

# **Packages in R**

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# PACKAGES

- ✓ **A set of R functions**
  - ✓ **A set of compiled code**
  - ✓ **A Set of sample data**
- 
- ✓ **Library directory in R environment**
  - ✓ **A few packages are installed by default and can be called directly**
  - ✓ **A few packages are installed but should be loaded explicitly**

- ✓ **CRAN: Official Repos**
- ✓ **Biconductor: Topic-specific Rep**
- ✓ **GitHub: Open source project repo**

## library()

### ✓ Lists all the R packages installed

Packages in library 'C:/Users/mpanc/AppData/Local/R/win-library/4.3':

```
cli                Helpers for Developing Command Line Interfaces
colorspace         A Toolbox for Manipulating and Assessing Colors and Palettes
distributions3     Probability Distributions as S3 Objects
ellipsis           Tools for Working with ...
fansl              ANSI Control Sequence Aware String Functions
farver             High Performance Colour Space Manipulation
ggplot2           Create Elegant Data Visualisations Using the Grammar of Graphics
glue              Interpreted String Literals
gtable            Arrange 'Grobs' in Tables
isoband           Generate Isolines and Isobands from Regularly Spaced Elevation Grids
labeling          Axis Labeling
lestat            A Package for Learning Statistics
lifecycle         Manage the Life Cycle of your Package Functions
magrittr          A Forward-Pipe Operator for R
munsell           Utilities for Using Munsell Colours
pillar           Coloured Formatting for Columns
pkgconfig         Private Configuration for 'R' Packages
R6               Encapsulated Classes with Reference Semantics
RColorBrewer     ColorBrewer Palettes
rlang            Functions for Base Types and Core R and 'Tidyverse' Features
scales           Scale Functions for Visualization
tibble          Simple Data Frames
utf8            Unicode Text Processing
vctrs           Vector Helpers
VGAM            Vector Generalized Linear and Additive Models
viridisLite     Colorblind-Friendly Color Maps (Lite Version)
withr           Run Code 'With' Temporarily Modified Global State
```

Packages in library 'C:/Program Files/R/R-4.3.3/library':

```
base              The R Base Package
boot             Bootstrap Functions (Originally by Angelo Canty for S)
class            Functions for Classification
cluster          "Finding Groups in Data": Cluster Analysis Extended Rousseeuw et al.
codetools        Code Analysis Tools for R
compiler         The R Compiler Package
datasets         The R Datasets Package
foreign          Read Data Stored by 'Minitab', 'S', 'SAS', 'SPSS', 'Stata', 'Systat',
                'Weka', 'dBase', ...
graphics         The R Graphics Package
grDevices        The R Graphics Devices and Support for Colours and Fonts
grid             The Grid Graphics Package
KernSmooth      Functions for Kernel Smoothing Supporting Wand & Jones (1995)
lattice          Trellis Graphics for R
MASS            Support Functions and Datasets for Venables and Ripley's MASS
Matrix          Sparse and Dense Matrix Classes and Methods
methods         Formal Methods and Classes
mgcv            Mixed GAM Computation Vehicle with Automatic Smoothness Estimation
nlme            Linear and Nonlinear Mixed Effects Models
nnet            Feed-Forward Neural Networks and Multinomial Log-Linear Models
parallel        Support for Parallel Computation in R
rpart           Recursive Partitioning and Regression Trees
spatial         Functions for Kriging and Point Pattern Analysis
splines         Regression Spline Functions and Classes
stats           The R Stats Package
stats4          Statistical Functions using S4 Classes
survival        Survival Analysis
tcltk           Tcl/Tk Interface
tools           Tools for Package Development
translations    The R Translations Package
utils           The R Utils Package
```

```
install.packages ("Package Name")  
install.packages ("VGAM")  
install.packages (c ("VGAM", "distributions3"))  
installed.packages ()  
update.packages ()
```

```
#you can load using library function  
library(graphics)  
Library(graphics,ggplot2,caret)  
#you can load using require function  
require(graphics)  
require(ggplot2)  
require(caret)
```

`library()` : Load a package and refers to a place where package is available

`packages()` : Collection functions bundled. You can share your functions with others using package



# BUILT-IN FUNCTIONS

# STATISTICS AND PROBABILITY

# *Built-in Functions-Normal Distribution*

- ✓ **R has various statistical probability functions to perform the statistical task**
- ✓ **You can find normal density, normal quartile, etc**
- ✓ **Density, distribution function, quantile function and random generation for the normal distribution with mean equal to mean and standard deviation equal to sd.**

$$f(x) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

# Built-in Functions-Normal Distribution

```
normal(expectation = 0, lambda, P = 1)
```

Function	Description
Expectation	The expectation of the distribution
<b>lambda</b>	The natural logarithm of the standard deviation of the distribution.
<b>P</b>	Precision of the distribution, inverse of the variance.

```
require(lestat)
dist <- normal(3, log(0.7))
variance(dist)
dist <- normal(5, log(0.49)/2)
variance(dist)
dist <- normal(7, P = 2)
variance(dist)
```

# *Built-in Functions-Normal Distribution*

```
Normal(mu = 0, sigma = 1)
```

Function	Description
Normal	It creates a normal distribution
<b>Mu</b>	Mean
<b>Sigma</b>	Standard deviation

# Built-in Functions-Normal Distribution

```
dnorm(x, mean = 0, sd = 1, log = FALSE)
pnorm(q, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
qnorm(p, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
rnorm(n, mean = 0, sd = 1)
```

Function	Description
dnorm	Density
pnorm	Distribution Function
qnorm	Quantile Function
rnorm	Random deviates

# Built-in Functions-Normal Distribution

```
dnorm(x, mean = 0, sd = 1, log = FALSE)
pnorm(q, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
qnorm(p, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
rnorm(n, mean = 0, sd = 1)
```

Argument	Description
x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations. If $\text{length}(n) > 1$ , the length is taken to be the number required.
mean	vector of means.
sd	vector of standard deviations.
log, log.p	logical; if TRUE, probabilities p are given as $\log(p)$ .
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ Otherwise $P[X > x]$

# Built-in Functions-Normal Distribution

```
require(graphics)
```

```
dnorm(0) == 1/sqrt(2*pi)
```

```
dnorm(1) == exp(-1/2)/sqrt(2*pi)
```

```
dnorm(1) == 1/sqrt(2*pi*exp(1))
```

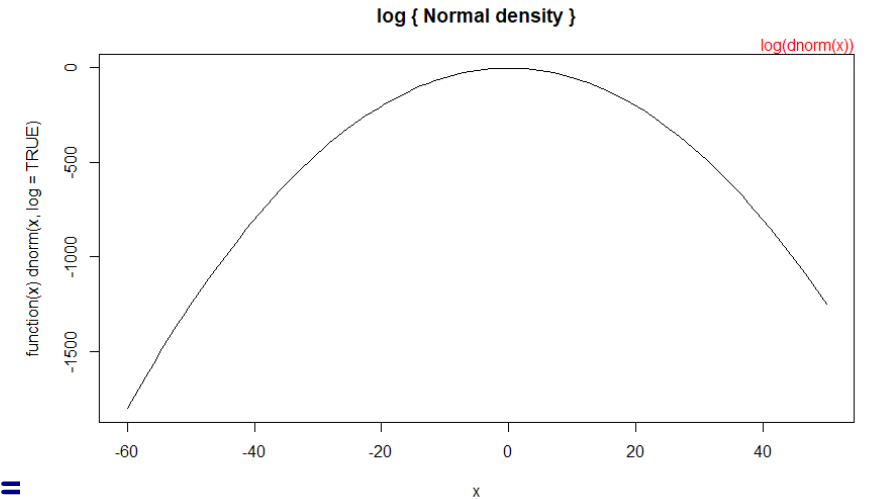
```
par(mfrow = c(2,1))
```

```
plot(function(x) dnorm(x, log = TRUE), -60, 50,  
      main = "log { Normal density }")
```

```
curve(log(dnorm(x)), add = TRUE, col = "red", lwd =
```

```
mtext("dnorm(x, log=TRUE)", adj = 0)
```

```
mtext("log(dnorm(x))", col = "red", adj = 1)
```



# *Built-in Functions-Bivariate Normal Distribution*

```
dbinorm(x1, x2, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0, log = FALSE)
pbinorm(q1, q2, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0)
rbinorm(n, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0)
pnorm2(x1, x2, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0)
```

Function	Description
dbinorm	Density
pbinorm	Distribution Function
rbinorm	Random Deviates (n by 2 matrix)



# Built-in Functions-Bivariate Normal Distribution

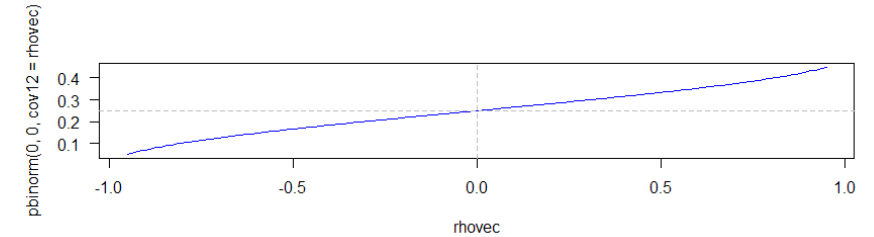
```
dbinorm(x1, x2, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0, log = FALSE)
pbnorm(q1, q2, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0)
rbinorm(n, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0)
pnorm2(x1, x2, mean1 = 0, mean2 = 0, var1 = 1, var2 = 1, cov12 = 0)
```

Argument	Description
x1,x2, q1,q2	vector of quantiles.
n	number of observations. If length(n) > 1, the length is taken to be the number required.
mean1,mean2,var1,var2	vector of means, variances
Cov12	Covariance
Log	Logical. If log = TRUE then the logarithm of the density is returned.

# Built-in Functions-Bivariate Normal Distribution

```
install.packages('VGAM')
require(VGAM)
require(graphics)
yvec <- c(-5, -1.96, 0, 1.96, 5)
ymat <- expand.grid(yvec, yvec)
cbind(ymat, pbinorm(ymat[, 1], ymat[, 2]))

rhovec <- seq(-0.95, 0.95, by = 0.01)
plot(rhovec, pbinorm(0, 0, cov12 = rhovec),
     type = "l", col = "blue", las = 1)
abline(v = 0, h = 0.25, col = "gray", lty = "dashed")
```



# Built-in Functions-Binormal Distribution

Maximum likelihood estimation of the five parameters of a bivariate normal distribution.

```
binormal(lmean1 = "identitylink", lmean2 = "identitylink",  
        lsd1   = "loglink",      lsd2   = "loglink",  
        lrho   = "rhobitlink",  
        imean1 = NULL,           imean2 = NULL,  
        isd1   = NULL,           isd2   = NULL,  
        irho   = NULL,           imethod = 1,  
        eq.mean = FALSE,        eq.sd   = FALSE,  
        zero   = c("sd", "rho"), rho.arg = NA)
```

Function	Description
lmean1,lmean2	Link functions applied to the means,
lsd1,lsd2	Link functions applied to the standard deviations,
Lrho	Link functions applied to the standard rho,

# *Built-in Functions-Bivariate Poisson Distribution*

```
dpois(x, lambda, log = FALSE)
ppois(q, lambda, lower.tail = TRUE, log.p = FALSE)
qpois(p, lambda, lower.tail = TRUE, log.p = FALSE)
rpois(n, lambda)
```

Function	Description
dpois	Density
ppois	Distribution Function
rpois	Random Deviates
qpois	Quantile Function

$$p(x) = \frac{\lambda^x e^{-\lambda}}{x!}$$

# Built-in Functions - Poisson Distribution

```
dpois(x, lambda, log = FALSE)
ppois(q, lambda, lower.tail = TRUE, log.p = FALSE)
qpois(p, lambda, lower.tail = TRUE, log.p = FALSE)
rpois(n, lambda)
```

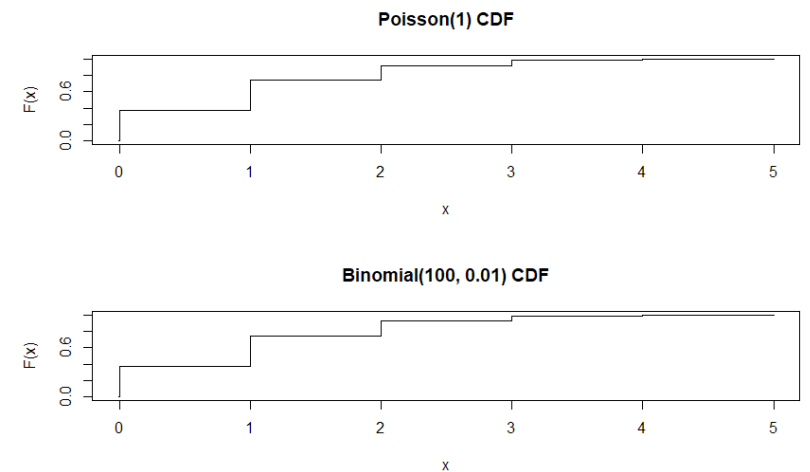
Argument	Description
x, q	vector of quantiles, where x is non-negative integer
p	vector of probabilities.
n	Number of random values to return
Lambda	vector of means (non-negative).
sd	vector of standard deviations.
log, log.p	logical; if TRUE, probabilities p are given as log(p).
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ Otherwise $P[X > x]$

# Built-in Functions - Poisson Distribution

```
require(graphics)

-log(dpois(0:7, lambda = 1) * gamma(1+ 0:7)) # == 1
Ni <- rpois(50, lambda = 4); table(factor(Ni, 0:max(Ni)))
1 - ppois(10*(15:25), lambda = 100) # becomes 0 (cancellation)
ppois(10*(15:25), lambda = 100, lower.tail = FALSE) # no cancellation

par(mfrow = c(2, 1))
x <- seq(-0.01, 5, 0.01)
plot(x, ppois(x, 1), type = "s", ylab = "F(x)", main = "Poisson(1) CDF")
plot(x, pbinom(x, 100, 0.01), type = "s", ylab = "F(x)",
      main = "Binomial(100, 0.01) CDF")
```



# Built-in Functions-Uniform Distribution

```
dunif(x, min = 0, max = 1, log = FALSE)
punif(q, min = 0, max = 1, lower.tail = TRUE, log.p = FALSE)
qunif(p, min = 0, max = 1, lower.tail = TRUE, log.p = FALSE)
runif(n, min = 0, max = 1)
```

Function	Description
dunif	Density
punif	Distribution Function
runif	Random Deviates
qunif	Quantile Function

$$f(x) = \frac{1}{max - min}$$

# Built-in Functions - Poisson Distribution

```
dunif(x, min = 0, max = 1, log = FALSE)
punif(q, min = 0, max = 1, lower.tail = TRUE, log.p = FALSE)
qunif(p, min = 0, max = 1, lower.tail = TRUE, log.p = FALSE)
runif(n, min = 0, max = 1)
```

Argument	Description
x, q	vector of quantiles
p	vector of probabilities.
n	number of observations. If $\text{length}(n) > 1$ , the length is taken to be the number required.
min,max	lower and upper limits of the distribution. Must be finite.
log, log.p	logical; if TRUE, probabilities p are given as $\log(p)$ .
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ Otherwise $P[X > x]$

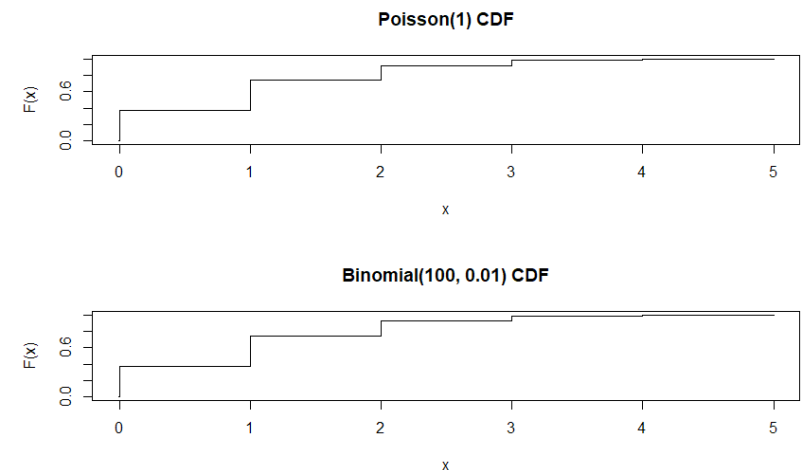


# Built-in Functions - Poisson Distribution

```
require(graphics)

-log(dpois(0:7, lambda = 1) * gamma(1+ 0:7)) # == 1
Ni <- rpois(50, lambda = 4); table(factor(Ni, 0:max(Ni)))
1 - ppois(10*(15:25), lambda = 100) # becomes 0 (cancellation)
ppois(10*(15:25), lambda = 100, lower.tail = FALSE) # no cancellation

par(mfrow = c(2, 1))
x <- seq(-0.01, 5, 0.01)
plot(x, ppois(x, 1), type = "s", ylab = "F(x)", main = "Poisson(1) CDF")
plot(x, pbinom(x, 100, 0.01), type = "s", ylab = "F(x)",
      main = "Binomial(100, 0.01) CDF")
```



# CHARTS

➤ **Values in data vector as height of the bars**

## ✓ **Default Syntax**

```
barplot(height, width = 1, space = NULL,  
        names.arg = NULL, legend.text = NULL, beside = FALSE,  
        horiz = FALSE, density = NULL, angle = 45,  
        col = NULL, border = par("fg"),  
        main = NULL, sub = NULL, xlab = NULL, ylab = NULL,  
        xlim = NULL, ylim = NULL, xpd = TRUE, log = "",  
        axes = TRUE, axisnames = TRUE,  
        cex.axis = par("cex.axis"), cex.names = par("cex.axis"),  
        inside = TRUE, plot = TRUE, axis.lty = 0, offset = 0,  
        add = FALSE, ann = !add && par("ann"), args.legend = NULL,  
        ...)
```

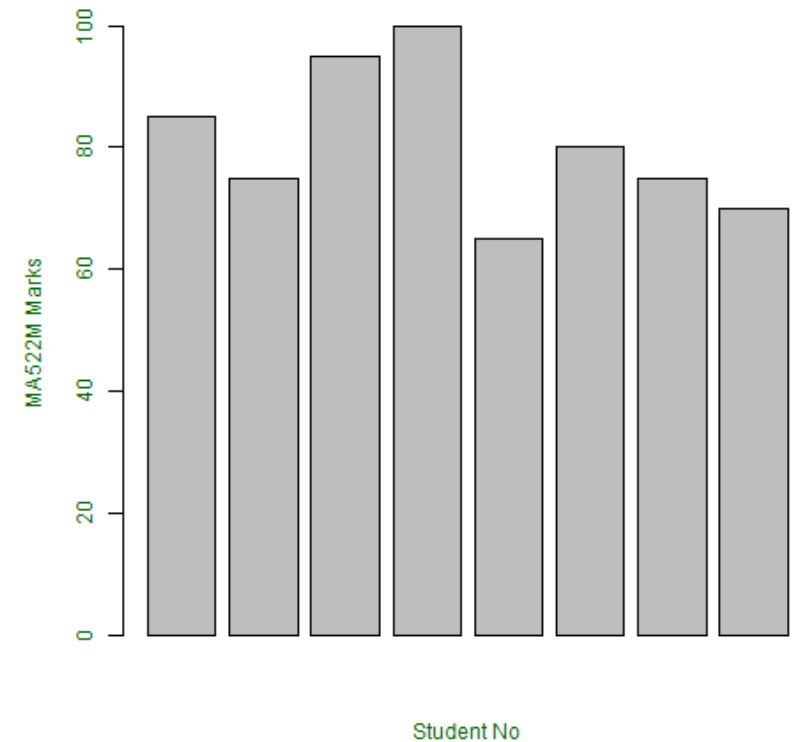
➤ *Values in data vector as height of the bars*

## ✓ For Formula

```
barplot(formula, data, subset, na.action,  
        horiz = FALSE, xlab = NULL, ylab = NULL, ...)
```

# Bar Plot/Chart-Example

```
marks=c(85, 75, 95, 100, 65, 80, 75, 70)
png(file="barplot.png")
barplot(marks,xlab="Student No",ylab="MA522M
Marks",col.axis="darkgreen",col.lab="darkgreen")
dev.off()
```



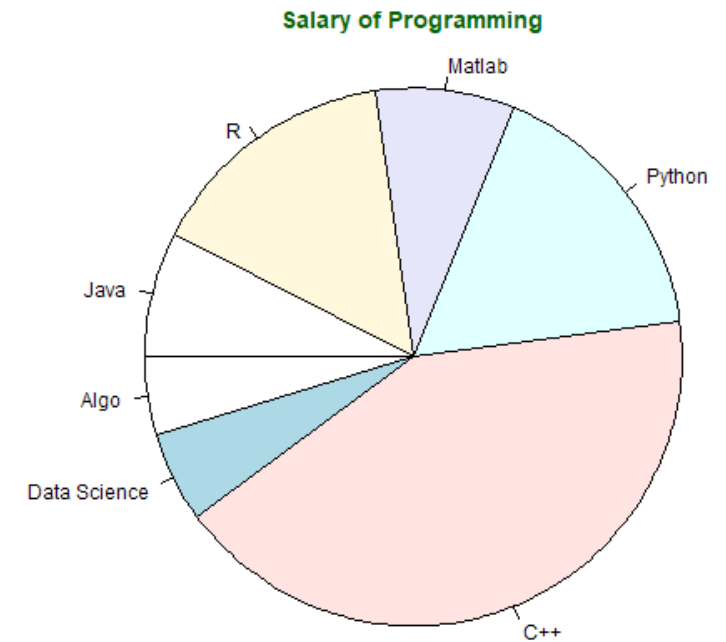
➤ **Circular chart divided into different segments according to the ratio**

## ✓ Syntax

```
pie(x, labels = names(x), edges = 200, radius = 0.8,  
    clockwise = FALSE, init.angle = if(clockwise) 90 else 0,  
    density = NULL, angle = 45, col = NULL, border = NULL,  
    lty = NULL, main = NULL, ...)
```

# Pie Plot/Chart-Example

```
salary=c(28, 32, 250, 100, 50, 90, 45)
names(salary)=c("Algo", "Data Science", "C++", "Python", "Matlab", "R", "Java")
png(file="pieplot.png")
pie(salary, labels = names(salary), main="Salary of Programming", radius = -
1, col.main="darkgreen")
dev.off()
```



➤ **Circular chart divided into different segments according to the ratio**

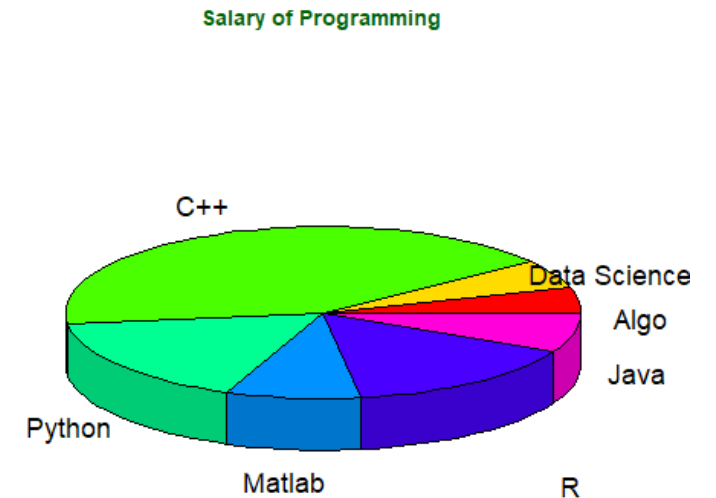
## ✓ Syntax

```
pie3D(x, edges=NA, radius=1, height=0.1, theta=pi/6, start=0, border=par("fg"),  
      col=NULL, labels=NULL, labelpos=NULL, labelcol=par("fg"), labelcex=1.5,  
      sector.order=NULL, explode=0, shade=0.8, mar=c(4, 4, 4, 4), pty="s", ...)
```



# Pie3D Plot/Chart-Example

```
install.packages("plotrix")
require(plotrix)
salary=c(28, 32, 250, 100, 50, 90, 45)
names(salary)=c("Algo", "Data Science", "C++", "Python", "Matlab", "R", "Java")
png(file="pieplot3D.png")
pie3D(salary, labels = names(salary), main="Salary of
Programming", col.main="darkgreen")
dev.off()
```

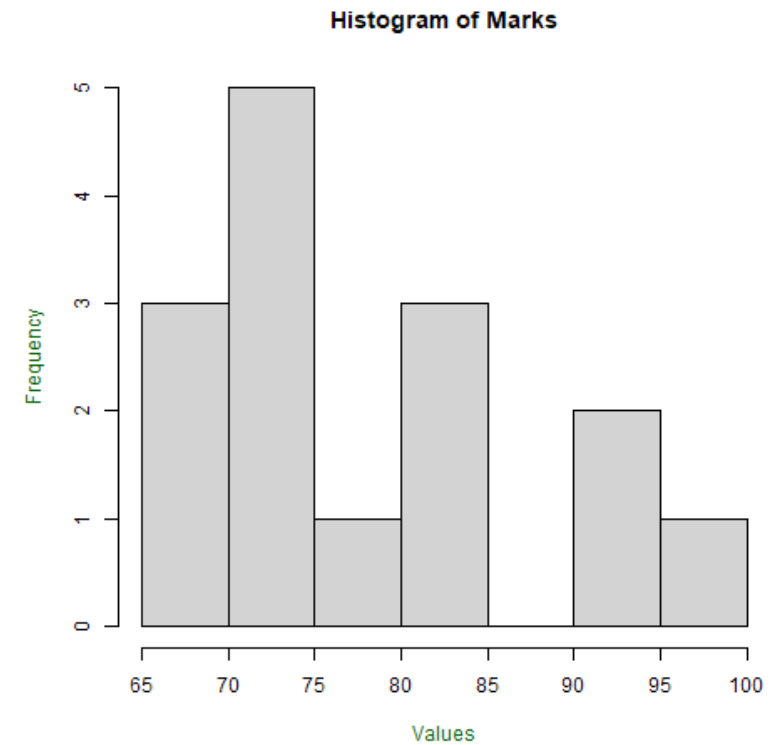


## ✓ Syntax

```
hist(x, breaks = "Sturges",  
     freq = NULL, probability = !freq,  
     include.lowest = TRUE, right = TRUE,  
     density = NULL, angle = 45, col = NULL, border = NULL,  
     main = paste("Histogram of" , xname),  
     xlim = range(breaks), ylim = NULL,  
     xlab = xname, ylab,  
     axes = TRUE, plot = TRUE, labels = FALSE,  
     nclass = NULL, warn.unused = TRUE, ...)
```

# Histogram-Example

```
marks=c(85, 75, 95, 100, 65, 80, 75, 70, 65, 85, 75, 95, 75, 85, 75)
png(file="histo.png")
hist(marks,main="Histogram of Marks",xlab="Values",col.lab="darkgreen")
dev.off()
```



## ➤ A matrix of scatterplots

### ✓ Default Syntax

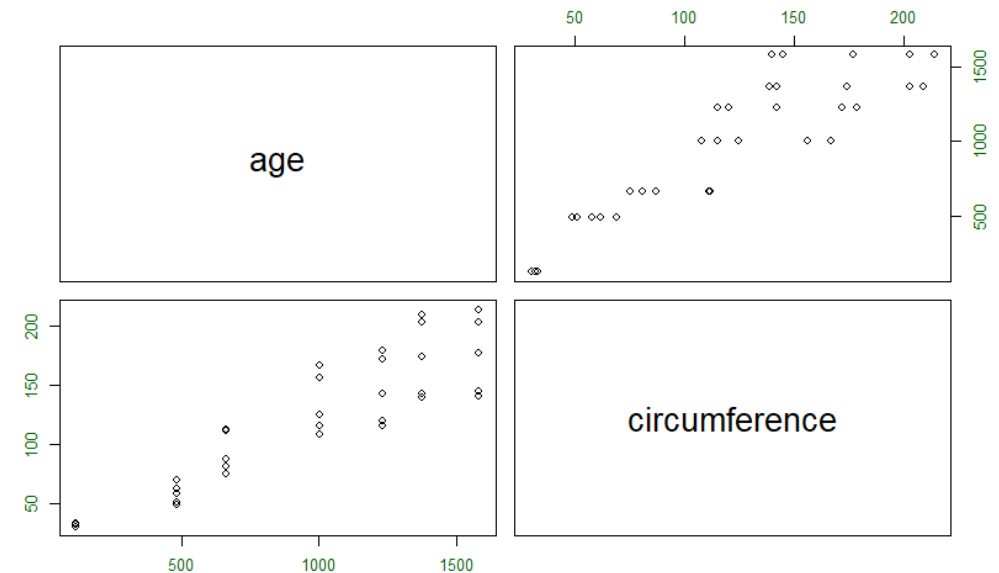
```
pairs(x, labels, panel = points, ...,  
      horInd = 1:nc, verInd = 1:nc,  
      lower.panel = panel, upper.panel = panel,  
      diag.panel = NULL, text.panel = textPanel,  
      label.pos = 0.5 + has.diag/3, line.main = 3,  
      cex.labels = NULL, font.labels = 1,  
      rowlattop = TRUE, gap = 1, log = "",  
      horOdd = !rowlattop, verOdd = !rowlattop)
```

➤ **Values in data vector as height of the bars**

## ✓ For Formula

```
pairs(formula, data = NULL, ..., subset,  
      na.action = stats::na.pass)
```

```
pairs(~age+circumference, data=Orange, col.axis="darkgreen")
```



## ➤ **Explore Other plots**

assocplot

abline

boxplot

curve

cdplot

coplot

contour

matplot

polygon

spineplot

stars

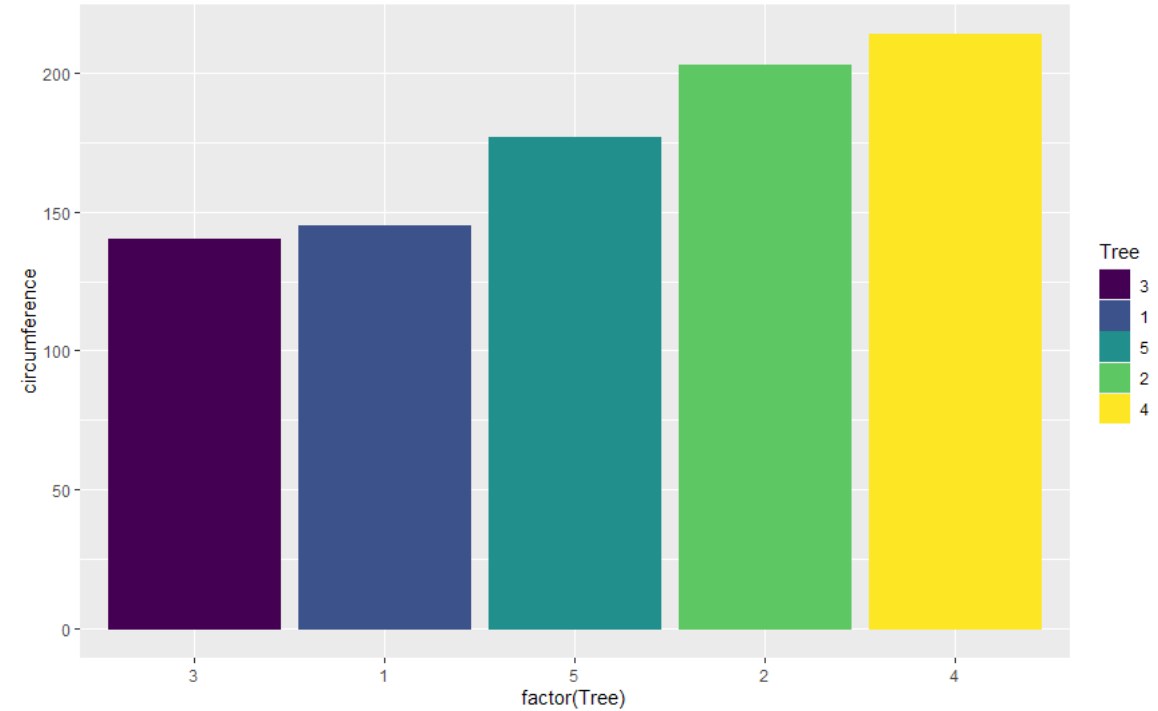
stem

sunflowerplot

<https://www.rdocumentation.org/packages/graphics/versions/3.6.2>

```
library(ggplot2)
ggplot(Orange, aes(x=factor(Tree), y=circumference, fill=Tree, color=Tree))
+geom_bar(stat='identity', position='dodge')
```

- **Ggplot2 is a system for declaratively creating graphics**
- **Map variables to aesthetics**
- <https://ggplot2.tidyverse.org/>



# BUILT-IN R DATA



- ***R has a collection of dataset***
- ***In fact, it has tons of built-in datasets that can be used for demo or benchmark purposes***
- ***There are few widely used datasets***
- ***For more details, please visit: <https://stat.ethz.ch/R-manual/R-devel/library/datasets/html/00Index.html>***

# Mostly used datasets

Dataset	Description
datasets	Base R datasets
airquality	New York Air Quality Measurements
ToothGrowth	Experiment studying the effect of Vitamin C
PlantGrowth	Yields obtained under a control of two different treatment conditions
USArrests	Statistics about violent crime rates by the US state
AirPassengers	Monthly Airline Passenger Numbers 1949-1960
mtcars	Motor Trend Car Road Tests
Iris	Edgara Anderson's Iris Data
Orange	Growth of Orange Trees
Quakes	Locations of Earthquakes in Fuji
JohnsonJohnson	Quarterly Earning of Johnson & Johnson Share
women	Average Heights and Weights for American Women
uspop	Populations recorded by the US Census

- **Data extracted from 1974 Motor Trend US Magazine**
- **32 observations, 11 columns**

mpg	Miles/(US) gallon
cyl	Number of cylinders
disp	Displacement (cu.in.)
hp	Gross horsepower
drat	Rear axle ratio
wt	Weight (1000 lbs)
qsec	1/4 mile time
vs	Engine (0 = V-shaped, 1 = straight)
am	Transmission (0 = automatic, 1 = manual)
gear	Number of forward gears
carb	Number of carburetors

- ***Classic Box & Jenkins airline data***
- ***Monthly totals of international airline passengers***
- ***1949-1960***

➤ **Daily airquality measurements of New York, May – Sep 1973**

Ozone	Ozone (ppb)
Solar.R	Solar R (lang)
Wind	Wind (mph)
Temp	Temperature (degrees F)
Month	Month (1--12)
Day	Day of month (1--31)

- Famous (Fisher's or Anderson's) iris data
- Measurements in centimeters of the variables sepal length and width
- Petal length and width
- 50 flowers from each of 3 species of iris



- Growth of orange trees
- 35 rows and 3 columns
- Tree, age, circumference



```
summary(mtcars[1])  
      mpg  
Min.   :10.40  
1st Qu.:15.43  
Median :19.20  
Mean   :20.09  
3rd Qu.:22.80  
Max.   :33.90
```



```
summary(airquality)
```

Ozone	Solar.R	Wind	Temp	Month	Day
Min. : 1.00	Min. : 7.0	Min. : 1.700	Min. : 56.00	Min. : 5.000	Min. : 1.0
1st Qu.: 18.00	1st Qu.: 115.8	1st Qu.: 7.400	1st Qu.: 72.00	1st Qu.: 6.000	1st Qu.: 8.0
Median : 31.50	Median : 205.0	Median : 9.700	Median : 79.00	Median : 7.000	Median : 16.0
Mean : 42.13	Mean : 185.9	Mean : 9.958	Mean : 77.88	Mean : 6.993	Mean : 15.8
3rd Qu.: 63.25	3rd Qu.: 258.8	3rd Qu.: 11.500	3rd Qu.: 85.00	3rd Qu.: 8.000	3rd Qu.: 23.0
Max. : 168.00	Max. : 334.0	Max. : 20.700	Max. : 97.00	Max. : 9.000	Max. : 31.0
NA's : 37	NA's : 7				

```
> print(AirPassengers)
```

	Jan	Feb	Mar	Apr	May	Jun	Jul	Aug	Sep	Oct	Nov	Dec
1949	112	118	132	129	121	135	148	148	136	119	104	118
1950	115	126	141	135	125	149	170	170	158	133	114	140
1951	145	150	178	163	172	178	199	199	184	162	146	166
1952	171	180	193	181	183	218	230	242	209	191	172	194
1953	196	196	236	235	229	243	264	272	237	211	180	201
1954	204	188	235	227	234	264	302	293	259	229	203	229
1955	242	233	267	269	270	315	364	347	312	274	237	278
1956	284	277	317	313	318	374	413	405	355	306	271	306
1957	315	301	356	348	355	422	465	467	404	347	305	336
1958	340	318	362	348	363	435	491	505	404	359	310	337
1959	360	342	406	396	420	472	548	559	463	407	362	405
1960	417	391	419	461	472	535	622	606	508	461	390	432

**READ CSV**

➤ ***It is part of utils packages***

```
read.csv(file, header = TRUE, sep = ",", quote = "\"",  
         dec = ".", fill = TRUE, comment.char = "", ...)
```

```
read.csv(tf, fill = TRUE) # 1 column  
ncol <- max(count.fields(tf, sep = ","))  
read.csv(tf, fill = TRUE, header = FALSE,  
         col.names = paste0("V", seq_len(ncol)))
```

```
data=read.csv("input.csv")  
print(data)  
  
print(is.data.frame(data))  
print(ncol(data))  
print(nrow(data))
```

```
read.table(file, header = FALSE, sep = "", quote = "\"'",  
          dec = ".", numerals = c("allow.loss", "warn.loss", "no.loss"),  
          row.names, col.names, as.is = !stringsAsFactors,  
          na.strings = "NA", colClasses = NA, nrow = -1,  
          skip = 0, check.names = TRUE, fill = !blank.lines.skip,  
          strip.white = FALSE, blank.lines.skip = TRUE,  
          comment.char = "#",  
          allowEscapes = FALSE, flush = FALSE,  
          stringsAsFactors = default.stringsAsFactors(),  
          fileEncoding = "", encoding = "unknown", text, skipNul = FALSE)
```

# FREQUENCY TABLE

- **A list of objects with the frequency of each**
  - **Useful for categorical data**
- ✓ **Default Syntax**

```
table(data)
```

```
marks=c(85, 75, 95, 100, 65, 80, 75, 70, 65, 85, 75, 95, 75, 85, 75)
```

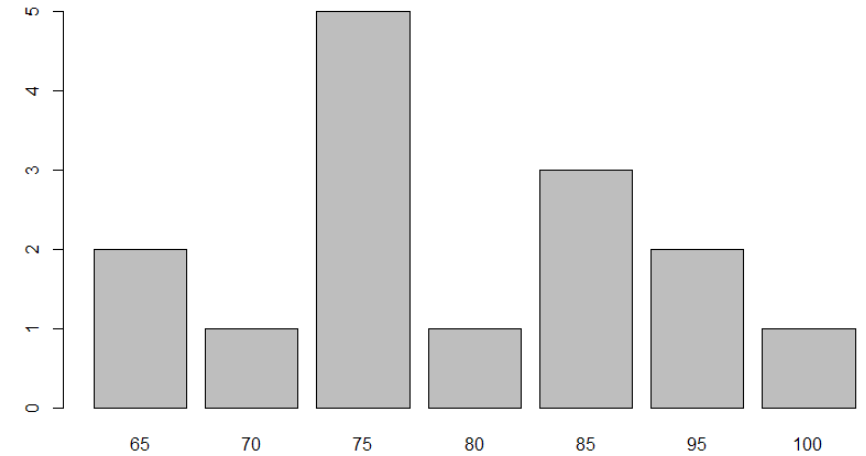
```
freq=table(marks)
```

```
print(freq)
```

```
marks
```

```
 65  70  75  80  85  95 100  
  2   1   5   1   3   2   1
```

```
barplot(freq)
```





# Frequency table with proportions

```
marks=c(85,75,95,100,65,80,75,70,65,85,75,95,75,85,75)
```

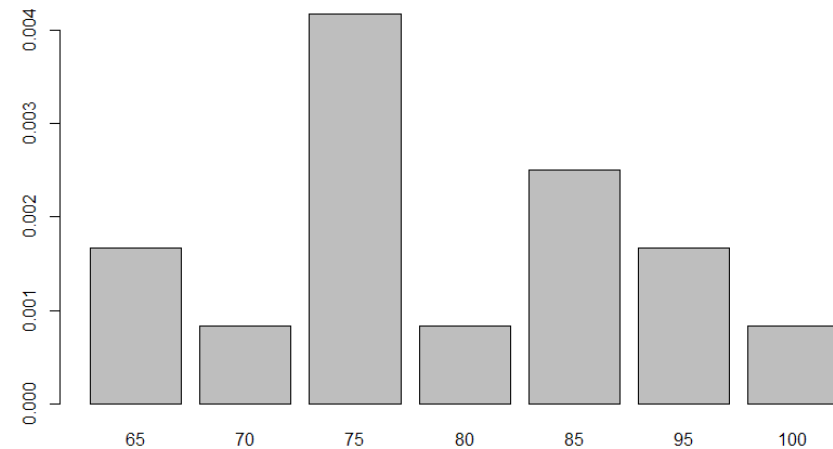
```
freq=table(marks)/sum(marks)
```

```
print(freq)
```

```
marks
```

```
          65          70          75          80          85
95          100
0.001666667 0.000833333 0.004166667 0.000833333 0.002500000
0.001666667 0.000833333
```

```
barplot(freq)
```



# Cumulative Frequency table

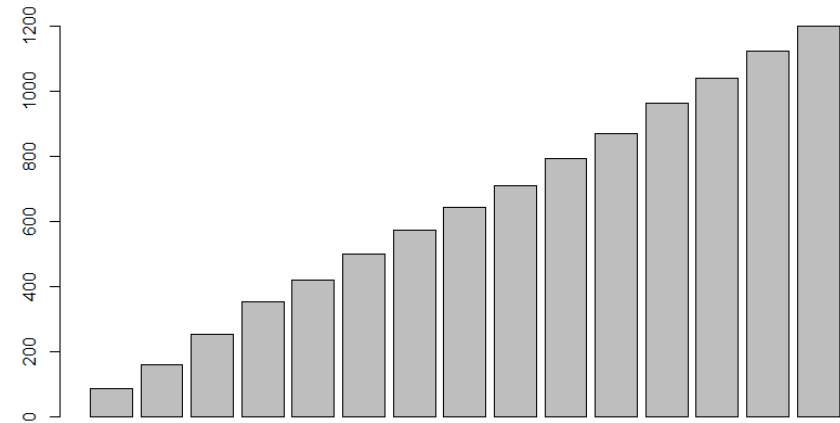
```
marks=c(85,75,95,100,65,80,75,70,65,85,75,95,75,85,75)
```

```
freq=cumsum(marks)
```

```
print(freq)
```

```
[1] 85 160 255 355 420 500 575 645 710 795 870 965  
1040 1125 1200
```

```
barplot(freq)
```



# Two-way Frequency Table

```
marks=c(85,75,95,100,65,80,75,70,65,85,75,95,75,85,75)
```

```
passfaile=c(1,1,1,1,0,1,1,1,0,1,1,1,1,1,1)
```

```
table(marks,passfaile)
```

```
      passfaile
```

```
marks 0 1
```

```
65    2 0
```

```
70    0 1
```

```
75    0 5
```

```
80    0 1
```

```
85    0 3
```

```
95    0 2
```

```
100   0 1
```

```
library(dplyr)
marks=c(85, 75, 95, 100, 65, 80, 75, 70, 65, 85, 75, 95, 75, 85, 75)
pf=c(1, 1, 1, 1, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1)
data=data.frame(Marks=marks, pf=pf)
tot=data %>% group_by(Marks) %>% summarize(tot=sum(Marks))
print(tot)
# A tibble: 7 × 2
  Marks tot
  <dbl> <dbl>
1     65  130
2     70   70
3     75  375
4     80   80
5     85  255
6     95  190
7    100  100
```

```
read.table(header = TRUE, text = "  
a b  
1 2  
3 4  
")  
# }
```

**WRITE CSV**

➤ ***It is part of utils packages***

```
write.csv(data,path)
x=1:5
name=c("MA23M001","MA23M002","MA23M003","MA23M004","MA23M005")
marks=c(100,98,77,89,67)
DF=data.frame(x,name,marks)
write.csv(DF,"students.csv")
```

# READ/WRITE XLSX



## ➤ **It is part of xlsx packages**

```
install.packages("xlsx")
```

```
require(xlsx)
```

```
readxlsx=read.xlsx(path, sheeIndex=1)
```

```
writexlsx=write.xlsx(data, path, col.names=TRUE, row.name=TRUE, sheetName="Sheet2", append=TRUE)
```

## ➤ It is part of utils packages

```
x=1:5
name=c("MA23M001", "MA23M002", "MA23M003", "MA23M004", "MA23M005")
marks=c(100, 98, 77, 89, 67)
DF=data.frame(x, name, marks)
write.xlsx(DF, file = "employee.xlsx", col.names=TRUE,
row.names=TRUE, sheetName="Sheet2", append = TRUE)
read.xlsx("employee.xlsx", sheetIndex = 1)
```

# READ/WRITE JSON

➤ ***It is part of rjson packages***

```
install.packages("rjson")  
require(rjson)  
readjson=fromJSON(path, shexIndex=1)  
jsondata=toJSON(data)  
write(jsondata,path)
```

## ➤ *It is part of rjson packages*

```
list1 <- vector(mode="list", length=2)
list1[[1]] <- c("sunflower", "guava", "hibiscus")
list1[[2]] <- c("flower", "fruit", "flower")

# creating the data for JSON file
jsonData <- toJSON(list1)

# writing into JSON file
write(jsonData, "result.json")

# Give the created file name to the function
result <- fromJSON(file = "result.json")

# Print the result
print(result)
```