

PRACTICAL NO :4

AIM: Applying conditional filters subset() or filter() in R.

OUTPUT:

The first screenshot shows the RStudio interface with the following code in the Source pane:

```
# Load libraries
library(dplyr)
library(readr)

# Load your BMI dataset
bmi <- read_csv("bmi.csv")
# Columns: 5
# Delimiter: ","
chr (1): BmiClass
dbl (4): Age, Height, Weight, Bmi

# Use 'spec()' to retrieve the full column specification for this data.
# Specify the column types or set 'show_col_types = FALSE' to quiet this message.

# View first few rows
head(bmi)
# A tibble: 6 x 5
  Age Height Weight Bmi BmiClass
  <dbl> <dbl> <dbl> <dbl> <chr>
1 61 1.85 109. 31.9 Obese Class 1
2 60 1.71 79.0 27.0 Overweight
3 60 1.55 74.7 31.1 Obese Class 1
4 60 1.46 35.9 16.8 Underweight
5 60 1.58 97.1 38.9 Obese Class 2
6 59 1.71 79.3 27.1 Overweight

# Filtering Examples on BMI Data
# Example 1: Filter people whose BMI > 30 (Obese)
obese_people <- subset(bmi, Bmi > 30)
# Number of obese people: 137
head(obese_people)
# A tibble: 6 x 5
  Age Height Weight Bmi BmiClass
  <dbl> <dbl> <dbl> <dbl> <chr>
1 61 1.85 109. 31.9 Obese Class 1
2 60 1.55 74.7 31.1 Obese Class 1
3 60 1.58 97.1 38.9 Obese Class 2
4 59 1.83 105. 31.3 Obese Class 1
5 57 1.75 110. 35.9 Obese Class 2
6 57 1.58 79.7 31.9 Obese Class 1
```

The second screenshot shows the RStudio interface with the following code in the Source pane:

```
# Example 2: Filter age > 50 AND Overweight/Obese
older_unhealthy <- subset(bmi, Age > 50 & Bmi > 25)
# Number of older_unhealthy: 44
head(older_unhealthy)
# A tibble: 6 x 5
  Age Height Weight Bmi BmiClass
  <dbl> <dbl> <dbl> <dbl> <chr>
1 61 1.85 109. 31.9 Obese Class 1
2 60 1.71 79.0 27.0 Overweight
3 60 1.55 74.7 31.1 Obese Class 1
4 60 1.58 97.1 38.9 Obese Class 2
5 59 1.71 79.3 27.1 Overweight
6 59 1.7 73.3 25.4 Overweight

# Example 3: Only "Normal" or "Overweight" class using %in%
normal_or_overweight <- subset(bmi, BmiClass %in% c("normal", "Overweight"))
# Number of normal or overweight individuals: 166
head(normal_or_overweight)
# A tibble: 6 x 5
  Age Height Weight Bmi BmiClass
  <dbl> <dbl> <dbl> <dbl> <chr>
1 60 1.71 79.0 27.0 Overweight
2 59 1.71 79.3 27.1 Overweight
3 59 1.7 73.3 25.4 Overweight
4 59 1.72 85.3 28.8 Overweight
5 58 1.71 79.3 27.1 Overweight
6 58 1.7 73.4 25.4 Overweight

# Same Filtering using dplyr
# People shorter than 1.6m
short_height <- bmi |>
  filter(Height < 1.6)
# Number of short height individuals (< 1.6m): 72
head(short_height)
```

MVLU

The screenshot displays the RStudio environment with the following components:

- Source Editor:** Contains R code for data manipulation. The code defines a tibble with columns Age, Height, Weight, Bmi, and BmiClass. It filters for BMI between 18.5 and 24.9, identifies healthy individuals, and then filters for Obese Class 1 individuals.
- Environment Panel:** Lists global environment objects including heart, heart_multi_sort, heart_sorted_age, heart_sorted_cholesterol, high_hr_sorted, normal_or_overweight, obese_class1, obese_people, older_unhealthy, and short_height, along with their respective observation counts and variable counts.
- Files Panel:** Shows a file explorer view of the user's home directory, listing various files and folders such as AIM.docx, amazon.csv, amazon.csv.zip, anisproject.sql, anti_ragging.pdf, AVLabs Video Enhancer AI, Blackmagic Design, bmi.csv, checksql, CN.pdf, connection.sql, Custom Office Templates, database_db.sql, desktop.ini, Experience-Salary.csv, graph scala m2 1.py, heart.csv, and HINDI ANSWERS.docx.

```
# A tibble: 6 x 5
  Age Height Weight Bmi BmiClass
<dbl> <dbl> <dbl> <dbl> <chr>
1 60 1.55 74.7 31.1 Obese Class 1
2 60 1.46 35.9 16.8 Underweight
3 60 1.58 97.1 38.9 Obese Class 2
4 59 1.46 36 16.9 Underweight
5 58 1.47 36 16.7 Underweight
6 57 1.58 79.7 31.9 Obese Class 1

> # BMI between 18.5 & 24.9 (Healthy range)
> healthy_bmi <- bmi |>
+ filter(Bmi >= 18.5, Bmi <= 24.9)
>
> cat("Healthy BMI individuals:", nrow(healthy_bmi), "\n")
Healthy BMI individuals: 338
> head(healthy_bmi)
# A tibble: 6 x 5
  Age Height Weight Bmi BmiClass
<dbl> <dbl> <dbl> <dbl> <chr>
1 48 1.73 74.1 24.8 Normal Weight
2 47 1.73 74.2 24.8 Normal Weight
3 46 1.73 74.3 24.8 Normal Weight
4 46 1.86 70 20.2 Normal Weight
5 45 1.73 74.4 24.9 Normal Weight
6 45 1.84 68 20.1 Normal Weight

> # People who are Obese Class 1
> obese_class1 <- bmi |>
+ filter(BmiClass == "Obese Class 1")
>
> cat("Obese Class 1 individuals:", nrow(obese_class1), "\n")
Obese Class 1 individuals: 20
> head(obese_class1)
# A tibble: 6 x 5
  Age Height Weight Bmi BmiClass
<dbl> <dbl> <dbl> <dbl> <chr>
1 61 1.85 109. 31.9 Obese Class 1
2 60 1.55 74.7 31.1 Obese Class 1
3 59 1.83 105. 31.3 Obese Class 1
4 57 1.58 79.7 31.9 Obese Class 1
5 56 1.8 98.5 30.4 Obese Class 1
6 55 1.72 95 32.1 Obese Class 1
```