

TUTORIALSDUNIYA.COM

Data Science Notes

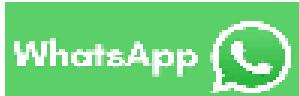
**Contributor: Kirtika
[SGGS CC DU]**

Computer Science Notes

Download **FREE** Computer Science Notes, Programs, Projects, Books for any university student of BCA, MCA, B.Sc, M.Sc, B.Tech CSE, M.Tech at
<https://www.tutorialsduniya.com>

Please Share these Notes with your Friends as well

facebook



21st Jan '19

Page No.		
Date		

Vectors

Creating Vectors

1) Using 'c' function

```
> x <- c(1, 7, 6, 9)
```

```
> x
```

```
[1] 1 7 6 9
```

class() & typeof() are some functn

```
> typeof(x)
```

```
[1] Numeric
```

eg. > y <- c(1, 2, TRUE, "a") # not error.

(`R' interpreter

precedence = logical → numeric → character

→ highest precedence.

```
> y
```

```
[1] "1" "2" "TRUE" "a"
```

2) Using infix function :

restrected works only on continuous values.

i.e. gap b/w data members = 1

If we want 10 values then using 'c' functn

```
> v <- c(1, 2, 3, ..., 10)
```

=

```
> v <- 1:10
```

means start with '1'

end .. '10'

:: means 1 gap b/w
members.

gap cannot be greater
than 1

eg > y <- 2:-2 ∴ y ≡ > y <- 2:-2
concatenates
two diff. stat.

but output of only 2 is shown.
O/P [1] 2 1 0 -1 -2

step size is still 1

eg $> z < \text{seq}(3, 4, 10, 2); z$

[1] 3.4 4.4 5.4 6.4 7.4 8.4 9.4

3) Using sequence functⁿ (seq())

Step size > 1

Syntax : $> \text{seq}(\text{from}=a, \text{to}=b, [\text{by}=\text{step-value}, \text{length.out}=\text{length-vector}])$

default step size = 1

(to - from
length.out - 1)

length
vector

eg $> x < \text{seq}(1, 7); x$ # $x \leftarrow 1:7$
[1] 1 2 3 ... 7 or

$x \leftarrow c(1, 2, 3, 4, 5, 6, 7)$

eg $> x < \text{seq}(1, 10, \text{by}=2); x$

[1] 1 3 5 7 9

eg $> x < \text{seq}(1, 10, \text{length.out}=5); x$
[1] 1 2 3 4 5 → x will have length of 5
(breaks into 5 equal parts &
assign 1:7 to x)

eg $> x < \text{seq}(1, 10, \text{length.out}=10)$

↳ x have length of 10

[1] 1.0 2.0 3.0 4.0 5.0 6.0 7.0 8.0 9.0 10.0

Note : If we specify all 4 parameters then seq() doesn't work as length.out breaks the values accordingly so step size() is not needed.

4) Using rep()

Syntax : $> f < \text{rep}(0, 5)$

value ↑, 5 ↘ how many times

[1] 0 0 0 0 0 value is repeated.

```
> f1 <- rep(1:3, 4); f1  
[1] 1 2 3 1 2 3 1 2 3
```

```
> length(f1)  
[1] 12
```

find no. of vectors

```
> h <- rep(4:6, 1:3); h  
[1] 4 5 5 6 6 6
```

* recycling

eg: h1 <- rep(1:10, 1:5)

$\overbrace{1 \ 2 \ 3 \ 4 \ 5}^{\text{if factor of 5}} \ \underbrace{6 \ 7 \ 8 \ 9 \ 10}_{\times 2}$

```
[1] 1 2 2 3 3 4 4 4 4 5 5 5 5 5
```

if not factor

eg: h1 <- rep(1:10, 1:4)

wanting is shown ↘

```
[1] 1 2 2 3 3 3 4 4 4 4  
5 6 6 7 7 7 8 8 8 8  
9 10 10
```

e → > x <- rep(1:2, each=2); x

↳ each individual element is repeated 2 times

OP: [1] 1 1 2 2 ↗ nms
> y <- rep(1:2, 2); y #rep(1:2, times=2)
[1] 1 2 1 2

⇒ Indexing a vector (Same as indexing array in any lang)

a[index]

① Doesn't start with '1'

② specify multiple index value.

e.g. a[[1, 2]]

returns two elements corr.
to 1st & 2nd index of a vector

or a[1]

> a[2]

③ Index values can be -ve'

e.g. > a ← c(1, 2, 3)

[1] [2] [3]
[1] 1 2 3

> a[-1]

[1] 2 3

all elements of vector

should be returned except
at index '1'.

possible
in
'R'

for (i=1; i>5; i++)
{
if (i%2 == 0) cout << a[i%2] = 0;
}

* ① > d ← seq(1, 5, by=0.5)

> d[3]

> d[5:7]

> d > 2.8

loops for every value in d for 2.8

> d[d > 2.8]

O/P:

d=[1.0 1.5 2.0 2.5 3.0 3.5 4.0 4.5 5.0]

[1] 2.0

[1] 3.0 3.5 4.0

[1] FALSE FALSE FALSE FALSE TRUE TRUE
TRUE TRUE

[1] 3.0 3.5 4.0 4.5 5.0

> length(d[d > 2.8])

Q)

$t \leftarrow c(0, 1, 2, 3, 4, 5, 6)$
 $x \leftarrow t[c(2, 3, 6)]$

print(x)

$v \leftarrow t[c(\text{TRUE}, \text{FALSE}, \text{FALSE})]$
print(v)

$x \leftarrow t[c(-2, -3)]$ print(x)
[1] 1 2 5

[1] TRUE FALSE FALSE 0
[1] 0 3 4 5 6

$v \leftarrow t[c(\text{TRUE}, \text{false}, \text{false}, \text{true})]$; q29.

eg.
eg.

q29 DIP [1] 0 3.

R-Studio

Write your script $x \leftarrow c(0, 1, 2)$	Variables in work environment. x.
Console DIP of current env	current packages, libraries, ...

> ls()

listing of all current variables (Object) in environment

> rm(x)

'x' is deleted from env.

> rm(c(x, y))

x, y deleted.

> rm(x)

> rm(y)

* > seq(1, 10, length.out = 5); d.

[1] 1.00 3.25 5.50 7.75 10.00

length.out includes 1 to 10 and after including these 2

3 more elements are to be included in a way
that all those are equally divided.

22 Jun '19

> assign ("y", c(1, 2)) # $y \leftarrow c(1, 2)$ or $y = c(1, 2)$

> y

[1] 1 2

> x <- c(1, 2, 9, 6, 5)

> sort(x); x

sort values in vector
order

[1] 1 2 5 6 9

> sort(x, decreasing = TRUE)

[1] 9 6 5 2 1

> sum(x)

sum of all elements of x

> mean(x)

give mean. sum(x)
(length(x))

> var(x)

st. deviation of x.

> y <- c(x, 0, x); y

[1] 1 2 9 6 5 0 1 2 9 6 5.

> 1/y. → 2 i z

[1] 1 1 1 1 1
2 9 6 5

> a <- c(1, 2, 3, 1, 2, 3)

> table(a)'

value of all values with
multiplicity.

a

1 2 3 → elements in vector

2 2 2 → multiplicity (freq) of these elements.

Inf → infinity

-Inf → -∞

Page No.

Date

> unique(a)

NA

give only unique vector
NA → not available

> x ← c(1, 2, NA, 3)

> length(x)

[1] 4

NA is element but not assigned

> x[1] ← 3. ; x

[1] 3 2 NA 3

> x[10] ←

NA

doesn't give errors for out of bound index.

* Vectors in R are dynamic i.e., length can be ↗

> length(x) ← 6 ; x # new length of x=6

[1] 3 2 NA 3 NA NA

NOTE *

The vectors consisting 'NA' as elements won't work with all functions

e.g. > x ← c(1, 2, NA, 3)

> sum(x)

[1] NA

(as NA+3 = NA)

→ to avoid NA use

check

> sum(x, na.rm=TRUE)

or
na.rm

[1] 6

NA is excluded.

> mean(x, na.rm=TRUE)

or
na.rm

> max(x) } both will work but
> sum(x) other function won't work

Ques ① > x = 1:8

> x[4] = -9 ; x

> y [cc(1,4)]

> x[x>3]

> x[x>3 & x<8]

> x[x %/% 2 == 0]

mod operator in R'

x = 1, 2, 3, 4, 5, 6, 7, 8

Ans. 2 1 8 3 4 5 6 7 8

> 1 2 3 -9 5 6 7 8

> 1 -9

> 4 5 6 7 8

> numeric(0)[1] 5 6 7

> [1] 2 6 8

② > y <- (1:5)^2 ; y → (1, 8, 25) ^ 2 = (1, 4, 9, 16, 25)

> y[2:4]

> y[-2:-4]

> y[cc(1, -1)] or y[(4, -1)] X not allowed.

> y[-1, 9]

y[-1]

> y[6]

NA

> y[1, 9]

1.9 is truncated to 1 y[1]

> y + 1

> y + 5.1

Mixing of (ve, -ve) indexing is not allowed.

③ $\begin{aligned}> a &\leftarrow \text{seq}(1:9, by=2); a \\&\geq \text{rep}(1:3, length.out=9) \rightarrow b; b \\&> a &\leftarrow a[1:4]; a \\&> a + \underbrace{2 * 1:5}_{2 \times (1, 2, 3, 4, 5)} - (2, 4, 6, 8, 10)\end{aligned}$

Soln ② $\begin{bmatrix} 1 \\ 4 \\ 1 \end{bmatrix}, \begin{bmatrix} 4 \\ 9 \\ 9 \end{bmatrix}, \begin{bmatrix} 16 \\ 16 \\ 25 \end{bmatrix}$

Error

$\begin{bmatrix} 1 \\ 4 \\ 9 \\ 16 \\ 25 \end{bmatrix}$

$\begin{bmatrix} NA \end{bmatrix}$

$\begin{bmatrix} 1 \end{bmatrix}$

$\begin{bmatrix} 2 \\ 5 \\ 10 \\ 17 \\ 26 \end{bmatrix}$

$\begin{bmatrix} 6.1 \\ 9.1 \\ 14.1 \\ 21.1 \\ 30.1 \end{bmatrix}$

③ $\begin{bmatrix} 1 \\ 3 \\ 5 \\ 7 \\ 9 \\ 11 \\ 13 \\ 15 \\ 17 \\ 19 \end{bmatrix}$

$\begin{bmatrix} 1 \\ 2 \\ 3 \\ 1 \\ 2 \\ 3 \\ 1 \\ 2 \\ 3 \end{bmatrix}$

$\begin{bmatrix} 8 \end{bmatrix}$

$\begin{bmatrix} 3 \\ 5 \\ 7 \\ 9 \\ 11 \end{bmatrix}$

⇒ functions

i) $\text{abs}(x)$ # absolute.

\sqrt{x}

$\text{ceiling}(x)$

$\text{floor}(x)$

$\text{round}(x, digits=n)$

$\text{trunc}(x)$

} works on vectors, arrays, list etc.

2018
Ques

Given a vector 'f' as $f \leftarrow c(0, 1, 1, 2, 3, 5, 8, 13, 21, 34)$
what is the output of following 'R' command?

- (i) $> f[1:3]$
(ii) $> f[-(1:3)]$
(iii) $> f < 10$
(iv) $f[f < 10]$
(v) $f[f[1:2] == 0]$.

Soln

- (i) 0 1 1
(ii) 2 3 5 8 13 21 34
(iii) T T T T T T F F F
(iv) 0 1 1 2 3 5 8
(v) 0 2 8 34

29th

Sep

29 Jan '19

Matrix \rightarrow 2D object

Syntax `matrix(data, nrow, ncol, byrow, dimnames)`

default is

defaut is "false"

eg: > a \leftarrow matrix(1:9, 3, 3, FALSE); a

$\begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}$

> a \leftarrow matrix(1:9, 3, 3, TRUE); a

$\begin{matrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{matrix}$

> c \leftarrow matrix(1:5, nrow=3, TRUE); c

$\begin{bmatrix} 1 \\ 2 \\ 3 \end{bmatrix}$

> c \leftarrow matrix(1:5, nrow=3, ncol=3, byrow=TRUE); c

Warning: In `matrix(1:5, nrow=3, ncol=3, byrow=TRUE)`
data length [5] is not a multiple of the
nrow * ncol [3]

$\begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}$

Call write out put warning

x[1,] 1 2 3
y[2,] 4 5 6
z[3,] 2 3 4

```
> row.names <- c("r1", "r2", "r3")  
> col.names <- c("c1", "c2", "c3")  
> p <- matrix(1:9, 3, 3, TRUE, dimnames = list(rownames, col.names))
```

dimnames is either a list/vector.

list() (c)
list(crownames, colnames)

if not col.names then only rows will be given name.

	c1	c2	c3
r1	9	8	7
r2	6	5	4
r3	3	2	1

```
> colnames(p)
```

[1] "c1" "c2" "c3"

```
> rownames(p)
```

[1] "r1" "r2" "r3"

```
> row.names <- c("A", "B", "C")
```

```
> col.names <- c("X", "Y", "Z")
```

```
> p
```

	X	Y	Z
A	9	8	7
B	6	5	4
C	3	2	1

```
> nrow(p)
```

[1] 3

takes matrix & returns no. of rows

```
> ncol(p)
```

[1] 3

takes matrix & returns no. of columns

```
> dim(p)
```

[1] 3 3

returns no. of dimensions

8

→ Changing vector into matrix (ways)

1) 1st method

> $x \leftarrow c(1, 2, 3, 4, 5, 6)$; x (Recycling allowed)
[1] 1 2 3 4 5 6
> $\text{dim}(x) \leftarrow c(2, 3)$ ← # default filled by col
> x
$$\begin{bmatrix} [1, 1] & [1, 2] & [1, 3] \\ [1] & 1 & 3 & 5 \\ [2, 1] & 2 & 4 & 6 \end{bmatrix}$$

> $\text{class}(x)$

[1] "matrix"

2) 2nd method [by default create a matrix type term $n \times 1$]

> $y \leftarrow 1:5$; y
[1] 1 2 3 4 5
> $\text{class}(y)$
[1] "numeric"
> $m \leftarrow y \text{ as.matrix}(y)$; m
[1] [1,]

[1] 1
[2] 2
[3] 3
[4] 4
[5] 5

TutorialsDuniya.com

Download FREE Computer Science Notes, Programs, Projects, Books PDF for any university student of BCA, MCA, B.Sc, B.Tech CSE, M.Sc, M.Tech at <https://www.tutorialsduniya.com>

- Algorithms Notes
- Artificial Intelligence
- Android Programming
- C & C++ Programming
- Combinatorial Optimization
- Computer Graphics
- Computer Networks
- Computer System Architecture
- DBMS & SQL Notes
- Data Analysis & Visualization
- Data Mining
- Data Science
- Data Structures
- Deep Learning
- Digital Image Processing
- Discrete Mathematics
- Information Security
- Internet Technologies
- Java Programming
- JavaScript & jQuery
- Machine Learning
- Microprocessor
- Operating System
- Operational Research
- PHP Notes
- Python Programming
- R Programming
- Software Engineering
- System Programming
- Theory of Computation
- Unix Network Programming
- Web Design & Development

Please Share these Notes with your Friends as well

facebook

WhatsApp 

twitter 

Telegram 

→ Combining vectors to form matrix

functions `cbind()` & `rbind()` are used to combine vectors to form a matrix.
`cbind()` will combine 'n' vectors by column
and `rbind()` " " " " row

```
> x <- 1:3  
> y <- 4:6
```

```
> m1 <- cbind(x, y) ; m1
```

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

```
> m2 <- rbind(x, y) ; m2
```

1 2 3
4 5 6

→ Accessing Elements of a matrix

Elements of a matrix can be accessed by using an index of the form `[i,j]`

```
> a
```

	[,1]	[,2]	[,3]
[1,]	1	2	3
[2,]	4	5	6
[3,]	7	8	9

> a[2,]

[1] 4

> a[2,]

[1] 4 5 6

> a[, 3]

[1] 3 6 9

$\frac{r=2\text{row}}{c=col}$

> a[c(1, 2), c(2, 3)]

#(multiple indexing in 1 statement)

- $\begin{matrix} [1, 1] & [1, 2] \\ [1, 2] & 3 \\ [2, 1] & 5 \end{matrix}$

→ it has nothing to do with
e.g

→ > a[c(3, 2),]

every col

~~c(1, 2) [2] [3]~~

$\begin{matrix} [1, 1] & 7 & 8 & 9 \\ [2, 1] & 4 & 5 & 6 \end{matrix}$

→ > a[c(3, 2)]

by default all col's w.r.t col '1' only

$\begin{matrix} [1, 1] & 7 \\ [2, 1] & 4 \end{matrix}$

> a[,] or a .

all rows & all cols

> a[-1,]

$\begin{matrix} [1, 1] & [1, 2] & [1, 3] \end{matrix}$

$\begin{matrix} [1, 1] & 4 & 5 & 6 \\ [2, 1] & 7 & 8 & 9 \end{matrix}$

* It is possible to index a matrix with a single vector while indexing in such a way it acts like a vector formed by stacking the cols of the matrix one after another.
The result is returned as a vector.

eg: $\geq a[1:4] \equiv a[1, 2, 3, 4]$

When vector is used for indexing then elements are stored in col wise

i.e., 1, 4, 7 2 5 8 3 6 9 → vector

[1] 1 4 7 2

* Using logical vectors as index

* Here rows & cols where the value is true will be returned. Also the logical vectors will be recycled if necessary.

* Logical vectors can be mixed with integer vectors for indexing

Recycling will be done in logical indexing on its own

eg: $\geq x$

	[1]	[2]	[3]
[1]	4	8	3
[2]	6	0	7
[3]	1	2	9

$\geq u[c(\text{TRUE}, \text{FALSE}, \text{TRUE}), c(\text{TRUE}, \text{TRUE}, \text{FALSE})]$
 \equiv
 $x[[1, 3], [1, 2]]$

[1,]	[1,]	[1,]
	4	8
[2,]	1	2

> $x[c(\text{TRUE}, \text{FALSE}), c(2,3)]$ # Recycling of index
=

$x[c(\text{TRUE}, \text{FALSE}, \text{TRUE}), c(2,3)]$
[1,]
[2,]

[1,]	8	3
[2,]	2	9

> $x[c(\text{TRUE}, \text{FALSE})]$

means all elements of x are stack together as element into a vector.

so vector will be 4 6 1 8 0 2 3 7 9.

so the recycling takes place as

T F T F T F T F T.

4 6 1 8 0 2 3 7 9.

[1] 4 1 0 3 9.

> $x[x > 5]$ #(store as 1-D & print elements greater than 5)

[1] 6 8 7 9

> x

[1,]	[1,]	[1,]
	2	3
[2,]	4	1

> $\det(x)$

determinant is returned

> Solve(x)

inverse of a matrix.

$$\begin{bmatrix} [1,1] & [1,2] \\ [2,1] & [2,2] \end{bmatrix}$$
$$\begin{bmatrix} 1.0 & -0.3 \\ -0.4 & 0.2 \end{bmatrix}$$

> sum(x)

$$[1] 20$$

sum of all elements of mat.

> prod(x)

$$[1]$$

product of all elements

> sum(x[1,])

$$[1] 5$$

sum of 1st row all cols

> y <- diag(3); y

used to create diag. mat.

$$\begin{bmatrix} [1,1] & [1,2] & [1,3] \\ [2,1] & 0 & 0 \\ [3,1] & 0 & 0 \end{bmatrix}$$

of 3x3 order.

> t(x)

transpose.

$$\begin{bmatrix} [1,1] & [1,2] \\ [2,1] & 2 & 4 \\ [2,2] & 3 & 1 \end{bmatrix}$$

> y <- matrix(c(1:4, 2, 2), 2, 2); y

$$\begin{bmatrix} [1,1] & [1,2] \\ [2,1] & 1 & 3 \\ [2,2] & 2 & 4 \end{bmatrix}$$

> x+y

element wise addit.

> x-y

" " " sub.

> x/y

" " " div.

> $x * y$

[Element wise matrix multiplication]

> $x \% * \% . y$

matrix multiplication

Ques:

write a R script to solve a system of linear eqn.

Soln.

$$a_1x_1 + b_1x_2 = c_1$$

$$a_2x_1 + b_2x_2 = c_2$$

$$\Rightarrow \begin{bmatrix} a_1 & b_1 \\ a_2 & b_2 \end{bmatrix} \begin{bmatrix} x_1 \\ x_2 \end{bmatrix} = \begin{bmatrix} c_1 \\ c_2 \end{bmatrix}$$

$$\begin{bmatrix} x_1 \\ x_2 \end{bmatrix} = \begin{bmatrix} a_1 & b_1 \\ a_2 & b_2 \end{bmatrix}^{-1} \begin{bmatrix} c_1 \\ c_2 \end{bmatrix}$$

1) $\rightarrow A \rightarrow$ matrix (, , ,)

2) $\rightarrow B \rightarrow$ matrix (, , ,)

{ 3) \leftarrow solve(A) } # inverse of A .
 { 4) ~~ans~~ $\leftarrow y \% * \% . B$ } //

OR solve(A, B) # (calculates inverse of A & multiplies it by B)

\Rightarrow Reading user Input

Prompt user with message & store value into N

* $n \leftarrow \text{as.integer}(\text{readline}(\text{prompt} = \text{"Enter no."}))$

Type casting used to read 1 line input \rightarrow input is returned as into integer. ↓ "string"
 either a no. or a string

* more than 1 element (no. or string)

> $x \leftarrow \text{scanf}()$ \leftarrow

1: 3 \leftarrow

2: 7
ctrl+z / ctrl+d }

to stop ~~input~~ entering input.

Read 2 inputs

msg shown as console

Ques

Take 2 vectors of size '2' as an input from the user and combine those vectors to form a matrix where the elements will be filled by row.

Soln

> a ← scan()

1: 1

2: 2

3: ctrl+Z / ctrl+D

read 2 items

> b ← scan()

1: 3

2: 4

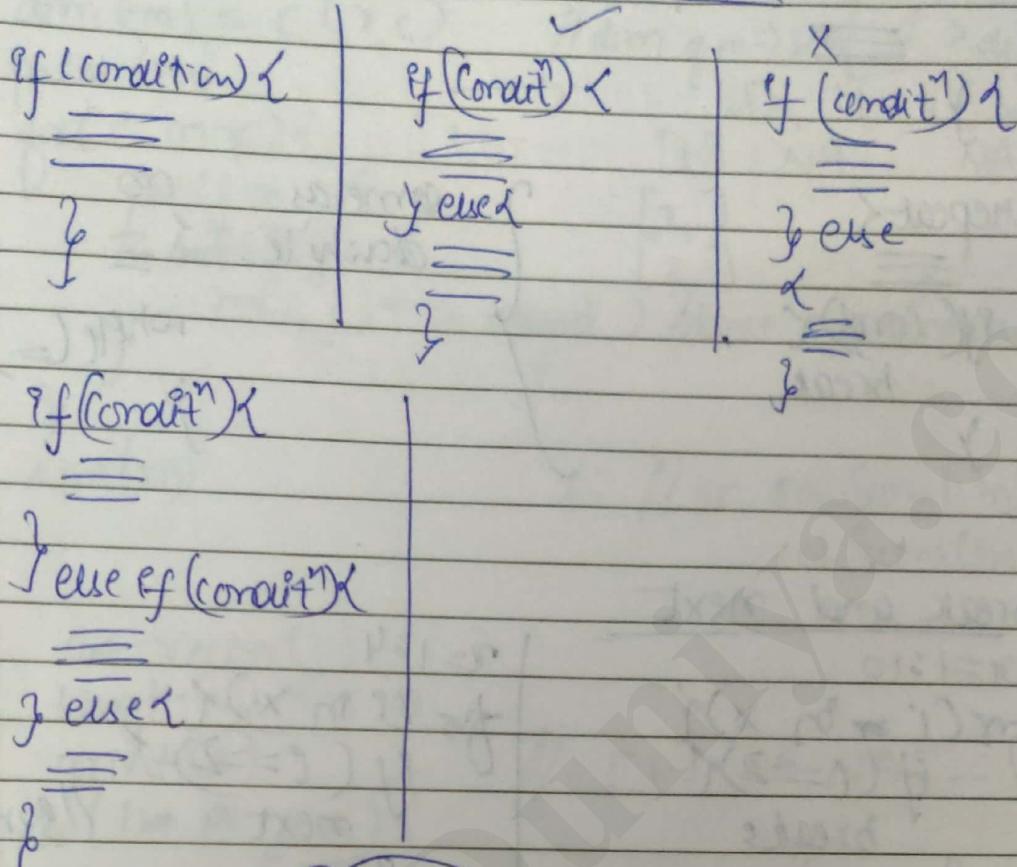
3:

read 2 items

> x ← rbind(a, b)

uth Fcb'19

Control structures in R



→ switch (exp, case1, case2, ...) → NO DEFAULT CASE

No multiple statements in 1 case.

No fall through since no specification of break.

If case1 is matched then execute case1.

→ Loops in R

`for (value in vector) {

}`

e.g. `for (i in 1:3) {
 print i
}`

→ while (test exp) {

 ≡

}.

→ repeat <

 ≡

if (cond) {
 break;

}.

}.

} same as
do
 while. <=

 where (=)

}

→ Break and next

n=1:10

for (i = 1; i < n; i++) {
 if (i == 2) {
 break;

}.

} print(i)

}.

[1] 1

x=1:4

for (c in x) {
 if (c == 2) {
 next

// ignore me

} print(c)

[1] 1

[1] 3

[1] 4

Q. Write a 'R' prog. to take a matrix as an input from user and print its sum. Also take the dimensions of the matrix as an input.

f← readline

#for (i ← 1; i < n; i++) {
 r← as.integer(readline(prompt="Enter rows"))
 c← as.integer(readline(prompt="Enter cols"))

for (i in 1:n)

length(m) $\leftarrow r * c$

m \leftarrow vector()

dim(m) = c(r, c)

print(m)

for(i in r){

for(j in c){

~~m[i][j]~~

m[i, j] \leftarrow a. integer

) // since no elements in vector.

}

sum(m)

// dim. of m \Rightarrow rxc
eg. >c = 1:4
dim(c) = c(2, 2)

[, 1]

[, 2]

[, 3]

[, NA]

[, NA]

[, NA]

[, ...]

c // or rowSums(m)

colSums(m)

II

a \leftarrow vector()

length(a) $\leftarrow 10$

print(a)

for(i ∞ in a){

}

a[i] $\leftarrow \dots$

// [1] NA NA ...

// input vector.

III

a, b vector

c vector

\rightarrow d sum

Q Write a prog. to input a matrix and
prog. that $A \cdot A^{-1} = I$.

Q write a prog. to find sum of main diag. of a matrix.

Q write a prog. to modify the main diag.
elements of a 3×3 matrix given as

$\begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}$ by the values entered by the user.

diag.(a) \leftarrow c(x, y, z)

Page No.	
Date	

- Q. WAP to find ~~whether~~ whether a no. is prime or not.
- Q. WAP to find factorial of no. point fibonnaci series upto ' n '.



Page No.			
Date			

12th feb '19

```
f←function (n){  
    s←0  
    m←n  
    while (n>0) {  
        r=n%10 # returns modulus  
        s=s+r*(r**r)  
        n=n//10 # integer div  
    }  
    print(m)  
    print(s)  
    if (m==s){  
        print("a")  
    } else {  
        print("b")  
    }  
}  
m←as.integer(
```

* by default whatever is the last variable used in the funct' (last statement) is returned irrespective of return statement.

Q/P: for n=

Arrays in R

$3 \times 4 \times 2$

2 arrays 3×4 .

Syntax: \rightarrow type can be anything. row \uparrow col matrix

array(data, dim = c(r, c, m), dimnames = list(rn, cn))

eg: $a \leftarrow \text{array}(1:24, \text{dim} = c(3, 4, 2))$ $\therefore 3 \times 4 \times 2 = 24$

eg $u_1 \leftarrow 1:12$

$u_2 \leftarrow 13:24$

$b \leftarrow \text{array}(c(u_1, u_2), \text{dim} = c(3, 4, 2))$ // may have used cbind()

1st array

$[1]$
 $[2]$
 $[3]$

, , 1
[1,1] [1,2] [1,3]

, , 2
[1,1] [1,2] [1,3]

[1]
[2]
[3].

\rightarrow 2nd array

Matrices using Array()

```
> p <- array(c(1:3, 3:1), dim = c(3, 2))
> row.names <- c("r1", "r2", "r3")
> col.names <- c("c1", "c2", "c3")
> row.names <- c("m1", "m2")
> b <- array(1:24, dim = c(3, 4, 2), dimname.list(row.names,
  col.names, m.names))
```

name(matrix) } checks.
attribute () }
mode () }

* `isNOMatrix()` → tells if anything is matrix or array

`a[4, 2, 1]` # 4
⁴ _{1st row & 2nd col q}
_{1st matrix}

, , m₁
 c₁ c₂ c₃
 r₁ 1 4 7
 r₂ 2 5 8
 r₃ 3 6 9

`a[, 2, 1]` # 456

`a[, ,]` # entire
 array

, , m₂
 c₁ c₂ c₃

`a[c(1), c(2, 3), 2]` # 13 16
^{1st row} _{2nd & 3rd col} _{2nd mat}
 tot

r₁ 10 13 16
 r₂ 11 14 17
 r₃ 12 15 18

`a[c(TRUE, FALSE, c(2, 3), 2)]` # 13, 16, 15, 19.

^{Recycling so}
_{1st & 3rd rows}

`a[-1, , , 1]` x.

`a[-1, , ,]`

functions available,

`sum(a)` # a[, 2, 1]

`nrow(a)`] check?

`ncol(a)`

2

~~if~~

→ Index Matrices.

using another array as index to an array

$x = \begin{bmatrix} 1 & 4 & 7 \\ 2 & 5 & 8 \\ 3 & 6 & 9 \end{bmatrix}$

$i = \begin{bmatrix} 2 & 1 \\ 1 & 2 \\ 3 & 3 \end{bmatrix}$

`x[i] = 4 5 4 9`

`i[x]`

// not possible

Negative indexes are not allowed!

TutorialsDuniya.com

Download FREE Computer Science Notes, Programs, Projects, Books PDF for any university student of BCA, MCA, B.Sc, B.Tech CSE, M.Sc, M.Tech at <https://www.tutorialsduniya.com>

- Algorithms Notes
- Artificial Intelligence
- Android Programming
- C & C++ Programming
- Combinatorial Optimization
- Computer Graphics
- Computer Networks
- Computer System Architecture
- DBMS & SQL Notes
- Data Analysis & Visualization
- Data Mining
- Data Science
- Data Structures
- Deep Learning
- Digital Image Processing
- Discrete Mathematics
- Information Security
- Internet Technologies
- Java Programming
- JavaScript & jQuery
- Machine Learning
- Microprocessor
- Operating System
- Operational Research
- PHP Notes
- Python Programming
- R Programming
- Software Engineering
- System Programming
- Theory of Computation
- Unix Network Programming
- Web Design & Development

Please Share these Notes with your Friends as well

facebook

WhatsApp 

twitter 

Telegram 

→ Lists Heterogeneous 1-D combination of data.

- * Vector $v \leftarrow c(1, 2, c(1, 2)) \neq v \leftarrow c(1, 2, 1, 2) X$
- * Vector homogeneous & list heterogeneous
- * list can have nested list but nested vectors are not available in R!

> list_1 \leftarrow list("Red", "Green", c(21, 31, 11), TRUE, 51.23)

> list_1

[1][]

[1] "Red"

1st element of list

1 O/P is Red.

[2][]

[1] "Green"

[3][]

[1] 21 32 11

> v
[1] 1, 2, ..., 20
[2] 21, 22, ..., 40

→ screen size can have 20 elements in 1 row

[1] → element corresponding to the
1st element in row.

[] → used to select element of 9th component
[[]] → " " " " components of list.

> list_1 [1]

[1][]

[1] "Red"

> list_1 [[1]]

[1] "Red"

→ Adding Elements to a list

'null' instead of 'NA'

> list[1][0] = c(1,1)
list[1][0] = c(1,1)

[1] <

> list[1]

[1]

;

;

[7]

NULL

[8]

[1] , 1

→ Naming Elements in List

stray

> list_data <- list(c("A", "B"), matrix(c(3, 9, 2), nrow = 2), + list("green", 12.3))

> names(list_data) <- c("1st quarter", "matrix", "List")

> list_data

\$ '1st quarter'
[1] "A" "B"

\$ 'matrix'

\$ 'List'

\$ 'list'[1]

[1] "green"

\$ 'list' [2]
[1] 12.3.

\$ 'list' [3]
[1]

~~2nd way~~

$x \leftarrow \text{list}(\text{day}=\text{"Tuesday"}, \text{lecture}=\text{"DS"}, \text{date}=c(13, 2, 2018))$
> x
\$ day
[1] "Tuesday"
\$ lecture.
[1] "DS"

\$ date
[1] 13 2 2018

> x\$date
[1] "Tuesday"
> x\$date
[1] 13 2 2018
> x\$date [1]
13

⇒ Merging lists

⊗ > is.list(x)
[1] TRUE

checks if n is matrix

> y = c(1)

> y = as.list(y)

> new_list = c(x, y)

changes vector into list
n elements comes first & then
y comes.

c() is also used to combine multiple list.

* no arithmetic operators on list-like $+,-,\ast, \%,$

>list1 + list2 #error

$\triangleright v_1 \leftarrow \text{unlist}(\text{list1})$

$\Rightarrow v2 \leftarrow \text{unlist}(\text{list} + 2)$

$$> v_1 + v_2$$

1st March in Al reg.

1) marking

factors

(^x DataScience)

factor(data_vector) or factor(data_vector, level_vector)

> data <- c("east", "west", "east", "north")

> is.factor(data)

[1] FALSE

> f1 <- Factor(data); f1

[1] east west east north.

or

Levels: east north west

levels are created alphabetically.

> f1 <- factor(data, levels = c("north", "east", "west"))

> class(f1) class(data)

numeric

> class(f1)

factor.

* functions

i) > levels(f1): returns level corr. to factor f1
O/P: "east" "north" "west"

> nlevels(f1): returns no. of levels.

[1] 3.

Factor

ordered

(Sorting is w.r.t to
levels)

unordered

(by default)

Mandatory to specify levels.

< if unordered
> $\max(f_1)$ or $\min(f_1)$
↓
gives max level of f_1
[1] min is not meaningful for factors

> is ordered (f_1) # tells if ordered or not
[1] FALSE

> $f_2 \leftarrow \text{factor}(\text{data}, \text{levels} = c("east", "north", "west"), \text{ordered} = \text{TRUE})$

> f_2

[1]

levels: east < north < west

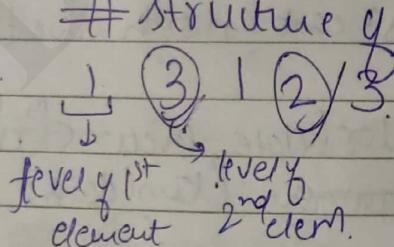
> $\max(f_2)$

[1] west

east → 1
west → 3
east → 2
north
west

> $\text{str}(f_1)$ # Structure of factor f_1

Factor w/ 3 levels



> $f_1[3]$

· east

> $f_1[c(2, 4)]$

· west north

> $f_1[c(\text{TRUE}, \text{FALSE})]$

· east east west

Recycling.

> $f_1[-1]$

> $\text{length}(f_1)$

[1] 5

> $f_1[6] \leftarrow \text{"south"}$

note as only 3 levels are available w.r.t new level # valid in R but not
met as only 3 levels are available so we can't

[1] Invalid factor level, NA generated.

```
> f1  
[1] ... - NA
```

→ Adding levels to existing factors.

```
> levels(f1) <- c(levels(f1), "south")  
old levels  
east north west
```

```
> nlevels(f1)  
[1] 4
```

```
> f1[6] <- "south" ; f1  
# south
```

→ Major application of factors (Interpretation of dataframes w.r.t visualization)

→ tapply(): this function is used to run a predefined function or a user defined function on a factor.

empid	region	income	income w.r.t each region
e	c	20/c	
s	w	30/c	
n			
w			

→ acc. to factor level mean is calculated & returned

```
> region <- c("c", "w", "e", "n", "s", "e", "n", "s")
```

```
> fr1 <- factor(region)
```

```
> income <- c(20, 40, 60, 80, 30, 30, 60, 40)
```

→ can be user defined function all

```
> mean_income <- tapply(income, fr1, mean)
```

→ data is independent on what factor

> mean - in same

e n s w
— — — —

→ avg value for each level.

Q. Differentiate b/w ordered / unorderd factors?

Ans

> items <- data.frame(color = c('red', 'blue', 'blue',
'red', 'green', 'green', 'yellow', 'blue',
'green', 'yellow', 'red', 'blue', 'blue', 'green',
'red', 'yellow'),
size = c(5, 10, 11, 6, 15, 16, 20, 9, 13, 18, 7, 14, 8,
13, 6, 18))

Items → factors

> tapply(items \$size, items \$color, sum)
blue green red yellow
52 57 24 56

> tapply(items \$size, items \$color, mean)

27th March '19

Unit 5 → 5-10 marks

Ch-10

> data()
list of whatever datasets are inbuilt
> data(iris)
for particular dataset in current environment
↳ naming dataset
> install.packages...
> str(iris)
structure of dataset
> summary(iris)
> head(iris)
first 5 records/rows
> head(iris, n=10)
10 .. "
> tail(iris)
last 5 rows.

In practical it runs 6 but in theory 5 rows.

VIVA

Airquality, pressure

6.2

> max(pressure & temperature)
[1] 360.

Quartile, decile, percentile

4 10 100

> quantile(pressure & temperature)

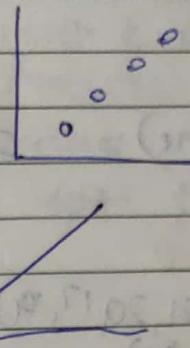
> Summary gives min, 1st quartile, median, mean,
3rd quart. max.
↳ summary(prsmes & temperature)

> nrow(iris)
> ncol(iris)
> sum(iris)

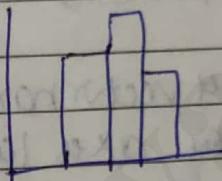
no. of rows
.. 150

⇒ Graphical (Ch-10)
⇒ $n \leftarrow 1 : 10$
⇒ $\text{plot}(n)$

Scattered plot graph
line graph



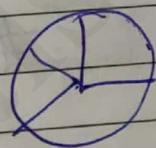
~~for plot~~ Bar plot



Box plot

Histogram

Pie chart



1) Scattered plot → (intervals can't be specified)
 $\text{plot}(n, y, \text{main}, \text{xlab}, \text{ylab}, \text{xlim}, \text{ylim}, \text{axes})$

$\text{xlab} = "xlabel"$

→ mandatory parameter

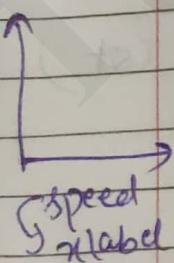
$\text{ylab} = "y"$

$\text{xlim} = "xlimits" \rightarrow \text{xlim} = c(0, 50)$

$\text{ylim} = "ylimits"$

$\text{axes} = "true" \rightarrow \text{to draw axes (default)}$

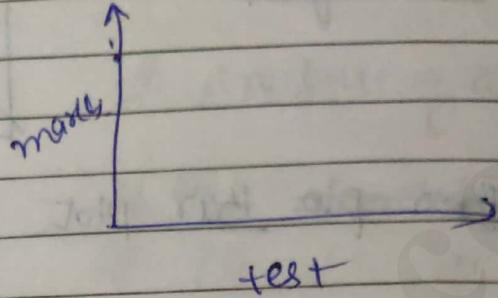
$\text{main} = "title of graph"$



```
> x <- 1:10  
> plot(x)
```

```
> plot(x, y)  
> plot(dataframe) # Graph b/w 1st 2 col by default
```

```
> test <- 1:5  
> marks <- c(23, 21, 20, 17, 19)  
> plot(test, marks)
```



by default vector name
is set as axes label

```
> test  
> marks  
> df = data.frame(test, marks)  
> df  
> plot(df)
```

} plot using
data frames.

always 1st 2 cols

```
> plot(df$c, df$d) # col c & col d are plotted
```

```
> data(mtcars) # Load current env. with  
mtcars dataset
```

~~> head(mtcars)~~

m
> plot(mtcars[, 1], mtcars[, 2]) (1st 1st & 2nd col)
or
plot(mtcars \$ mpg, mtcars \$ cyl)

Page No.		
Date		

If want to we > plot(mpg, cyl)

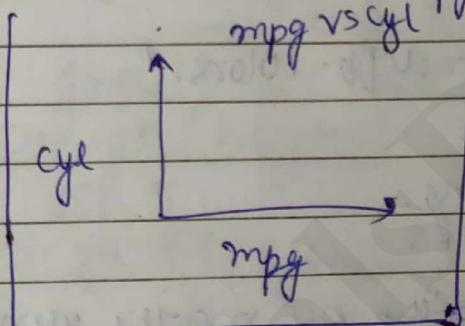
R will look for mpg ~~not~~ but
dataset is not known so error.
but if > attach(mtcars)
thus attaches the mtcars dataset
to the search path.

> attach(mtcars)

> plot(mpg, cyl)

→ adding titles (use)

> plot(mtcars[, 1], mtcars[, 2], main = "mpg vs cyl",
xlab = "mpg", ylab = "cyl")



Note :- > plot(x, y) is same as ~~plot~~ plot(y ~ x)

df <- read.csv("C:\\Users\\ved\\Desktop\\data.csv")

> str(df)

> nrow(df)

> ncol(df)

> dim(df)

>

2) Barcharts

barplot (H, xlab, ylab, main, names.arg, col)

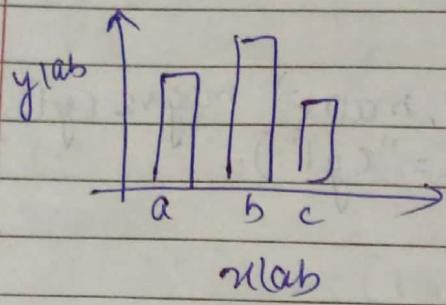
H → vector or matrix.

xlab, ylab → label of x & y - axis.

main → title

names.arg = is a vector of names appearing under each bar

col → is used to give colors to bars in graph.



names.arg = c("a", "b", "c")

col = c("red", "-")

for different colors.

→ color (col) has recycling.

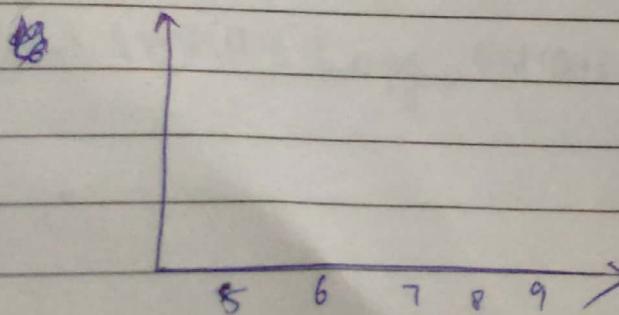
Q4 plot barchart of mean temp by months using airquality.

```
> data(airquality)
```

```
> head(airquality)
```

1
2
3
4
5

	Month	temp	day
1	5	30	1
2	6	27	2
3	7		
4	8		
5	9		
6	5		



tapply → apply functⁿ on complete dataset
tapply (vector, factor to divide, mean)

heights ← tapply (airquality \$ Temp, airquality \$ Month, mean)

vector

factor

functⁿ to
be applied.

> barplot (heights, main = "mean temp by month",
names.arg = c("May", "June", "July", "Aug", "Sep"),
ylab = "Temp (deg. F)")

they shade bars acc. to rank.

shorter → light, taller → darker.

range is [0-1]

gray() is used → to find color value

→ > length(heights)

[1] 5

→ > rank(heights)

[1] 6 7 8 9

[1] 3 4 5 2

(lowest value) ↑ (highest value)

→ > rel.heights ← rank(heights) / length(heights)

> grays ← gray(1 - rel.heights)

color value acc. to rank.

> barplot (heights, col = grays, ...)

TutorialsDuniya.com

Download FREE Computer Science Notes, Programs, Projects, Books PDF for any university student of BCA, MCA, B.Sc, B.Tech CSE, M.Sc, M.Tech at <https://www.tutorialsduniya.com>

- Algorithms Notes
- Artificial Intelligence
- Android Programming
- C & C++ Programming
- Combinatorial Optimization
- Computer Graphics
- Computer Networks
- Computer System Architecture
- DBMS & SQL Notes
- Data Analysis & Visualization
- Data Mining
- Data Science
- Data Structures
- Deep Learning
- Digital Image Processing
- Discrete Mathematics
- Information Security
- Internet Technologies
- Java Programming
- JavaScript & jQuery
- Machine Learning
- Microprocessor
- Operating System
- Operational Research
- PHP Notes
- Python Programming
- R Programming
- Software Engineering
- System Programming
- Theory of Computation
- Unix Network Programming
- Web Design & Development

Please Share these Notes with your Friends as well

facebook

WhatsApp 

twitter 

Telegram 

2018 (Semester)

Q

Write a code in R to plot a graph to depict the relation b/w temp both in $^{\circ}\text{C}$ (on x-axis) and in $^{\circ}\text{F}$ (on y-axis) using the formula $T(F) = T(C) * \frac{9}{5} + 32$:

Label the axis and give title as Celsius vs Fahrenheit graph.

Soln:

generate code to take values for $^{\circ}\text{C}$ & use formula for 'F' or vice versa.

Ques

Store the following data in a dataframe & perform the following.

	A	B
alpha	100	
Beta	120	
gamma	80	
Delta	110	

- (i) Display the contents of dataframe.
- (ii) Draw barplot and name the bars according to the values of col A.

28th March '19

⇒ Plotting a line from x & y points.

> `plot(x, y, type = "l")`
cause bar plot & not scattered.

> `plot(df, type = "l")`

⇒ Boxplot

→ mean

→ Standard deviation $\sqrt{\sum (x - \bar{x})^2}$

→ max, min

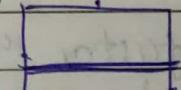
→ outliers × remove

→ quartile.

③ Outliers.

→ max ~~****~~

Q1



median.

Q3

—

→ min.

`boxplot(x, data, notch, varwidth, names, main)`

→ vector or a formula (from where we want to pick data)

data → `dataframe(factor)` (categorical data)

notch → logical value. Set TRUE to draw a notch.

varwidth: logical value. Set as true to draw width of box proportionate to sample space.

names:

main:

(Q) -

std	marks	test
1	30	1
2	20	2
3	25	3
1	20	2
1	30	3

factor .

boxplot(df\$marks, df\$test)

input <- mtcars[, c('mpg', 'cyl')]

new data frame with 1st & 2nd col

boxplot(mpg, cyl) → 2 box plots

* * boxplot(cyl, mpg) → 3 box plots.

If ~~test~~ parameter is factor then
mpg levels = no. of boxplots.

? table(input\$cyl)

0 1 2 3 4 5 6 7 8

11 7 14

11 + 7 + 14 = 32 → no. of rows

freq. of cyl in given

→ Histogram → depict freq. corr. to something.

hist(v, main, nclass, nlim, ylim, breaks, col, border)
v → vector whose freq. is to be plotted

main

breaks → width of each bar.

col → colors for bars

border → used to set border color for each bar
border = c(black, --)

histogram is ~~not~~ never used for more than 1 attribute

→ names(df)

names every col.

→ Boxplot v miaka 'final' w.r.t grade
means final marks & grade bad mein

Section 2.6 → to be covered. (Result of foll. funct'ns with diff data structures.)

→ mean(x)

→ median(x)

→ sd(x)

standard deviation

→ var(x)

→ cor(x, y)

correl

→ cov(x, y)

covariance

Unit 5
Section 6:

Ques: 1) data given \rightarrow write program for
variance, etc.
2) mean, etc used; asking for
O/P.

Ques: $x \leftarrow c(0, 1, 1, 2, 3, 5, 8, 13, 21, 34)$
> mean(x)
> median(x)
> var(x)
> sd(x) # for

i) $\bar{x} = \frac{\sum_{i=1}^n x_i}{n}$ n: length of x

ii) $n \begin{cases} \text{odd} & \frac{n+1}{2} + \frac{n}{2} \\ \text{even} & \frac{n}{2} \end{cases}$

iii) var $\rightarrow \frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})^2$

iv) sd $\rightarrow \sqrt{\text{var}(x)}$

* In case of NA if m. NA not used
 then ans is NA. If used, NA term
 not calculated in total.
 $\text{mean}(n) \rightarrow 8.8$

$\text{median} \rightarrow 4$

$$\text{iii) } \text{var}(x) = \frac{1}{10-1} \sum_{i=1}^{10} (x_i - \bar{x})^2$$

$$= \frac{1}{9} \left[(10-8.8)^2 + (1-8.8)^2 + (1-8.8)^2 + (2-8.8)^2 + (3-8.8)^2 + (5-8.8)^2 + (8-8.8)^2 + (13-8.8)^2 + (21-8.8)^2 + (34-8.8)^2 \right]$$

$$= 121.73.$$

$\text{sd} = 11.03 \text{ km}$

Ques: Consider the vector x defined
 as follows and write the
 o/p of the following
 questions:

$x \leftarrow \text{seq}(1, 10, \text{by}=2)$

~~so~~ $n = 1, 3, 5, 7, 9$

i) find mean n & median x

i) $x[n[6]] \leftarrow 2$

$\rightarrow x[7] \leftarrow 4$

$\rightarrow x[c(8, 9, 10)] \leftarrow 2$

$\rightarrow print(length(x))$

print

Consider print length of x & the final mode of x .

↳ has no predefined function.

iii) find st. dev. & variance of x .

iv) $x[12] \leftarrow NA$

\rightarrow now, print mean(x)

\rightarrow also state the effect of the introduction of the NA value on this vector x .

\rightarrow Rectify the syntax so as to calculate the actual mean of vector x .

Q1^m: i) mean - 5
median - 5

ii) 1, 3, 5, 7, 9, 2, 4, 2, 2, 2.
 $length(x) \rightarrow 10$

preliminary mode of (x)

where, $x = c(1, 3, 5, 7, 9, 2, 4, 2, 2, 2)$

Q) ans $\leftarrow y \left[\text{which} \cdot \text{max}(\text{tabulate}(\text{match}(x, y))) \right]$
 ① $y \leftarrow \text{unique}(x); y: o/p: [1] 12345+9$

unique(x): This func returns a row vector that contains the sorted set of unique values of its argument.

* Some versions of R return a set of unique values w/o sorting it.

match(x, y): This func returns the first occurrence of the first argument in the second arg.

tabulate(): takes the integer valued vector and counts the no. of times each integer occurs in it.

iii) $\text{val} \rightarrow \frac{1}{q} \left[q - \text{which}([] \rightarrow \text{TRUE}) \right]$: returns the index of the logical ~~true~~ object when it is true.

val $\rightarrow 6.67$.

sd $\rightarrow 2.58$

iv) $\text{mean}(x) \rightarrow \text{NA}$

mean can't be cal.

$\text{mean}(x, m. \text{NA} = \text{True})$

1, 3, 5, 7, 9, 2, 4, 2, 2, NA, NA

length $\rightarrow 12$

unique(x) /: match: 123456+666

1 3 5 + 9 2 4

① (2) (3) 24 (1) [(1) 87]

$\text{tabulate}(1, 3, 4, 5, 6, 7, 6, 6, 6)$

O/P: 1 1 1 1 4 1

↓

frequency of each no.

$\max(\text{tabulate}(\dots)) \rightarrow 4 \checkmark$

i.e., $\text{ans} \leftarrow y[\text{which}(y)]$.

v)

CORRELATION: $y \leftarrow \text{rep}(1:n, \text{each}=2)$
& covariance i) find correlation (n, y) and
 covariance (n, y)

vii) remove NA values from x & create
 a vector $x1$ from x with values < 4 .

viii) $y \leftarrow 1:6$

now find $\text{cor}(x1, y1)$ & $\text{cov}(x1, y1)$

$$\text{cov}(n, y) = \frac{\sum_{i=1}^n (n_i - \bar{x})(y_i - \bar{y})}{n-1}$$

$$\text{cor}(x, y) = \frac{\text{cov}(x, y)}{\text{var}(x) \cdot \text{var}(y)}$$

where, x & y have to be vectors of same dimensions.

Sol: \checkmark $x \rightarrow [1, 3, 5, 7, 9, 2, 4, 2, 2, 2, \text{NA}, \text{NA}]$

$y \leftarrow [9, 2, 4, 4, 6, 6, 8, 8, 10, 10, 3, 3, 5, 5, 3, 3, 3, 3, \text{NA}, \text{NA}, \text{NA}]$

i) has mismatch in dimensions.

vii) $\checkmark x \leftarrow x[1:12] < 4$
 $\checkmark n \leftarrow n[n == \text{NA}]$.

$y \leftarrow x[! \text{is.na}(x)]$] removes NA values.
 $y \leftarrow$

[1] 1 3 5 7 9 2 4 2 2 2.

$\checkmark n \leftarrow x[x < 4]$

$\checkmark n$

[1] 1 3 2 2 2

viii) $y \leftarrow (1, 2, 3, 4, 5, 6)$

$$\begin{aligned}& > \text{cov}(x, y) \\& > \text{cov}(x, y_1)\end{aligned}$$

values:

$$\bar{x} = 2, \bar{y} = 3.5$$

$$\text{cov}(x, y) = \frac{(1-2)(1-3.5) + (3-2)(2-3.5)}{5}$$

$$+ \frac{(2-2)(3-3.5)}{5} + (0+0+0)$$

$$= 0.5 + (-0.3)$$

$$= 0.2.$$

$$\text{var}(x) = \frac{1}{5} [(1-2)^2 + (3-2)^2 + 0^2]$$

$$= \frac{1}{5} [18 + 1^2] = \frac{2}{5} = 0.4.$$

$$\text{var}(y) = \frac{1}{5} [(1-3.5)^2 + (2-3.5)^2 + (3-3.5)^2 + (4-3.5)^2 + (5-3.5)^2 + (6-3.5)^2]$$

$$= 3.5$$

$$\text{cor} = \frac{0.2}{\sqrt{0.4 \times 3.5}} = 0.1493. \quad \underline{\underline{= 0.169}}$$

Dataframes:

1) mean and standard deviation of a df works by column
and these functions will only work for the numerical cols of a dataframe.

$\text{df} \rightarrow \text{df}$

small medium large

2	5	9
3	6	10
4	7	8

1) mean(df)

small medium large

mean value

m.v.

m.v.

2) std(df)

small medium large

3) median function doesn't apply directly for a df; needs to be applied individually.

- 1) median (df & small)
median (df & medium)
median (df & large)

or
using `apply()` or `lapply()`
to apply through function
→ med by mode also.

- 2) variance & covariance remains
a covariance matrix for the
dataname.

`> var(df)`
`> cov(df)`

O/P:

	small	medium	large
small	small unit small	one unit medium units	large unit
medium			
large		many	few

→ covariance matrix

- 4) `cor(df)` - same.

Pg 31 - Q. 6: apply these functions.

TutorialsDuniya.com

Download FREE Computer Science Notes, Programs, Projects, Books PDF for any university student of BCA, MCA, B.Sc, B.Tech CSE, M.Sc, M.Tech at <https://www.tutorialsduniya.com>

- Algorithms Notes
- Artificial Intelligence
- Android Programming
- C & C++ Programming
- Combinatorial Optimization
- Computer Graphics
- Computer Networks
- Computer System Architecture
- DBMS & SQL Notes
- Data Analysis & Visualization
- Data Mining
- Data Science
- Data Structures
- Deep Learning
- Digital Image Processing
- Discrete Mathematics
- Information Security
- Internet Technologies
- Java Programming
- JavaScript & jQuery
- Machine Learning
- Microprocessor
- Operating System
- Operational Research
- PHP Notes
- Python Programming
- R Programming
- Software Engineering
- System Programming
- Theory of Computation
- Unix Network Programming
- Web Design & Development

Please Share these Notes with your Friends as well

facebook

WhatsApp 

twitter 

Telegram 