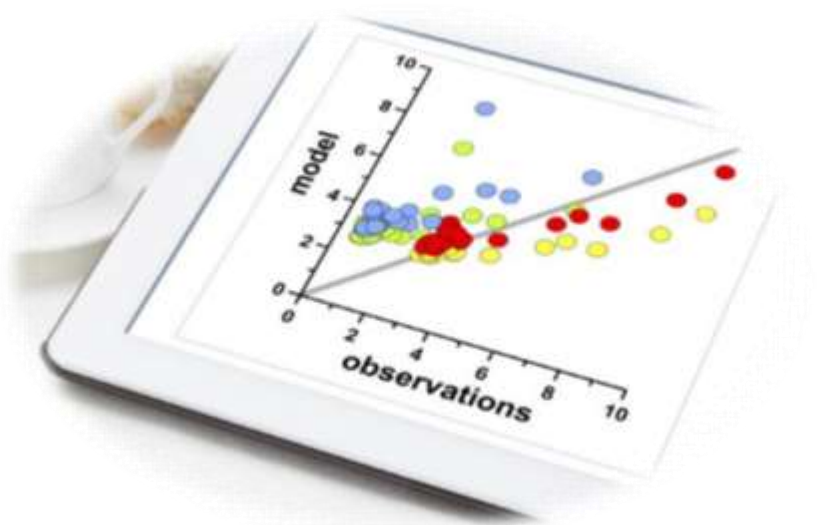
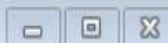


CORRELATION



Correlation is a statistic that measures the degree to which two variables move in relation to each other. Correlation shows the strength of a relationship between two variables and is expressed numerically by the correlation coefficient. The correlation coefficient's values range between -1.0 and 1.0.



```
> x=c(41, 19, 23, 40, 55, 57, 33)
> y=c(94, 60, 74, 71, 82, 76, 61)
> cor.test(x,y,method="pearson")#which gives the karlpearson correlation coeffici$
```

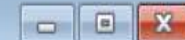
Pearson's product-moment correlation

```
data: x and y
t = 1.434, df = 5, p-value = 0.211
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
 -0.3592689  0.9192115
sample estimates:
      cor
0.5398442
```

```
> x=c(41, 19, 23, 40, 55, 57, 33)
> y=c(94, 60, 74, 71, 82, 76, 61)
> cor.test(x,y,method="spearman")#which gives the spearman rank correlation corre$
```

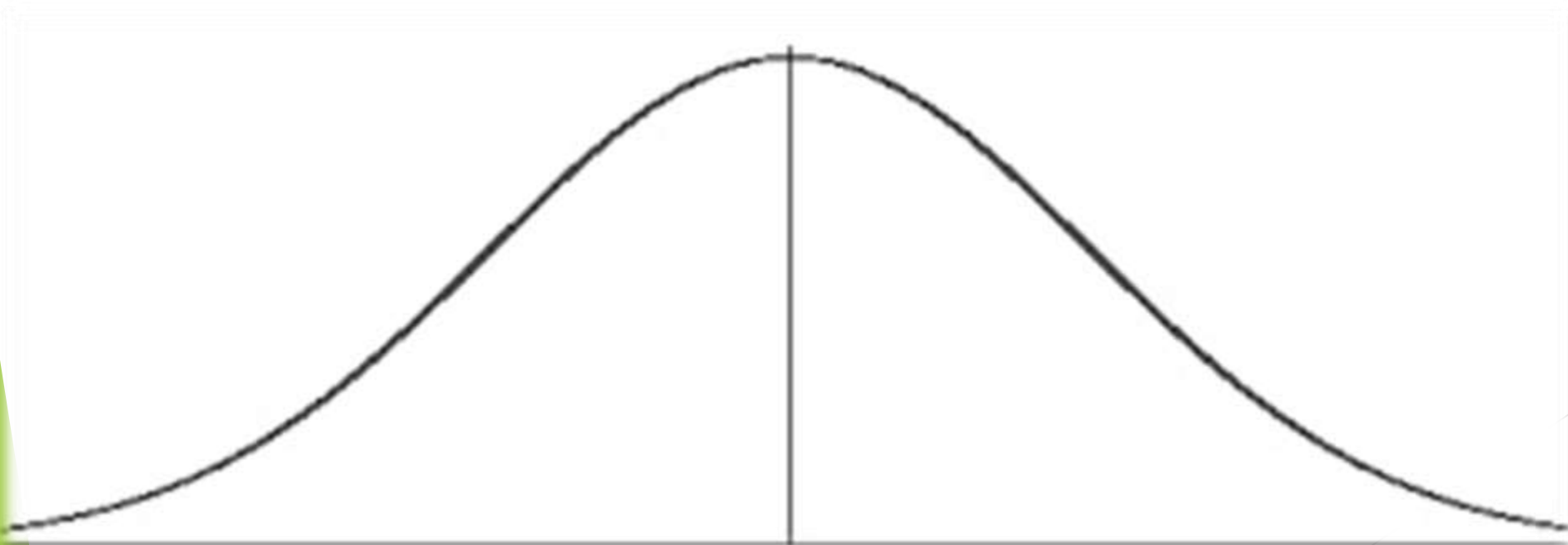
Spearman's rank correlation rho

```
data: x and y
S = 14, p-value = 0.06627
alternative hypothesis: true rho is not equal to 0
sample estimates:
 rho
0.75
```



```
x=c(41, 19, 23, 40, 55, 57, 33)
y=c(94, 60, 74, 71, 82, 76, 61)
cor.test(x,y,method="pearson")#which gives the karlpearson correlation coefficient
x=c(41, 19, 23, 40, 55, 57, 33)
y=c(94, 60, 74, 71, 82, 76, 61)
cor.test(x,y,method="spearman")#which gives the spearman rank correlation correlation coefficient
```

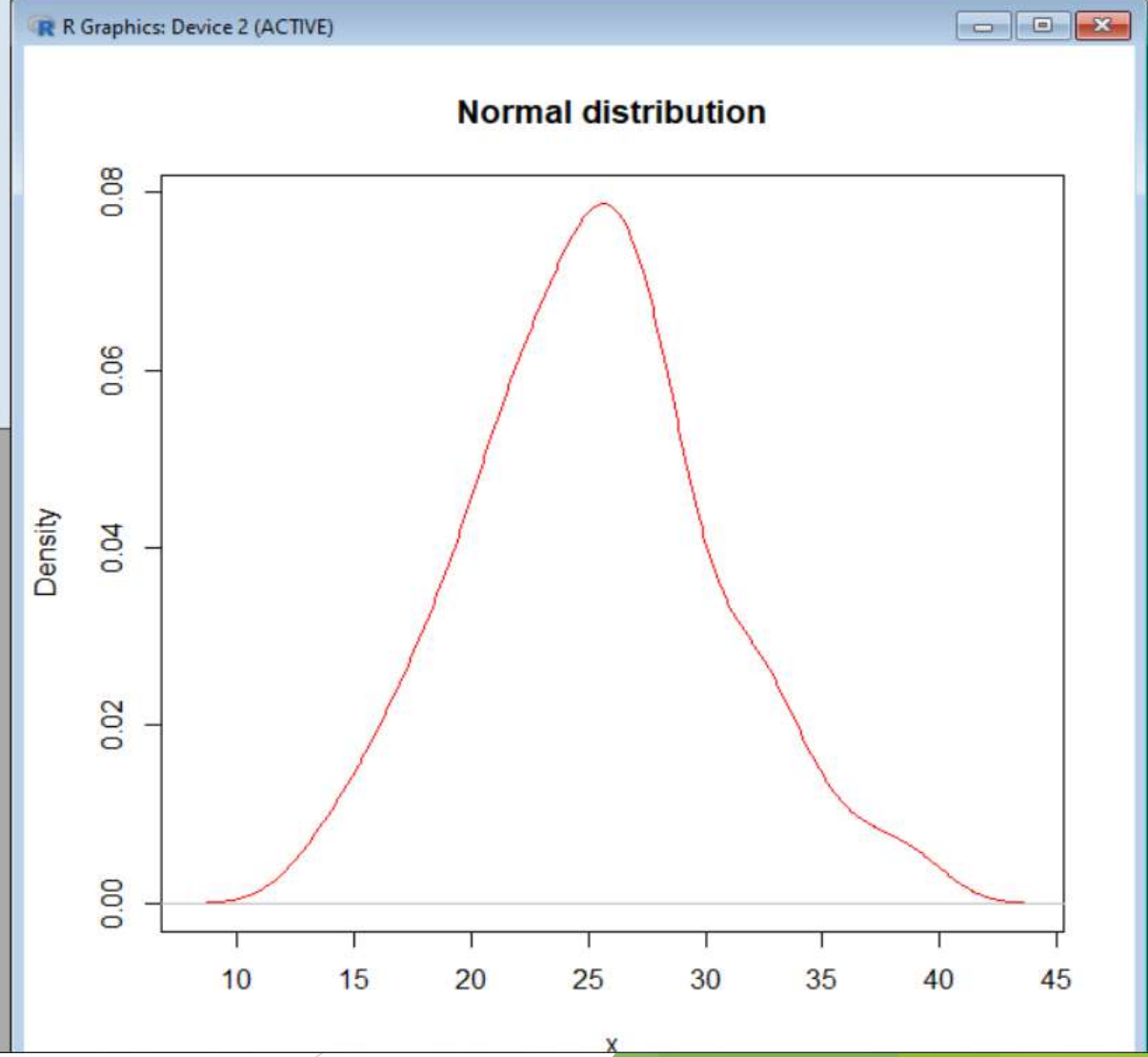
NORMAL DISTRIBUTION





```
R Console
> x=rnorm(100,25,5)
> x
 [1] 20.81912 27.04226 21.83944 28.74034 22.49530 17.64128 26.12654 19.80359
 [9] 32.23006 22.79251 13.15147 24.23004 26.75500 27.42123 23.32643 19.93035
[17] 23.06309 25.42483 23.51121 19.81128 30.73146 25.96814 17.26478 21.39583
[25] 20.95771 30.01278 22.46398 21.57285 27.28210 19.42773 19.34907 27.87450
[33] 25.08025 27.68925 27.93027 22.49029 17.71134 16.58586 19.32949 21.07117
[41] 24.83860 26.65687 17.05034 28.81423 28.87760 33.07609 36.33339 24.87211
[49] 38.95004 24.07807 34.19551 24.95435 15.43695 14.63349 29.12675 30.63966
[57] 25.37408 33.01092 31.58248 25.05012 27.49078 36.15386 27.89803 32.78314
[65] 29.34952 23.37823 27.28611 27.61882 27.83632 26.67948 22.31832 23.53783
[73] 25.47758 38.54592 26.14368 19.86300 28.57875 25.27992 32.59681 14.90846
[81] 23.48397 32.12400 30.18208 22.66758 18.24439 22.64950 21.10928 34.42324
[89] 25.64916 26.04689 27.77377 24.68229 25.17532 21.85420 31.45554 24.91702
[97] 24.68167 17.79900 26.23592 20.73721
> plot(density(x),xlab="x",main="Normal distribution",col="red")
> |
```

```
Untitled - R Editor
x=rnorm(100,25,5)
x
plot(density(x),xlab="x",main="Normal distribution",col="red")
```



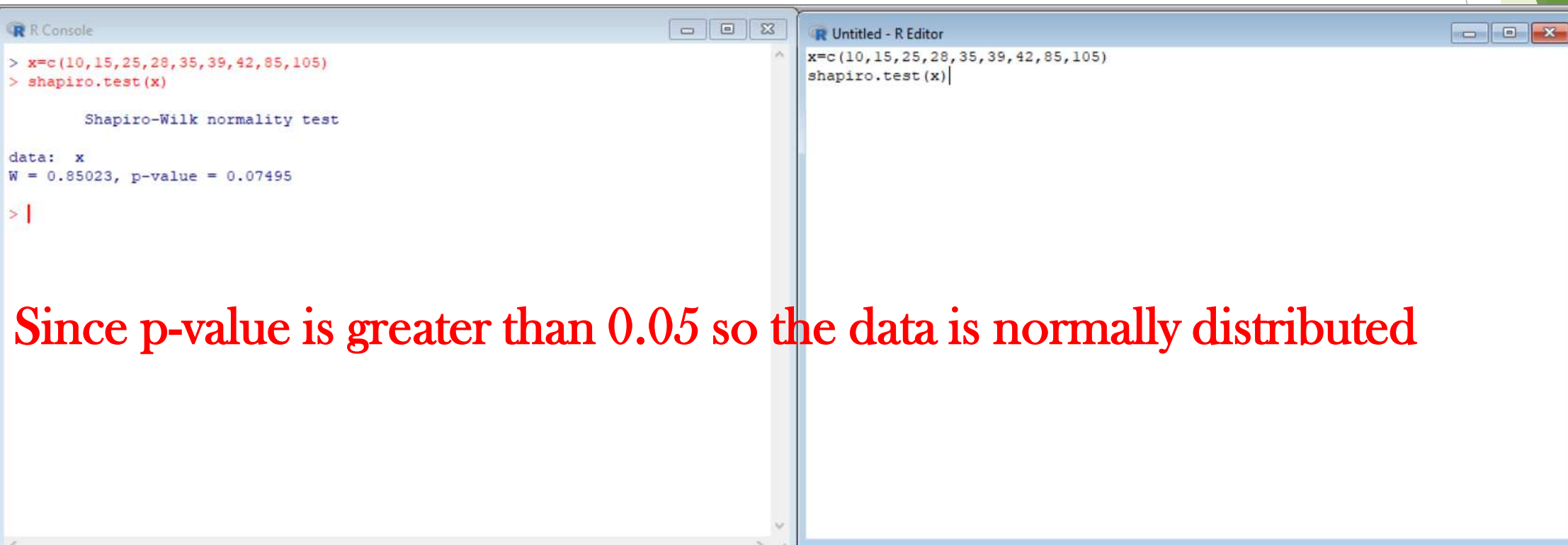
Tests for Normality are:

- ▶ **Shapiro-Wilk's test :** The Shapiro-Wilk test is a way to tell if a random sample comes from a normal distribution. The test gives you a **W** value; small values indicate your sample is *not* normally distributed (you can reject the null hypothesis that your population is normally distributed if your values are under a certain threshold)
- ▶ **Normal probability Plot :** A normal probability plot is a graphical representation of the data. A normal probability plot is used to check if the given data set is normally distributed or not. It is used to compare a data set with the normal distribution. If a given data set is normally distributed then it will reside in a shape like a straight line.

Shapiro-Wilk's test

H0: The given data set is Normally distributed

H1: The given data set is not normally distributed



The image shows two windows from an R environment. The left window, titled 'R Console', displays the following output:

```
> x=c(10,15,25,28,35,39,42,85,105)
> shapiro.test(x)

      Shapiro-Wilk normality test

data:  x
W = 0.85023, p-value = 0.07495
> |
```

The right window, titled 'Untitled - R Editor', shows the code used to perform the test:

```
x=c(10,15,25,28,35,39,42,85,105)
shapiro.test(x)
```

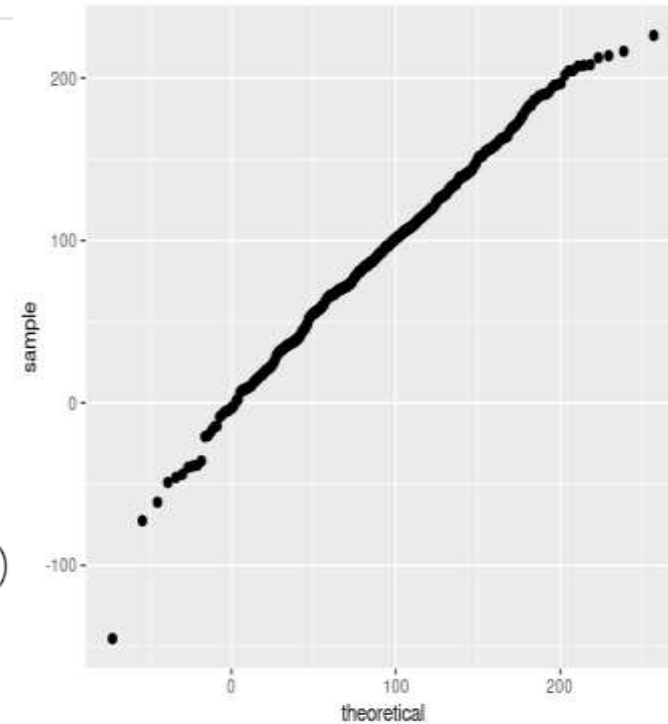
Since p-value is greater than 0.05 so the data is normally distributed

Normal probability Plot

```
# importing libraries
library(ggplot2)
library(qqplotr)

# creating random data
random_values = rnorm(500, mean = 90, sd = 50)

# plotting data without line and labels
ggplot(mapping = aes(sample = random_values)) + stat_qq_point(size = 2)
```

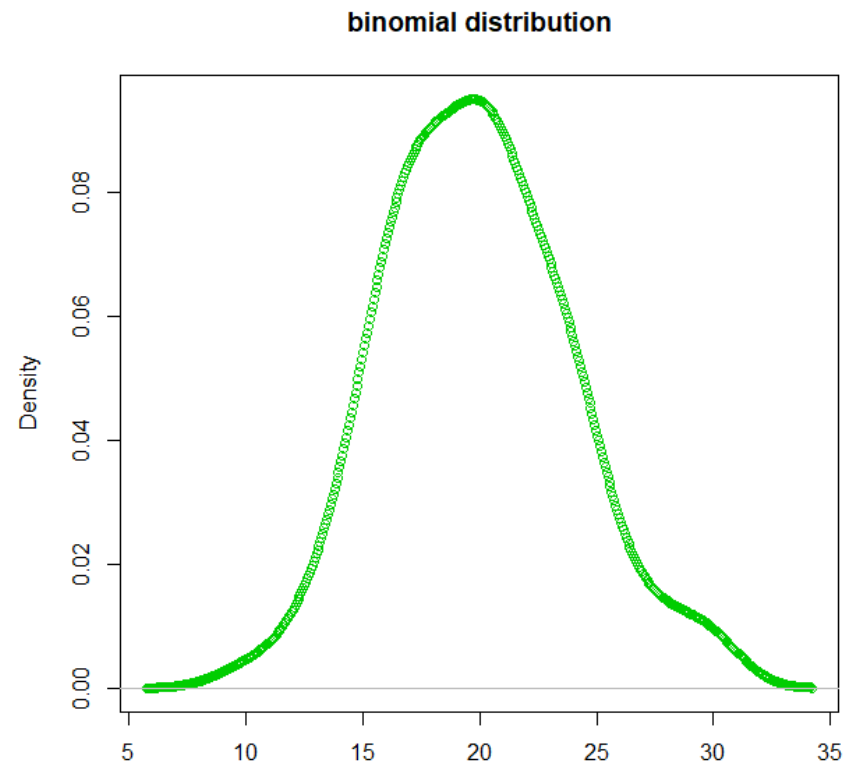


Plotting different types of distributions

➤ Binomial Distribution

```
R Console
> x=rbinom(100,50,.4)
> plot(density(x),type="b",xlab="x",main="binomial distribution",col="green3")
>

Untitled - R Editor
x=rbinom(100,50,.4)
plot(density(x),type="b",xlab="x",main="binomial distribution",col="green3")
```



- **Poisson Distribution : `rpois (n, lambda)`**
- **Exponential Distribution : `rexp(n,rate)`**
- **Normal Distribution : `rnorm(n,mean,sd)`**