Mean

It is calculated by taking the sum of the values and dividing with the number of values in a data series.

The function **mean()** is used to calculate this in R.

Syntax

```
The basic syntax for calculating mean in R is - mean(x, trim = 0, na.rm = FALSE, ...)
```

Following is the description of the parameters used -

- x is the input vector.
- trim is used to drop some observations from both end of the sorted vector.
- na.rm is used to remove the missing values from the input vector.

Example

```
# Create a vector.
x <- c(12,7,3,4.2,18,2,54,-21,8,-5)

# Find Mean.
result.mean <- mean(x)
print(result.mean)
When we execute the above code, it produces the following result -
[1] 8.22</pre>
```

Applying Trim Option

When trim parameter is supplied, the values in the vector get sorted and then the required numbers of observations are dropped from calculating the mean.

When trim = 0.3, 3 values from each end will be dropped from the calculations to find mean.

In this case the sorted vector is (-21, -5, 2, 3, 4.2, 7, 8, 12, 18, 54) and the values removed from the vector for calculating mean are (-21, -5, 2) from left and (12, 18, 54) from right.

```
# Create a vector.
x <- c(12,7,3,4.2,18,2,54,-21,8,-5)
# Find Mean.
result.mean <- mean(x,trim = 0.3)
print(result.mean)</pre>
```

When we execute the above code, it produces the following result – [1] 5.55

Applying NA Option

If there are missing values, then the mean function returns NA.

To drop the missing values from the calculation use na.rm = TRUE. which means remove the NA values.

```
# Create a vector.
x <- c(12,7,3,4.2,18,2,54,-21,8,-5,NA)

# Find mean.
result.mean <- mean(x)
print(result.mean)

# Find mean dropping NA values.
result.mean <- mean(x,na.rm = TRUE)
print(result.mean)

When we execute the above code, it produces the following result -
[1] NA
[1] 8.22</pre>
```

Median

The middle most value in a data series is called the median. The **median()** function is used in R to calculate this value.

Syntax

The basic syntax for calculating median in R is - median(x, na.rm = FALSE)

Following is the description of the parameters used -

- x is the input vector.
- na.rm is used to remove the missing values from the input vector.

Example

```
# Create the vector.
x <- c(12,7,3,4.2,18,2,54,-21,8,-5)

# Find the median.
median.result <- median(x)
print(median.result)
When we execute the above code, it produces the following result --</pre>
```

Mode

The mode is the value that has highest number of occurrences in a set of data.

Unike mean and median, mode can have both numeric and character data.

R does not have a standard in-built function to calculate mode. So we create a user function to calculate mode of a data set in R.

This function takes the vector as input and gives the mode value as output.

Example

```
# Create the function.
getmode <- function(v) {</pre>
   uniqv <- unique(v)</pre>
   uniqv[which.max(tabulate(match(v, uniqv)))]
}
# Create the vector with numbers.
v \leftarrow c(2,1,2,3,1,2,3,4,1,5,5,3,2,3)
# Calculate the mode using the user function.
result <- getmode(v)</pre>
print(result)
# Create the vector with characters.
charv <- c("o","it","the","it","it")</pre>
# Calculate the mode using the user function.
result <- getmode(charv)</pre>
print(result)
When we execute the above code, it produces the following result -
[1] "it"
```