

**23BCE9360 – C N S SWAROOP**

**LAB – L57/L58**

# **FDA LAB ASSIGNMENT - 4**

**Dr. Voddelli Sri Lakshmi**

## FDA LAB ASSIGNMENT - 4

**Q1: Iris Flowers Description** This famous (Fisher or Anderson) iris data set gives the measurements in

centimetres of the variables sepal length and width and petal length and width, respectively, for 50

flowers from each of 3 species of iris. The species are *Iris setosa*, *versicolor*, and *virginica*.

Open iris data set in R and do the following

Print the dataset iris

Print the structure of the dataset iris

Print the summary of all the variables of the dataset iris (Hint: Use function `summary()`)

How many of the variables (columns) are in the dataset iris

How many observations (rows) are in the dataset iris

Print last 3 rows without using `tail()` function

Print the elements of 3rd column

Print all the elements of 2nd row

Print all the elements of 2nd, 5th and 13th row

Print all odd indexed rows (rows 1,3,5,...) (Hint: Use function - `seq`)

Print first three columns

Print the elements of columns from 5 to 10

Print the elements of columns from 3 to 7, 9 and 11

Print the elements of rows from 15 to 20

Print First, Second Row and Second and Third Column

Print element at 2nd row, third column

Print all the rows having "Sepal.Width" value greater than 1.4

Print all the rows having "Sepal.Width" value less than 2.5

Sort the observations of the dataset "iris" in increasing order based on the values in the column

"Sepal.Width"

Sort the observations of the dataset "iris" by column "Sepal.Width" in increasing order and column

"Petal.Width" in decreasing order

Use `duplicated()` function to print the logical vector indicating the duplicate values present in the

dataset iris

Extract duplicate elements from the dataset iris

Extract unique elements from the dataset iris

Print the indices of duplicate elements in the dataset iris

Print the indices of unique elements in the dataset iris

How many unique elements are in the dataset iris

How many duplicate elements are in the dataset iris

### CODE:

```
# Load dataset  
data(iris)
```

```
# 1. Print the dataset  
print(iris)
```

```
# 2. Print the structure of the dataset  
str(iris)
```

```
# 3. Print summary of all variables  
summary(iris)
```

```
# 4. Number of variables (columns)  
num_cols <- ncol(iris)  
print(num_cols)
```

```
# 5. Number of observations (rows)  
num_rows <- nrow(iris)  
print(num_rows)
```

```
# 6. Print last 3 rows without using tail() function  
print(iris[(nrow(iris) - 2):nrow(iris), ])
```

```
# 7. Print elements of 3rd column  
print(iris[, 3])
```

```
# 8. Print all elements of 2nd row  
print(iris[2, ])
```

```
# 9. Print all elements of 2nd, 5th, and 13th row  
print(iris[c(2, 5, 13), ])
```

```
# 10. Print all odd indexed rows  
print(iris[seq(1, nrow(iris), 2), ])
```

```
# 11. Print first three columns  
print(iris[, 1:3])
```

```
# 12. Print elements of columns from 5 to 10 (iris has only 5 columns, so this  
will show only the 5th column)  
print(iris[, 5, drop=FALSE])
```

```
# 13. Print elements of columns from 3 to 7, 9, and 11 (iris has only 5  
columns, so it cannot extract beyond 5)  
print(iris[, 3:5, drop=FALSE])
```

```
# 14. Print elements of rows from 15 to 20  
print(iris[15:20, ])
```

*# 15. Print First and Second Row, and Second and Third Column*

```
print(iris[1:2, 2:3])
```

*# 16. Print element at 2nd row, 3rd column*

```
print(iris[2, 3])
```

*# 17. Print all rows where "Sepal.Width" is greater than 1.4*

```
print(iris[iris$Sepal.Width > 1.4, ])
```

*# 18. Print all rows where "Sepal.Width" is less than 2.5*

```
print(iris[iris$Sepal.Width < 2.5, ])
```

*# 19. Sort dataset by "Sepal.Width" in increasing order*

```
print(iris[order(iris$Sepal.Width), ])
```

*# 20. Sort dataset by "Sepal.Width" (ascending) and "Petal.Width" (descending)*

```
print(iris[order(iris$Sepal.Width, -iris$Petal.Width), ])
```

*# 21. Use duplicated() to print logical vector for duplicate values*

```
print(duplicated(iris))
```

*# 22. Extract duplicate elements*

```
print(iris[duplicated(iris), ])
```

*# 23. Extract unique elements*

```
print(unique(iris))
```

*# 24. Print indices of duplicate elements*

```
print(which(duplicated(iris)))
```

*# 25. Print indices of unique elements*

```
print(which(!duplicated(iris)))
```

*# 26. Count unique elements*

```
num_unique <- nrow(unique(iris))
```

```
print(num_unique)
```

*# 27. Count duplicate elements*

```
num_duplicates <- nrow(iris) - num_unique
```

```
print(num_duplicates)
```

OUTPUT:

	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
7	4.6	3.4	1.4	0.3	setosa
8	5.0	3.4	1.5	0.2	setosa
9	4.4	2.9	1.4	0.2	setosa
10	4.9	3.1	1.5	0.1	setosa
11	5.4	3.7	1.5	0.2	setosa
12	4.8	3.4	1.6	0.2	setosa
13	4.8	3.0	1.4	0.1	setosa
14	4.3	3.0	1.1	0.1	setosa
15	5.8	4.0	1.2	0.2	setosa
16	5.7	4.4	1.5	0.4	setosa
17	5.4	3.9	1.3	0.4	setosa
18	5.1	3.5	1.4	0.3	setosa
19	5.7	3.8	1.7	0.3	setosa
20	5.1	3.8	1.5	0.3	setosa
21	5.4	3.4	1.7	0.2	setosa
22	5.1	3.7	1.5	0.4	setosa
23	4.6	3.6	1.0	0.2	setosa
24	5.1	3.3	1.7	0.5	setosa
25	4.8	3.4	1.9	0.2	setosa
26	5.0	3.0	1.6	0.2	setosa
27	5.0	3.4	1.6	0.4	setosa
28	5.2	3.5	1.5	0.2	setosa
29	5.2	3.4	1.4	0.2	setosa
30	4.7	3.2	1.6	0.2	setosa
31	4.8	3.1	1.6	0.2	setosa
32	5.4	3.4	1.5	0.4	setosa
33	5.2	4.1	1.5	0.1	setosa
34	5.5	4.2	1.4	0.2	setosa
35	4.9	3.1	1.5	0.2	setosa
36	5.0	3.2	1.2	0.2	setosa
37	5.5	3.5	1.3	0.2	setosa
38	4.9	3.6	1.4	0.1	setosa
39	4.4	3.0	1.3	0.2	setosa
40	5.1	3.4	1.5	0.2	setosa
41	5.0	3.5	1.3	0.3	setosa
42	4.5	2.3	1.3	0.3	setosa
43	4.4	3.2	1.3	0.2	setosa
44	5.0	3.5	1.6	0.6	setosa
45	5.1	3.8	1.9	0.4	setosa
46	4.8	3.0	1.4	0.3	setosa
47	5.1	3.8	1.6	0.2	setosa
48	4.6	3.2	1.4	0.2	setosa
49	5.3	3.7	1.5	0.2	setosa
50	5.0	3.3	1.4	0.2	setosa
51	7.0	3.2	4.7	1.4	versicolor
52	6.4	3.2	4.5	1.5	versicolor
53	6.9	3.1	4.9	1.5	versicolor
54	5.5	2.3	4.0	1.3	versicolor
55	6.5	2.8	4.6	1.5	versicolor
56	5.7	2.8	4.5	1.3	versicolor
57	6.3	3.3	4.7	1.6	versicolor
58	4.9	2.4	3.3	1.0	versicolor
59	6.6	2.9	4.6	1.3	versicolor
60	5.2	2.7	3.9	1.4	versicolor
61	5.0	2.0	3.5	1.0	versicolor
62	5.9	3.0	4.2	1.5	versicolor
63	6.0	2.2	4.0	1.0	versicolor
64	6.1	2.9	4.7	1.4	versicolor
65	5.6	2.9	3.6	1.3	versicolor
66	6.7	3.1	4.4	1.4	versicolor
67	5.6	3.0	4.5	1.5	versicolor
68	5.8	2.7	4.1	1.0	versicolor

```
'data.frame': 150 obs. of 5 variables:
 $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
 $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 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 1.5 ...
 $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
 $ Species : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1 ...
 Sepal.Length Sepal.Width Petal.Length Petal.Width
Min. :4.300 Min. :2.000 Min. :1.000 Min. :0.100
1st Qu.:5.100 1st Qu.:2.800 1st Qu.:1.600 1st Qu.:0.300
Median :5.800 Median :3.000 Median :4.350 Median :1.300
Mean :5.843 Mean :3.057 Mean :3.758 Mean :1.199
3rd Qu.:6.400 3rd Qu.:3.300 3rd Qu.:5.100 3rd Qu.:1.800
Max. :7.900 Max. :4.400 Max. :6.900 Max. :2.500
 Species
setosa :50
versicolor:50
virginica :50

[1] 5
[1] 150
 Sepal.Length Sepal.Width Petal.Length Petal.Width Species
148 6.5 3.0 5.2 2.0 virginica
149 6.2 3.4 5.4 2.3 virginica
150 5.9 3.0 5.1 1.8 virginica
 [1] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 1.5 1.6 1.4 1.1 1.2 1.5 1.3 1.4
 [19] 1.7 1.5 1.7 1.5 1.0 1.7 1.9 1.6 1.6 1.5 1.4 1.6 1.6 1.5 1.5 1.4 1.5 1.2
 [37] 1.3 1.4 1.3 1.5 1.3 1.3 1.3 1.6 1.9 1.4 1.6 1.4 1.5 1.4 4.7 4.5 4.9 4.0
 [55] 4.6 4.5 4.7 3.3 4.6 3.9 3.5 4.2 4.0 4.7 3.6 4.4 4.5

=== Execution Halted ===|
```