

	1	2	3	4	
			[1]	[2]	[3]
[1,]		1	3	5	
[2,]	2		4	6	

```
function(x)
{
  x[2]
  3
}
```

DAY 2 – LAB ASSESSMENT

Reg No:

Name:

19
20

1. Write a R program to create an array of two 3x3 matrices each with 3 rows and 3 columns from two given two vectors. Print the second row of the second matrix of the array and element in the 3rd row and 3rd column of the 1st matrix.

code:

```
mat1 <- matrix(1:9, nrow=3, byrow=TRUE)
mat2 <- matrix(10:18, nrow=3, byrow=TRUE)
```

```
arr <- array(c(m01, m01b), dim = c(3, 3, 2))
print(arr[2, , 2])
print(arr[2, 3, 1])
```

output:

12 24 16

2. Write a R program to combine three arrays so that the first row of the first array is followed by the first row of the second array and then first row of the third array.

code:

```
arr1 <- array(1:9, dim = c(3, 3))
arr2 <- array(10:18, dim = c(3, 3))
arr3 <- array(19:27, dim = c(3, 3))
combined_arr <- cbind(arr1[, ], arr2[, ], arr3[, ])
print(combined_arr)
```

output:

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]
[1,]	1	2	3	10	11	12	19	20	21

3. Write a R program to create an array using four given columns, three given rows, and two given tables and display the content of the array.

code:

```
col1 <- c(1, 2, 3)
col2 <- c(4, 5, 6)
col3 <- c(7, 8, 9)
col4 <- c(10, 11, 12)
row_names <- c("Row1", "Row2", "Row3")
table_names <- c("Table1", "Table2")
```

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```
my_array <- array (data = c (col1, col2, col3, col4), dim = c (3, 4, 2), dimnames = list (row_names, c ("col1", "col2", "col3", "col4"), table_names))
print (my_array).
```

output:

	col1	col2	col3	col4
Row1	1	4	7	10
Row2	2	5	8	11
Row3	3	6	9	12

	col1	col2	col3	col4
Row1	1	4	7	10
Row2	2	5	8	11
Row3	3	6	9	12

4. Write a R program to create a two-dimensional 5x3 array of sequence of even integers greater than 50.

code:

```
my_array <- matrix (seq (from = 52, by = 2, length.out = 15), nrow = 5)
```

```
print (my_array).
```

output:

	[, 1]	[, 2]	[, 3]
[1,]	52	62	72
[2,]	54	64	74
[3,]	56	66	76
[4,]	58	68	78
[5,]	60	70	80

Use Below Data frame from question 5 to 9

```
exam_data = data.frame(
  name = c('Anastasia', 'Dima', 'Katherine', 'James', 'Emily', 'Michael', 'Matthew', 'Laura', 'Kevin', 'Jonas'),
  score = c(12.5, 9, 16.5, 12, 9, 20, 14.5, 13.5, 8, 19),
  attempts = c(1, 3, 2, 3, 2, 3, 1, 1, 2, 1),
  qualify = c('yes', 'no', 'yes', 'no', 'no', 'yes', 'yes', 'no', 'no', 'yes')
)
```

5. Write a R program to extract 3rd and 5th rows with 1st and 3rd columns from a given data frame

6. Write a R program to add a new column named country in a given data frame

```
Country<-c("USA","USA","UK","USA","USA","USA")
example_data$country<-c("USA","USA","UK","USA","USA","USA")
```

extracted -	print (extracted - data)	output:	name	attempts
5 Emily			3 Katherine	2
				2

name	3 Katherine	5 Emily
extracted -		
print (extracted - data)		
output:		
attempts	2	2

name	score	attempts	qualify
Anastasia	12.5	1	Yes
Dema	9.0	3	Yes
Katherine	16.5	2	No
James	12.0	3	Yes
Emily	9.0	2	No
Michael	20.0	3	Yes
Matthew	14.0	1	Yes
Laura	13.0	1	no
Kevin	8.0	2	no
Jonas	19.0	1	Yes
Robert	10.0	1	Yes

8. Write a R program to sort a given data frame by name and score

name	score	attempts	qualify
Anastasia	12.5	1	Yes
Dema	9.0	3	No
Emily	9.0	2	no
James	12.0	3	no
Katherine	19.0	1	Yes
Kevin	16.5	2	Yes
Jonas	8.0	2	no
Laura	13.5	1	no
Matthew	14.5	1	Yes
Michel	20.5	3	Yes

[order(exam.data\$name, exam.data\$score)]

print(exam.data)

9. Write a R program to save the information of a data frame in a file and display the output.

code:
write.csv(exam.data, "exam.data.csv",
row.names = FALSE)

Exam.data -> from-file = read.csv
("exam.data.csv")

print(Exam.data -> from-file)

10. Write a R program to call the (built-in) dataset airquality. Check whether it is a data frame or not? Order the entire data frame by the first and second column. remove the variables

'Solar.R' and 'Wind' and display the data frame.

data(airquality)
if (is.data.frame(airquality))

{
print("airquality is a data frame.")

airquality -> ordered -> airquality[order(airquality\$Month, airquality\$Day)]
airquality -> ordered -> airquality[order(c("Ozone", "Month", "Day", "Temp", "Month.1"))]
print(airquality -> ordered)

output:

"airquality is a data frame"

	ozone	Month	Day	Temp	Month.1
1	41	5	1	67	5.1
2	36	5	2	72	5.1
3	12	5	3	74	5.1
4	18	5	4	62	5.1
5	NA	5	5	56	5.1
6	28	5	6	66	5.1
7	23	5	7	65	5.1
8	19	5	8	59	5.1
9	8	5	9	61	5.1
10	NA	5	10	69	5.1

11. Write a R program to create a factor corresponding to height of women data set, which inbuilt in R, contains height and weights for a sample of women.

code:

data(women)

```
women$height_factor <- cut
(women$height, breaks =
  quantile(women$height, probs =
    0:4/4),
  labels = c("short", "Medium-short",
    "Medium-Tall", "Tall"))
```

head(women)

output:

	height	weight	height_factor
1	58	115	Medium-short
2	59	117	Medium-short
3	60	120	Medium-short
4	61	123	Medium-Tall
5	62	126	Medium-Tall
6	63	129	Tall

12. Write a R program to extract the five of the levels of factor created from a random sample from the LETTERS (Part of the base R distribution.)

code:

set.seed(123)

letters_sample <- sample(LETTERS, 20)

letters_factor <- factor(letters_sample)

five_levels <- levels(letters_factor)[1:5]

print(five_levels)

output:

"A" "B" "C" "D" "E"

13. Iris dataset is a very famous dataset in almost all data mining, machine learning courses, and it has been an R build-in dataset. The dataset consists of 50 samples from each of three species of Iris flowers (Iris setosa, Iris virginica and Iris versicolor). Four features (variables) were measured from each sample, they are the length and the width of sepal and petal, in centimetres. Perform the following EDA steps.

(i) Find dimension, Structure, Summary statistics, Standard Deviation of all features.

(ii) Find mean and standard deviation of features grouped by three species of Iris flowers (Iris setosa, Iris virginica and Iris versicolor)

(iii) Find quantile value of sepal width and length

(iv) create new data frame named iris1 which have a new column name Sepal.Length.Cate that categorizes "Sepal.Length" by quantile

(v) Average value of numerical variables by two categorical variables: Species and Sepal.Length.Cate:

(vi) Average mean value of numerical variables by Species and Sepal.Length.Cate

(vii) Create Pivot Table based on Species and Sepal.Length.Cate.

code:

```
data(iris)
```

```
dim(iris)
```

```
str(iris)
```

```
summary(iris)
```

```
sapply(iris, sd)
```

```
aggregate(~Species, data=iris, mean)
```

```
aggregate(iris[, 1:4], by=list(Species=iris$Species), sd)
```

```
quantile(iris$Sepal.Width)
```

```
quantile(iris$Sepal.Length)
```

```
iris1 = iris %>% mutate(Sepal.Length.Cate = cut(Sepal.Length, breaks=quantile(Sepal.Length)))
```

```
aggregate(iris1[, 1:4], by=list(Species=iris1$Species, Sepal.Length.Cate = iris1$Sepal.Length.Cate), mean)
```

```
aggregate(iris1[, 1:4], by=list(Species=iris1$Species, Sepal.Length.Cate), FUN=function(x) mean(x))
```

```
iris1 %>% pivot_table(rows = Sepal.Length.Cate, cols = Species, value = c(Sepal.Width, Sepal.Length, Petal.Width, Petal.Length), agg.fun = mean)
```

output

150 5

setosa

sepal.length 5.1 4.9 4.7 4.6 5 5.4
4.6 5 4.4 4.9

sepal.width 3.5 3 3.2 3.1 3.6 3.9 3.4
3.4 2.9 3.1

sepal.length 1.4 1.4 1.3 1.5 1.4 1.7
1.4 1.5 1.4 1.5

sepal.width 0.2 0.2 0.2 0.2 0.2
0.4 0.3 0.2 0.2 0.1

Species : factor 10/3 levels : 1 1 1 1 1

Summary (vars) sepal.width species

Min.: 4.300 Min.: 2.000 setosa : 50
Q1: 5.100 Q1: 2.800 versicolor: 50
Median: 5.800 Median: 3.000 virginica: 50
Mean: 5.843 Mean: 3.051

1 setosa 5.006 3.428 1.462 0.246
2 versicolor 5.936 2.770 4.260 1.326
3 virginica 6.588 2.974 5.552 2.026

1 setosa 0.3525 0.3791 0.1737 0.1054
2 versicolor 0.5162 0.3138 0.4699 0.1978
3 virginica 0.6359 0.3225 0.5519 0.2744

0% 25% 50% 75% 100%
2.000 2.800 3.000 3.300 4.400

0% 25% 50% 75% 100%
4.300 5.100 5.800 6.400 7.900

15.13.8 1.4 0.2 setosa (4.3, 5.1)
24.9 3.0 1.4 0.2 setosa (4.3, 5.1)
3 4.7 3.2 1.3 0.2 setosa (4.3, 5.1)
4 4.6 3.1 1.5 0.2 setosa (4.3, 5.1)
5 5.0 2.6 1.4 0.2 setosa (4.3, 5.1)

species sepal.length petal.length
1 setosa (3.4, 5.1) 0.246
2 versicolor (4.2, 5.5) 1.000
3 virginica (4.2, 5.5) 1.3
4 setosa (5.1, 5.8) 0.2

14. Titanic Casualties – Use the standard ‘Titanic’ dataset which is part of R Base to answer the following questions.

- Use an appropriate apply function to get the sum of males vs females aboard.
- Get a table with the sum of survivors vs sex.
- Get a table with the sum of passengers by sex vs age

code:

data(Titanic)

apply(Titanic, c("sex", "Survived"), sum)

xtabs(Freq ~ Survived + sex, data = Titanic)

xtabs(Freq ~ Age + sex, data = margin(Titanic, 2))

survived

No

Yes

1364

367

126

344

survived

sex

Male

Female

No

1364

126

Yes

367

344

sex

Male

Female

1731

470

DAY 3 - LAB ASSESSMENT

Reg No:

No

Write a function in R programming to print generate Fibonacci sequence using

version in R

code

fib

function (n)

(=0)

Invalid input. n should be a positive integer."

if (n==1)

19
20
81