### **CORELATION ANALYSIS:**

- \* Corelation is a statistical measuare that extends to which two variable are linearly in relation
- $^{st}$  It refers to the process of establishing the relationship between two variables

Types of Corelations:

```
Values = -1, 0, -1
```

- $\ ^{*}$  -1 indicates strong negative corelation which means x increases, y decreases
- \* 0 no association between two variables
- \* +1 indicates strong positive corelation which means x increases, y increases

Methods of Corelations:

- \* Pearson (measures the linear dependancy of two variables)
- \* Spearman (compute the corelation between the rank of x and y variables)
- $^{st}$  Kendal (measures the correspondence between the x & y varibales)

# Correlation Coefficient Formula

$$r = \frac{n(\Sigma xy) - (\Sigma x)(\Sigma y)}{\sqrt{[n\Sigma x^2 - (\Sigma x)^2][n\Sigma y^2 - (\Sigma y)^2]}}$$

#### **Pearson Method:**

```
In [1]:
```

```
x = c(4,8,12,16)
y = c(5,10,15,20)
```

if method is not specified by default pearson method is used

### In [2]:

```
cor.test(x,y, method="pearson")
```

```
Pearson's product-moment correlation
```

```
data: x and y
t = Inf, df = 2, p-value < 2.2e-16
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
    1 1
sample estimates:
cor
    1</pre>
```

```
In [6]:
a = cor.test(x,y)
if (a$estimate ==1)
    print("Strong Positive")
} else if(a$estimate==-1){
   print("Strong Negative")
} else{
    print("No Relation")
[1] "Strong Positive"
In [7]:
a = c(4,8,12,26)
b = c(5,10,15,30)
In [8]:
v = cor.test(a,b)
if (v$estimate ==1)
    print("Strong Positive")
} else if(v$estimate==-1){
   print("Strong Negative")
} else{
    print("No Relation")
[1] "No Relation"
In [12]:
r = c(4,8,12,0)
t = c(5,0,5,30)
In [16]:
o = cor.test(r,t)
if (o$estimate ==1)
   print("Strong Positive")
} else if(o$estimate<1){</pre>
   print("Strong Negative")
} else{
   print("No Relation")
[1] "Strong Negative"
```

#### In [ ]:

## **Statistical Summary:**

```
Exploring Data
          -> Summarization
          -> Visualisation
          -> Normalization
Data Summarization:
      * Mean -> Sum of X / number of values
      * Median -> Center number
      * Mode \rightarrow The number which is repeated multiple times
```

Replacing missing values:

If there is a null value in the dataset then go for data summarisation to fill the null value.

```
Measure the Central Dispertion:
-> Standard Deviation
-> Variance
-> Range

Measure of Central Dispertion:
The measure of central dispertion express the scattering of the datapoints in terms of distance such as range or in terms of deviation from the central value such as variance and standard deviation.
```

# **Measure of Central Tendency:**

```
In [41]:
a = c(5,52,53,64,65,86,86,92)
mean = mean(a)
mode <- function(a) {</pre>
   uniav <- unique(a)
   uniqv[which.max(tabulate(match(a, uniqv)))]
median = median(a)
result <- mode(a)</pre>
print(paste("Mean: ",mean))
print(paste("Mode: ",result))
print(paste("Median: ",median))
[1] "Mean: 62.875"
[1] "Mode: 86"
[1] "Median: 64.5"
In [43]:
# creating a data frame:
df = data.frame(mean, result, median)
df
```

```
        mean
        result
        median

        62.875
        86
        64.5
```

# Measure of Dispersion or Variability:

```
In [35]:
A = c(5,52,53,64,65,86,86,92)
a = sort(A)
# Range:
range= max(a) - min(a)
# Variance:
variance= var(a)
# Standard Deviation:
std = sd(a)
# Dispersion:
dispersion = mean(a) - sd(a)
print(paste("Range: ", range))
print(paste("Variance: ", variance))
print(paste("Standard Deviation: ", std))
print(paste("Dispersion: ", dispersion))
[1] "Range: 87"
[1] "Variance: 784.125"
[1] "Standard Deviation: 28.0022320538917"
[1] "Dispersion: 34.8727679461083"
```

```
In [24]:
```

```
# quartile:
b= quantile(a)
b

0%
5
25%
52.75
50%
64.5
75%
86
100%
92
```

#### In [37]:

```
df1 = data.frame(range, variance, dispersion, Q1=b[2], Q2=b[3], Q3=b[4], row.names="Data Set 1")
df1
```

	range	variance	dispersion	Q1	Q2	Q3	
Data Set 1	87	784.125	34.87277	52.75	64.5	86	

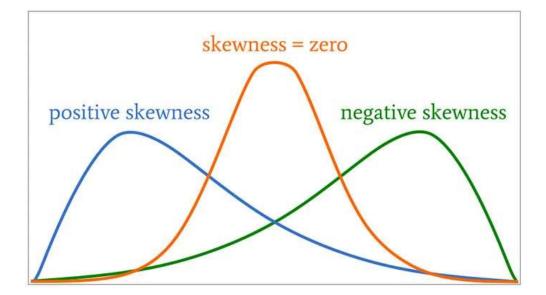
# **Symmetrical Distribution:**

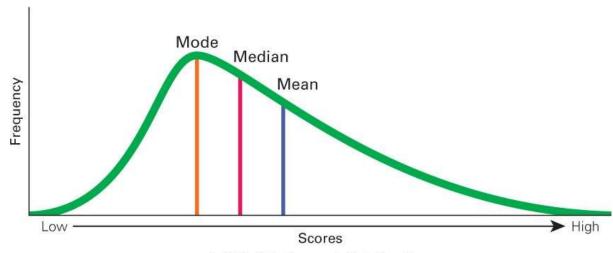
- \* A frequency distribution is said to be symmetrical if the frequencies are equally distributed on both the sides of central values.
- \* A symmetrical distibution may be either bell-shaped or U-shaped.
- \* Mean = Median = Mode

# **Skewed Distribution:**

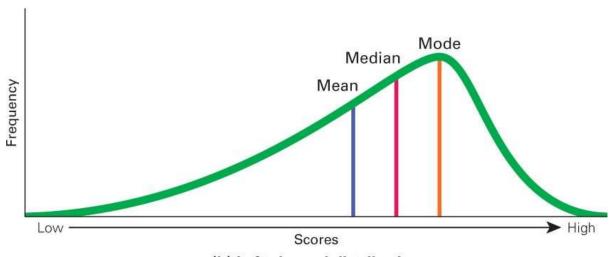
Types of Skewness:

- \* Positive Skewness (Mean exceeds Mode and Median)
- \* Negative Skewness (Mode exceeds Mean and Median)
- \* No Skewness (Mean Median Mode are equal)





# (a) Right-skewed distribution



# (b) Left-skewed distribution

## **Pearson's Coefficient of Skewness:**

\* This method is most frequently used for measuring skewness.

Formula:

Formula: (When Mode cannot be determined we can use this formula)

Mean - (3\*Median - 2\*Mean)

Pearsons Coefficient = ------
Standard Deviation

#### In [46]:

```
install.packages("e1071")
```

```
There is a binary version available but the source version is later:
    binary source needs_compilation
e1071 1.7-6 1.7-12 TRUE

Binaries will be installed

Warning message:
"package 'e1071' is in use and will not be installed"
```

```
In [47]:
```

```
library("e1071")
```

### In [48]:

```
a = c(41, 52, 53, 64, 65, 86, 86, 92)
skewness(a)
```

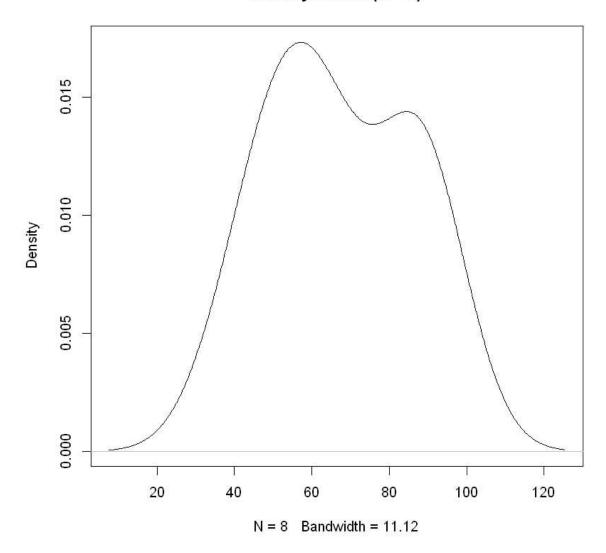
0.0542062901835262

### In [49]:

```
skewness(a)
plot(density(a))
```

0.0542062901835262

# density.default(x = a)



### In [54]:

data()

## In [60]:

data(mtcars)

## In [61]:

## mtcars\$mpg

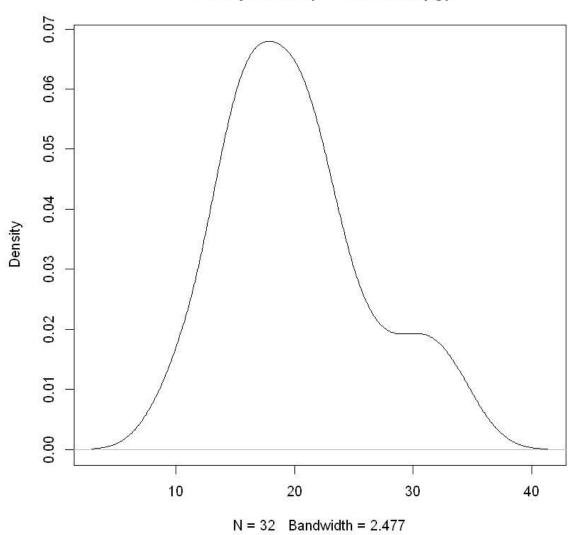
21 21 22.8 21.4 18.7 18.1 14.3 24.4 22.8 19.2 17.8 16.4 17.3 15.2 10.4 10.4 14.7 32.4 30.4 33.9 21.5 15.5 15.2 13.3 19.2 27.3 26 30.4 15.8 19.7 15 21.4

### In [65]:

skewness(mtcars\$mpg)
plot(density(mtcars\$mpg))

0.610655017573288

# density.default(x = mtcars\$mpg)



	Measure o	of Tendency	y			Measure of Dispersi	ion or Varia	ability				
	Mean	Median	Mode		Range	Standard Deviation	Variable	Q1	Q2	Q3	Dispression	
Dataset 1	67.4	64.5	86		51	18.7	350.8	52.5	64.5	86	48.7	
Dataset 1	62.8	64.5	86		87	28	784.1	52.5	64.5	86	34.8	

# In [ ]: