	11	5	No	Yes
13	0	23	Very High	5	9	Unhealthy	Yes	2	1	Yes	No
14	1	25	Very Low	1	5.5	Moderate	Yes	12	3	Yes	Yes
15	1	21	Very High	1	9	Unhealthy	Yes	3	5	Yes	Yes
16	1	28	Very High	3	5.5	Healthy	Yes	8	3	Yes	Yes
.7	1	23	Very High	2	9	Moderate	No	6	4	No	No
8	0	23	Very Low	3	9	Healthy	Yes	0	3	No	No