

## **Advance R**

### **DataFrame in R :**

A data frame is a collection of 2 vectors means 2-dimensional array-like structure or a tabular form in which a column contains values of one variable, and rows contains one set of values from each column. A data frame is a type of the list in which each component has equal length.

A data frame is used to store data table and the vectors which are present in the form of a list in a data frame, are of equal length.

In a simple way, it is a list of equal length vectors. A matrix can contain one type of data, but a data frame can contain different data types such as numeric, character, factor, etc.

DataFrame is the most important data structure in the field of Data Analytics and Data Science.

Data can be collected from various sources like Database, Cloud, CSV, etc, but is ultimately stored in a data frame.

There are following characteristics of a data frame.

- The columns name should be non-empty.
- The rows name should be unique.
- The data which is stored in a data frame can be a factor, numeric, or character type.
- Each column contains the same number of data items

## Creating data frames

Data frame can be created using `data.frame()` function

```
v1 <- 1:5
# Letter is an inbuilt list
letters

#output
'a' 'b' 'c' 'd' 'e' 'f' 'g' 'h' 'i' 'j' 'k' 'l' 'm' 'n' 'o' 'p' 'q' 'r' 's' 't' 'u' 'v' 'w'
'x' 'y' 'z'
```

```
v2 <- letters[1:5]
df <- data.frame(v1,v2)
df
```

output:

```
v1 v2
1  a
2  b
3  c
4  d
5  e
```

## Creating dataframe with column name

```
c1 <- c('Tendulkar','Kohli','Dohni','Bumrah','Chahal')
c2 <- c(10000,7100,5800,890,870)
c3 <- c(11,0,0,370,420)
cricket <- data.frame(players=c1,runs=c2,wickets=c3)
cricket
```

```
output :
Players    runs  wickets
Tendulkar 10000    11
Kohli      7100     0
Dohni      5800     0
```

Bumrah	890	370
Chahal	870	420

## Inbuilt data frames

View inbuilt data frames using data() function

```
data()  
state.x77
```

## Understanding data

Use head() & tail() function to see first 6 and last 6 rows respectively

```
head(state.x77)
```

```
tail(state.x77)
```

Use summary() and str() function to get basic description of data like mean, median, quartiles etc.

```
summary(state.x77)
```

Output:

```
Population Income Illiteracy Life Exp  
Min. : 365 Min. :3098 Min. :0.500 Min. :67.96  
1st Qu.: 1080 1st Qu.:3993 1st Qu.:0.625 1st Qu.:70.12  
Median : 2838 Median :4519 Median :0.950 Median :70.67  
Mean : 4246 Mean :4436 Mean :1.170 Mean :70.88  
3rd Qu.: 4968 3rd Qu.:4814 3rd Qu.:1.575 3rd Qu.:71.89  
Max. :21198 Max. :6315 Max. :2.800 Max. :73.60  
Murder HS Grad Frost Area  
Min. : 1.400 Min. :37.80 Min. : 0.00 Min. : 1049  
1st Qu.: 4.350 1st Qu.:48.05 1st Qu.: 66.25 1st Qu.: 36985  
Median : 6.850 Median :53.25 Median :114.50 Median : 54277  
Mean : 7.378 Mean :53.11 Mean :104.46 Mean : 70736  
3rd Qu.:10.675 3rd Qu.:59.15 3rd Qu.:139.75 3rd Qu.: 81163  
Max. :15.100 Max. :67.30 Max. :188.00 Max. :566432
```

```
str(state.x77)
```

Output :

```
num [1:50, 1:8] 3615 365 2212 2110 21198 ...  
- attr(*, "dimnames")=List of 2  
..$ : chr [1:50] "Alabama" "Alaska" "Arizona" "Arkansas" ...  
..$ : chr [1:8] "Population" "Income" "Illiteracy" "Life Exp" ...
```

```
nrow(state.x77)
```

```
#output  
50
```

```
ncol(state.x77)
```

```
#output  
8
```

```
rownames(state.x77)
```

```
colnames(state.x77)
```

```
#output  
'Population' 'Income' 'Illiteracy' 'Life Exp' 'Murder' 'HS Grad' 'Frost' 'Area'
```

## Indexing and slicing

### 1) Selecting cells

*Creating dataframe with column name*

```
c1 <- c('Tendulkar','Kohli','Dohni','Bumrah','Chahal')  
c2 <- c(10000,7100,5800,890,870)  
c3 <- c(11,0,0,370,420)  
cricket <- data.frame(players=c1,runs=c2,wickets=c3)  
cricket
```

```
Cricket[1,2]
```

```
#output  
10000
```

```
Cricket[1:3,1:2]
```

```
#output  
players    runs  
Tendulkar 10000  
Kohli      7100  
Dohni      5800
```

```
cricket[c(1,4),c(1,3)]
```

```
#output  
  players  wickets  
1 Tendulkar   11  
4 Bumrah    370
```

```
cricket[1:3,'wickets']
```

```
#output  
11 0 0
```

## 2) Selecting rows

```
cricket[1,]
```

```
#output  
players    runs  wickets  
Tendulkar 10000   11
```

```
Cricket[1:3,]
```

```
#output  
players    runs  wickets  
Tendulkar 10000   11  
Kohli      7100    0
```

```
Dohni      5800  0
```

```
cricket[c(1,3),]
```

```
#output
  players    runs  wickets
1 Tendulkar 10000      11
3 Dohni      5800       0
```

### 3) Selecting columns

```
Cricket[,3]
```

```
#output
11 0 0 370 420
```

```
Cricket[,1:2]
```

```
#output
 players    runs
Tendulkar  10000
Kohli       7100
Dohni       5800
Bumrah      890
Chahal      870
```

```
cricket[,c(1,3)]
```

```
#output
 players  wickets
Tendulkar  11
Kohli      0
Dohni      0
Bumrah    370
Chahal    420
```

```
Cricket[, 'runs']
```

```
#output
10000 7100 5800 890 870
```

```
cricket[,c('players','wickets')]
```

```
#output
players    wickets
Tendulkar   11
Kohli       0
Dohni       0
Bumrah     370
Chahal     420
```

```
cricket[['wickets']]
```

```
#output
11 0 0 370 420
```

```
cricket$players
```

```
#output
Tendulkar Kohli Dohni Bumrah Chahal
```

### Conditional selection with subset() function

```
subset(cricket,subset=runs>5000)
```

```
#output
Players    runs    wickets
Tendulkar 10000     11
Kohli      7100      0
Dohni      5800      0
```

```
subset(cricket,subset=wickets>300)
```

```
#output
  players  runs  wickets
4  Bumrah   890   370
5  Chahal   870   420
```

## Ordering dataframe

Ordering is done with order() function

```
order(cricket['runs'])
```

```
#output  
5 4 3 2 1
```

```
runs.order <- order(cricket['runs'])  
cricket[runs.order,]
```

```
#output  
  players  runs  wickets  
5 Chahal   870   420  
4 Bumrah   890   370  
3 Dohni   5800    0  
2 Kohli   7100    0  
1 Tendulkar 10000   11
```

```
runs.order <- order(-cricket['runs'])  
cricket[runs.order,]
```

```
#output  
players  runs  wickets  
Tendulkar 10000  11  
Kohli      7100   0  
Dohni      5800   0  
Bumrah     890  370  
Chahal     870  420
```

## Renaming Names

```
mat <- matrix(1:20,nrow=5)  
print(mat)
```

```
df <- data.frame(mat)  
Df
```

```
#output  
X1 X2 X3 X4  
1  6 11 16
```



```
2  7 12 17
3  8 13 18
4  9 14 19
5 10 15 20
```

```
# renaming single column
colnames(df)[1] <- 'index'
df
```

```
#output
index  X2  X3  X4
1      6   11 16
2      7   12 17
3      8   13 18
4      9   14 19
5     10   15 20
```

```
# renaming multiple column
colnames(df) <- c('A','B','C','D')
df
```

```
#output
A  B  C  D
1  6 11 16
2  7 12 17
3  8 13 18
4  9 14 19
5 10 15 20
```

### *Adding new rows and columns*

```
# Adding row
df2 <- data.frame(A=20,B=34,C=67,D=56)
df <- rbind(df,df2)
df
```

```
#output
A  B  C  D
1  6 11 16
2  7 12 17
3  8 13 18
```

```
4  9 14 19
5 10 15 20
20 34 67 56
```

```
# adding columns using replicate function
df$E <- rep(NA,nrow(df))
df
```

```
#output
A  B  C  D  E
1   6 11 16 NA
2   7 12 17 NA
3   8 13 18 NA
4   9 14 19 NA
5  10 15 20 NA
20 34 67 56 NA
```

```
df$F <- 10:15
df
```

```
#output
A  B  C  D  E  F
1   6 11 16 NA 10
2   7 12 17 NA 11
3   8 13 18 NA 12
4   9 14 19 NA 13
5  10 15 20 NA 14
20 34 67 56 NA 15
```

```
#copying another column
df$G = df$E
df
```

```
#output
A  B  C  D  E  F  G
1   6 11 16 NA 10 NA
2   7 12 17 NA 11 NA
3   8 13 18 NA 12 NA
4   9 14 19 NA 13 NA
5  10 15 20 NA 14 NA
20 34 67 56 NA 15 NA
```

```
# cbind function
V <- c(10,20,30,40,50,60)
df <- cbind(df,V)
df
#output
A  B  C  D  E  F  G  V
1   6 11 16 NA 10 NA 10
2   7 12 17 NA 11 NA 20
3   8 13 18 NA 12 NA 30
4   9 14 19 NA 13 NA 40
5  10 15 20 NA 14 NA 50
20 34 67 56 NA 15 NA 60
```

## Handling missing values

check presence of missing values with any() and is.na() function

```
# Check in specific column
any(is.na(df$A))
```

```
#output
FALSE
```

```
any(is.na(df$E))
```

```
#output
TRUE
```

```
# Check entire dataframe
any(is.na(df))
```

```
#output
TRUE
```

```
#Replacing missing values
df$G[is.na(df$G)] <- 0
df
```

```
#output
A  B  C  D  E  F  G  V
```

1	6	11	16	NA	10	0	10
2	7	12	17	NA	11	0	20
3	8	13	18	NA	12	0	30
4	9	14	19	NA	13	0	40
5	10	15	20	NA	14	0	50
20	34	67	56	NA	15	0	60

```
df[is.na(df)] <- -1
df
```

```
#output
A  B  C  D  E  F  G  V
1   6 11 16 -1 10 0 10
2   7 12 17 -1 11 0 20
3   8 13 18 -1 12 0 30
4   9 14 19 -1 13 0 40
5  10 15 20 -1 14 0 50
20 34 67 56 -1 15 0 60
```

Note : Missing data must be replaced with mean, median or mode.

## List in R

In R, lists are the second type of vector. Lists are the objects of R which contain elements of different types such as number, vectors, string and another list inside it. It can also contain a function or a matrix as its elements. A list is a data structure which has components of mixed data types. We can say, a list is a generic vector which contains other objects.

Example

```
vec <- c(3,4,5,6)
char_vec <- c("shubham","nishka","gunjan","sumit")
logic_vec <- c(TRUE,FALSE,FALSE,TRUE)
out_list <- list(vec,char_vec,logic_vec)
```

out\_list

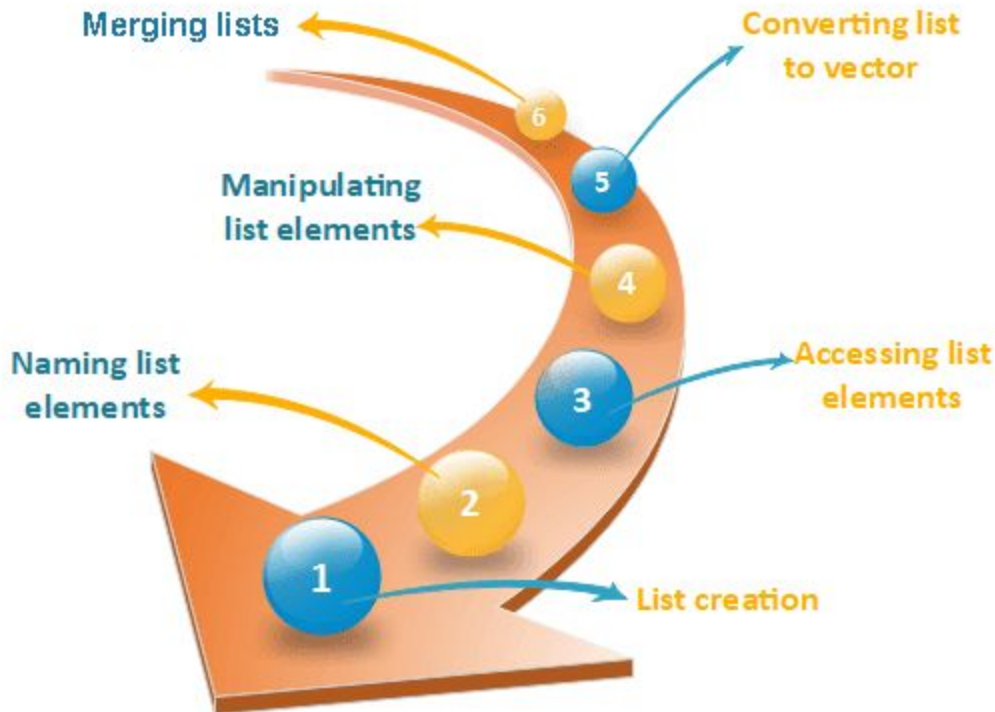
Output:

```
[[1]]
[1] 3 4 5 6

[[2]]
[1] "shubham" "nishka" "gunjan" "sumit"

[[3]]
[1] TRUE FALSE FALSE TRUE
```

# Lists in R programming



## Lists creation

The process of creating a list is the same as a vector. In R, the vector is created with the help of `c()` function. Like `c()` function, there is another function, i.e., `list()` which is used to create a list in R. A list avoid the drawback of the vector which is data type. We can add the elements in the list of different data types.

Syntax

```
list()
```

Example 1: Creating list with same data type

```
list_1<-list(1,2,3)
```

```
list_2<-list("Shubham","Arpita","Vaishali")
```

```
list_3<-list(c(1,2,3))
```

```
list_4<-list(TRUE,FALSE,TRUE)
```

```
list_1
```

```
list_2
```

```
list_3
```

```
list_4
```

Output:

```
[[1]]
```

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

```
[[1]]
[1] "Shubham"
[[2]]
[1] "Arpita"
[[3]]
[1] "Vaishali"
```

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

```
[[1]]
[1] TRUE
[[2]]
[1] FALSE
[[3]]
[1] TRUE
```

Example 2: Creating the list with different data type

```
list_data<-list("Shubham","Arpita",c(1,2,3,4,5),TRUE,FALSE,22.5,12L)
print(list_data)
```

In the above example, the list function will create a list with character, logical, numeric, and vector element. It will give the following output

Output:

```
[[1]]
[1] "Shubham"
[[2]]
[1] "Arpita"
[[3]]
[1] 1 2 3 4 5
[[4]]
[1] TRUE
[[5]]
[1] FALSE
[[6]]
[1] 22.5
[[7]]
[1] 12
```

Giving a name to list elements

R provides a very easy way for accessing elements, i.e., by giving the name to each element of a list. By assigning names to the elements, we can access the element easily. There are only three steps to print the list data corresponding to the name:

Creating a list.

Assign a name to the list elements with the help of `names()` function.

Print the list data.

Let see an example to understand how we can give the names to the list elements.

Example

# Creating a list containing a vector, a matrix and a list.

```
list_data <- list(c("Shubham","Nishka","Gunjan"), matrix(c(40,80,60,70,90,80),
nrow = 2),
  list("BCA","MCA","B.tech"))
```

# Giving names to the elements in the list.

```
names(list_data) <- c("Students", "Marks", "Course")
```

# Show the list.

```
print(list_data)
```

Output:

\$Students

```
[1] "Shubham" "Nishka" "Gunjan"
```

\$Marks

```
  [,1] [,2] [,3]
[1,] 40 60 90
[2,] 80 70 80
```

\$Course

```
$Course[[1]]
```

```
[1] "BCA"
```

```
$Course[[2]]
```

```
[1] "MCA"
```

```
$Course[[3]]
```

```
[1] "B. tech."
```

Accessing List Elements

R provides two ways through which we can access the elements of a list. First one is the indexing method performed in the same way as a vector. In the second one, we can access the elements of a list with the help of names. It will be possible only

with the named list.; we cannot access the elements of a list using names if the list is normal.



Let see an example of both methods to understand how they are used in the list to access elements.

Example 1:

**Accessing elements using index**

# Creating a list containing a vector, a matrix and a list.

```
list_data <- list(c("Shubham","Arpita","Nishka"), matrix(c(40,80,60,70,90,80), nrow = 2),  
  list("BCA","MCA","B.tech"))
```

# Accessing the first element of the list.

```
print(list_data[1])
```

# Accessing the third element. The third element is also a list, so all its elements will be printed.

```
print(list_data[3])
```

Output:

```
[[1]]
```

```
[1] "Shubham" "Arpita" "Nishka"
```

```
[[1]]
```

```
[[1]][[1]]
```

```
[1] "BCA"
```

```
[[1]][[2]]
```



```
[1] "MCA"
```

```
[[1]][[3]]  
[1] "B.tech"
```

Example 2:

### **Accessing elements using names**

# Creating a list containing a vector, a matrix and a list.

```
list_data <- list(c("Shubham","Arpita","Nishka"), matrix(c(40,80,60,70,90,80), nrow  
= 2),list("BCA","MCA","B.tech"))
```

# Giving names to the elements in the list.

```
names(list_data) <- c("Student", "Marks", "Course")
```

# Accessing the first element of the list.

```
print(list_data["Student"])
```

```
print(list_data$Marks)
```

```
print(list_data)
```

Output:

```
$Student
```

```
[1] "Shubham" "Arpita" "Nishka"
```

```
      [,1] [,2] [,3]  
[1,]  40  60  90  
[2,]  80  70  80
```

```
$Student
```

```
[1] "Shubham" "Arpita" "Nishka"
```

```
$Marks
```

```
      [,1] [,2] [,3]  
[1,]  40  60  90  
[2,]  80  70  80
```

```
$Course
```

```
$Course[[1]]
```

```
[1] "BCA"
```

```
$Course[[2]]
```

```
[1] "MCA"
```

```
$Course[[3]]
```

```
[1] "B. tech."
```

## Manipulation of list elements

R allows us to add, delete, or update elements in the list. We can update an element of a list from anywhere, but elements can add or delete only at the end of the list. To remove an element from a specified index, we will assign it a null value. We can update the element of a list by overriding it from the new value. Let see an example to understand how we can add, delete, or update the elements in the list.

Example

```
# Creating a list containing a vector, a matrix and a list.
```

```
list_data <- list(c("Shubham","Arpita","Nishka"), matrix(c(40,80,60,70,90,80), nrow = 2),  
  list("BCA","MCA","B.tech"))
```

```
# Giving names to the elements in the list.
```

```
names(list_data) <- c("Student", "Marks", "Course")
```

```
# Adding element at the end of the list.
```

```
list_data[4] <- "Moradabad"
```

```
print(list_data[4])
```

```
# Removing the last element.
```

```
list_data[4] <- NULL
```

```
# Printing the 4th Element.
```

```
print(list_data[4])
```

```
# Updating the 3rd Element.
```

```
list_data[3] <- "Masters of computer applications"
```

```
print(list_data[3])
```

Output:

```
[[1]]
```

```
[1] "Moradabad"
```

```
$<NA>
```

```
NULL
```

```
$Course
```

```
[1] "Masters of computer applications"
```

## Converting list to vector

There is a drawback with the list, i.e., we cannot perform all the arithmetic operations on list elements. To remove this, drawback R provides `unlist()` function. This function converts the list into vectors. In some cases, it is required to convert a list into a vector so that we can use the elements of the vector for further manipulation.

The `unlist()` function takes the list as a parameter and change into a vector. Let see an example to understand how to `unlist()` function is used in R.

Example

```
# Creating lists.
```

```
list1 <- list(10:20)
```

```

print(list1)
list2 <- list(5:14)
print(list2)
# Converting the lists to vectors.
v1 <- unlist(list1)
v2 <- unlist(list2)
print(v1)
print(v2)
#adding the vectors
result <- v1+v2
print(result)
Output:
[[1]]
[1] 1 2 3 4 5

```

```

[[1]]
[1] 10 11 12 13 14

```

```

[1] 1 2 3 4 5
[1] 10 11 12 13 14
[1] 11 13 15 17 19

```

## Merging Lists

R allows us to merge one or more lists into one list. Merging is done with the help of the `list()` function also. To merge the lists, we have to pass all the lists into `list` function as a parameter, and it returns a list which contains all the elements which are present in the lists. Let see an example to understand how the merging process is done.

Example

```

# Creating two lists.
Even_list <- list(2,4,6,8,10)
Odd_list <- list(1,3,5,7,9)
# Merging the two lists.
merged.list <- list(Even_list,Odd_list)
# Printing the merged list.
print(merged.list)

```

Output:

```

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

```

```

[[1]][[2]]
[1] 4

```

```

[[1]][[3]]

```

```
[1] 6
```

```
[[1]][[4]]  
[1] 8
```

```
[[1]][[5]]  
[1] 10
```

```
[[2]]  
[[2]][[1]]  
[1] 1
```

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

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

```
[[2]][[4]]  
[1] 7
```

```
[[2]][[5]]  
[1] 9
```

## **R Built-in Features**

`seq()` - creates a sequence

```
seq(1,10)  
  
#output  
1 2 3 4 5 6 7 8 9 10
```

```
# by argument for stepsize  
seq(1,10,by=2)  
  
#output  
1 3 5 7 9
```

```
seq(1,10,2)  
  
#output
```

```
1 3 5 7 9
```

sort() to sort a vector

```
v1 <- c(10,2,11,4,6,5,12,7)
sort(v1)
```

```
#output
2 4 5 6 7 10 11 12
```

```
sort(v1, decreasing=T)
```

```
#output
12 11 10 7 6 5 4 2
```

rev() to reverse elements of an object

```
rev(v1)
#output
7 12 5 6 4 11 2 10
```

str() describes structure of an object

```
str(v1)

#output
num [1:8] 10 2 11 4 6 5 12 7
```

```
mat <- matrix(1:15,nrow=3)
mat
#output
1 4 7 10 13
2 5 8 11 14
3 6 9 12 15
```

```
str(mat)

#output
int [1:3, 1:5] 1 2 3 4 5 6 7 8 9 10 ...
```

append() merges objects together

```
append(v1,90)
```

```
#output  
10 2 11 4 6 5 12 7 90
```

```
v2 <- c(1,2,3,4)  
append(v1,v2)
```

```
#output  
10 2 11 4 6 5 12 7 1 2 3 4
```

sample() gives a random value from a sequence

```
sample(1:20,1)
```

```
#output  
17
```

```
sample(1:20,2)
```

```
#output  
1 17
```

```
sample(v1,1)
```

```
#output  
10
```

is.\* checks class of an object

```
is.vector(v1)
```

```
#output  
TRUE
```

```
is.list(v1)
```

```
#output  
FALSE
```

as.\* converts object type

```
as.list(v1)
```

```
as.matrix(v1)
```

*Regular expression*

grepl() returns logical output

```
text <- "I am data scientist"  
grepl("data",text)
```

```
#output  
TRUE
```

```
grepl("outlier",text)
```

```
#output  
FALSE
```

grep() returns index

```
v <- c(10,20,30)  
v
```

```
#output  
10 20 30
```

```
grep(20,v)
```

```
#output  
2
```

```
grep(40,v)
```

```
#output  
# No output
```

## Date Time

`Sys.Date()` get system's date

```
Sys.Date()  
  
#output  
2019-09-06
```

`as.Date()` convert string to date object

```
as.Date('1997-02-19')  
#output  
1997-02-19
```

## Formatting

Code	Value
%d	Day of the month
%m	Month
%b	Abbreviated month
%B	Full month
%y	2 digit year
%Y	4 digit year

```
as.Date('28-November-1992',format="%d-%B-%Y")  
  
#output  
1992-11-28
```

```
as.Date('12-Jan-86',format="%d-%b-%y")  
  
#output  
1986-01-12
```

`Sys.time()` to get time

```
Sys.time()  
  
#output  
"2019-09-06 16:49:38 IST"
```

`as.POSIXct` to convert string to date and time



```
as.POSIXct("08:30:03",format="%H:%M:%S")
```

```
#output  
"2019-09-06 08:30:03 IST"
```

```
as.POSIXct("28-November-1992 09:25:04",format="%d-%B-%Y %H:%M:%S")
```

```
#output  
"1992-11-28 09:25:04 IST"
```

## Math functions

### **abs()**

```
In [29]: abs(-10)
```

```
10
```

### **round()**

```
In [30]: round(10.23)
```

```
10
```

```
In [31]: round(10.232343,2)
```

```
10.23
```

### **sqrt()**

```
In [33]: sqrt(144)
```

```
12
```

### **exp()**

```
In [35]: exp(1)
```

```
2.71828182845905
```

## Logarithm

In [43]: `log(10)`

2.30258509299405

In [44]: `log10(12)`

1.07918124604762

## Trigonometry

In [47]: `sin(90)`

0.893996663600558

In [48]: `cos(90)`

-0.44807361612917

In [49]:

`tan(90)`

## apply function in R

`apply()` is a inbuilt function which takes matrix or Data Frame as an input and provides the output in list, array or vector. `apply()` Function is used to avoid explicit uses of constructs of loop. It is the most basic of all collections can be used over a matrix. The simplest example is to sum a matrix over all the columns.

This function takes 3 arguments:

`apply(X, MARGIN, FUN)`

Here:

-x: an array or matrix

-MARGIN: take a value or range between 1 and 2 to define where to apply the function:

-MARGIN=1: the manipulation is performed on rows

-MARGIN=2: the manipulation is performed on columns

-MARGIN=c(1,2) the manipulation is performed on rows and columns

-FUN: tells which function to apply. Built functions like mean, median, sum, min, max and even user-defined functions can be applied

The simplest example is to sum a matrix over all the columns. The code `apply(m1, 2, sum)` will apply the sum function to the matrix 5x6 and return the sum of each column accessible in the dataset.

```
m1 <- matrix(C<-(1:10),nrow=5, ncol=6)
```

```
m1
```

```
a_m1 <- apply(m1, 2, sum)
```

```
a_m1
```

Output:

```
> m1
      [,1] [,2] [,3] [,4] [,5] [,6]
[1,]    1    6    1    6    1    6
[2,]    2    7    2    7    2    7
[3,]    3    8    3    8    3    8
[4,]    4    9    4    9    4    9
[5,]    5   10    5   10    5   10
> a_m1 <- apply(m1, 2, sum)
> a_m1
[1] 15 40 15 40 15 40
>
```

*Sum of column*







