

Sheth L.U.J & Sir M.V College  
SAS/SPSS/R  
Practical no.3

Aim: Exploring data: View() or print() (R).

Output:

The screenshot shows the RStudio interface. The top menu bar includes File, Edit, Code, View, Run, Session, Build, Debug, Profile, Tools, Help. The title bar says "RStudio - 0 R Scripts". The left sidebar has "Project" and "File Explorer". The main area has "Environment", "History", "Connections", and "Tutorial". In the Environment pane, datasets are listed: "0 book1" (15 obs. of 16 variables), "0 heart" (1025 obs. of 14 variables), "0 iris" (150 obs. of 5 variables), and "0 my\_data" (130 obs. of 3 variables). The "Script" pane contains R code and its output. The code is:

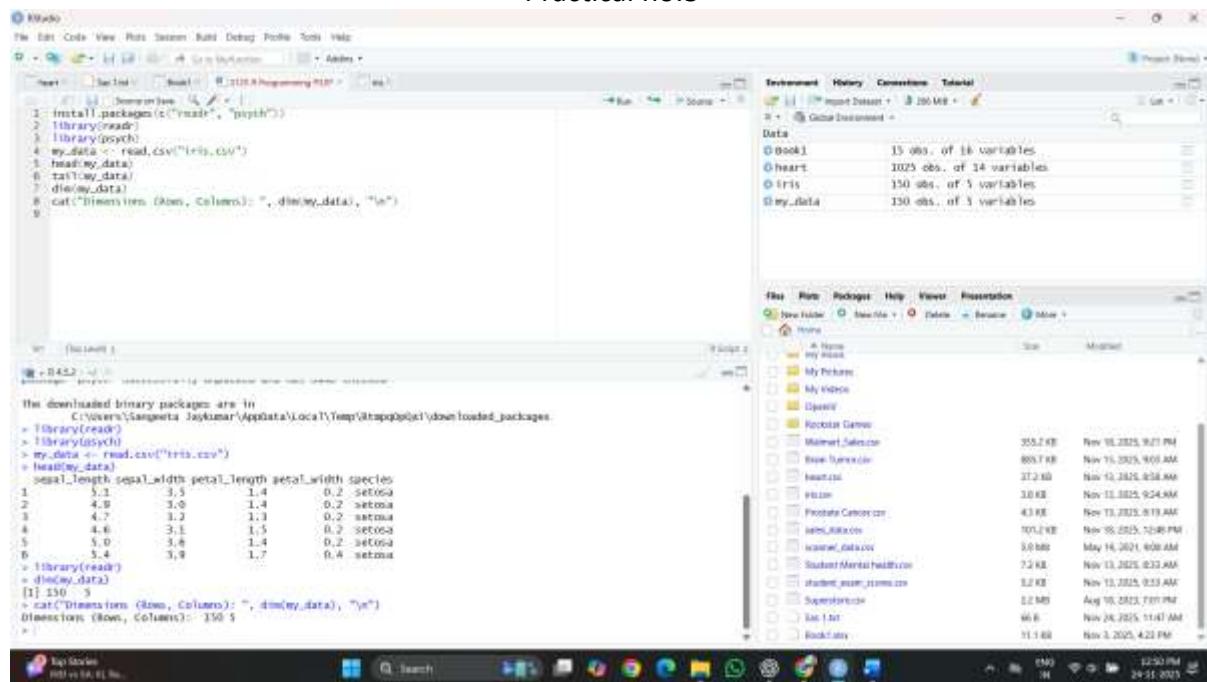
```
1 library(tidyverse)
2 library(psych)
3 my_data <- read.csv("iris.csv")
4 head(my_data)
```

The output shows the first few rows of the iris dataset:

	sepal.length	sepal.width	petal.length	petal.width	species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa

This screenshot is identical to the one above, showing the RStudio interface with the same project, environment, and script content. It displays the same dataset information and the same output for the iris dataset.

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```

install.packages("readr", "psych")
library(readr)
library(psych)
my_data <- read.csv("iris.csv")
tail(my_data)
display(my_data)
cat("Dimensions (Row, Columns): ", dim(my_data), "\n")

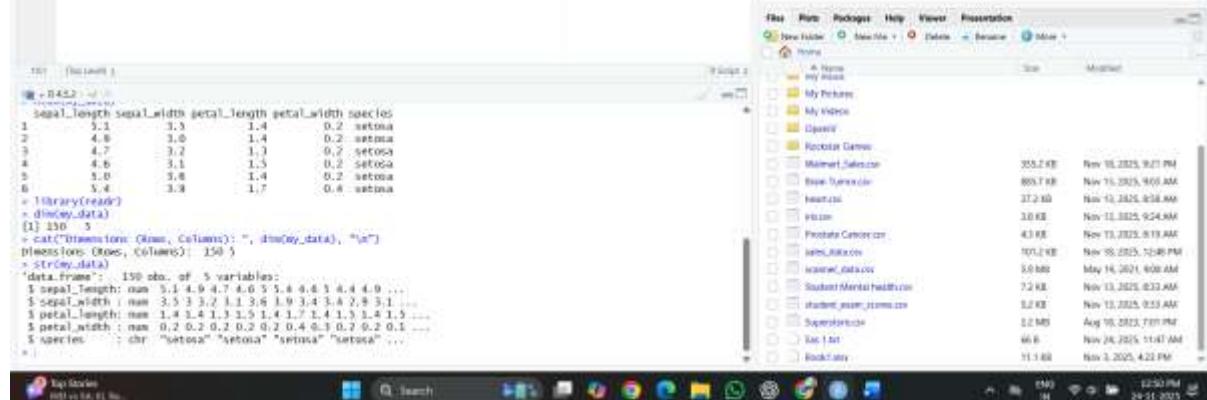
```



```

install.packages("readr", "psych")
library(readr)
library(psych)
my_data <- read.csv("iris.csv")
head(my_data)
sepal_length sepal_width petal_length petal_width species
1 5.1 3.5 1.4 0.2 setosa
2 4.9 3.0 1.4 0.2 setosa
3 4.7 3.2 1.3 0.2 setosa
4 4.6 3.1 1.5 0.2 setosa
5 5.0 3.6 1.4 0.2 setosa
6 5.4 3.9 1.7 0.4 setosa

```



```

install.packages("readr", "psych")
library(readr)
library(psych)
my_data <- read.csv("iris.csv")
head(my_data)
tail(my_data)
display(my_data)
cat("Dimensions (Row, Columns): ", dim(my_data), "\n")
dim(my_data)

```

Dimensions (Row, Columns): 150 5

str(my\_data)

sepal.length sepal.width petal.length petal.width species

1 5.1 3.5 1.4 0.2 setosa

2 4.9 3.0 1.4 0.2 setosa

3 4.7 3.2 1.3 0.2 setosa

4 4.6 3.1 1.5 0.2 setosa

5 5.0 3.6 1.4 0.2 setosa

6 5.4 3.9 1.7 0.4 setosa

10 |

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```

install.packages("readr", "magrittr")
library(readr)
iris_my_data<-read_csv("iris.csv")
head(my_data)
tail(my_data)
dim(my_data)
cat("Dimension (Row, Columns): ", dim(my_data), "\n")
str(my_data)
summary(my_data)

'data.frame': 150 obs. of 5 variables:
$ sepal_length: num 5.1 4.9 4.7 4.6 5.5 4.4 4.9 ...
$ sepal_width : num 3.5 3.2 3.1 3.0 3.4 3.4 2.9 3.1 ...
$ petal_length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.3 ...
$ petal_width : num 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
$ species     : chr "setosa" "setosa" "setosa" "setosa" ...
names(my_data)
summary(my_data)

```

```

install.packages("readr", "magrittr")
library(readr)
iris_my_data<-read_csv("iris.csv")
head(my_data)
tail(my_data)
dim(my_data)
cat("Dimension (Row, columns): ", dim(my_data), "\n")
str(my_data)
summary(my_data)
names(my_data)

```

```

'data.frame': 150 obs. of 5 variables:
$ sepal_length: num 5.1 4.9 4.7 4.6 5.5 4.4 4.9 ...
$ sepal_width : num 3.5 3.2 3.1 3.0 3.4 3.4 2.9 3.1 ...
$ petal_length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.3 ...
$ petal_width : num 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
$ species     : chr "setosa" "setosa" "setosa" "setosa" ...
names(my_data)
summary(my_data)

```

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