

assignment la1

2023-05-15

`help()`-Brings up a help entry for the specified command

`help.start()`-Opens the help system in the browser

`apropos()`-shows all the commands that contain the “partword”

`install.packages(“pkg”)`-Installs a library of commands from the CRAN website

`install.packages()`-Shows a list of packages that are installed.

`library(pkg)`-Loads a package of commands,making them available for use

`search()`-Shows a list of packages that are available and loaded for use

`example()`-View some examples on the use of a command

`ls()`-Lists memory of contents

`ls.str(pattern=‘data’)`-This lists all the objects with “data” in their name and shows you the structure of each.

`ls(pattern=‘b’)`-Here the pattern looks for everything containing a “b”

`ls(pattern=“^be”)`-If you want to search for objects beginning with a certain letter you use the ^ character

`ls(pattern=“1”)`-letters are enclosed in square brackets the square brackets is to isolate the letters; each is treated as a separate item, hence objects beginning with “b” or “e” are matched

`ls(pattern=‘m$’)`-To find objects ending with a specific character you use a dollar sign at the end

`ls(pattern=‘a.e’)`-You can use the period as a wildcard and R will match any character

`class()`-You can obtain information about the type of object

`sort()`-The `sort()` function is used to sort the elements of a vector or a data frame in ascending order. It returns a new vector or data frame with the elements arranged in ascending order.

`rank()`-The `rank()` function computes the ranks of the elements in a vector. It assigns a unique rank to each element based on its value, where the smallest element receives a rank of 1.

`method()`: The `method()` function is not a built-in R function. The term “method” typically refers to a specific implementation or algorithm for solving a problem in R.

`row.names()`-function in R is used to retrieve or set the row names of a matrix or a data frame.

`history()`-You can view the current list of the history items

`savehistory(file=‘.Rhistory’)`-You can save the current history to a desk

`loadhistory(file=‘.Rhistory’)`-You can load the file from the history

`c(),scan()`-Enter data manually to a vector in R

`rep()`-Make Vector of repeated values

`seq()`-Make arithmetic progression vector

`View()`-View dataset in a spreadsheet-type format

`data()`-Load built-in dataset

`dim()`-See dimensions(rows/columns)of a dataframe

`str()`-Display internal structure of an R-object

`read.csv(),read.table()`-load into data frame an existing datafile

`require()`-Make available an R add-on package

`names()`-List names of variables in a data frame

`length()`-Give length of a vector

`rm()`-removes an item from memory

`hist()`-Command for producing a histogram

`histogram()`-Lattice command for producing a histogram

`table()`-List all values of a variables with frequencies

`stem()`-Make a stem plot

`xtabs()`-Cross tabulation tables using formulas

`mosaicplot()`-Make a mosaicplot

cut()-Group values of a variable into larger bins
median()-Identify "center" of distribution
mean()-To find the avg of the given data
by()-Apply function to a column split by factors
summary()-Display 5-number summary and mean
var()-To find the variance of a given values
sd()-To find the standard variation of the given values in the vector
sum()-Add up all vlaues in vector
quantile()-Find the position of a quantile in a dataset
barplot()-Produce a bargraph
barchart()-Lattice command for producing bar graphs
boxplot()-Produce a boxplot
bwplot()-Lattice command for producing boxplots
plot()-To produce a scatter plot
xyplot()-Lattice command for producing a scatter plot
lm()-Determine the least-squares regression line
anova()-Analyzing of variance
predict()-Obtained predicted values from linear model
nls()-Estimate the parameters of the nonlinear model
residuals()-Gives for a model-fit to data
sample()-Take a smaple from vector of data
replicate()-repeat process a set number of times
cumsum()-produce running total of values for an input vector
ecdf()-Buittls empricial cumulative distribution function
dbinom()-Tools for binomial distribution
dpois()-Tools for poisson distribution
pnorm()-Tools for normal distribution
qt()-Tools for student t distributions
pchisq()-Tools for chi-square test distributions
binom.test()-Hypothesis test and confident interval for one proportion
prop.test()-inferencce for one portion using normal approx
chisq.test()-Carries out a chi-square test
fisher.test()-Fisher test for contingency table
t.test()-Studnet t test for inference on population mean
qqnorm(),qqline()-Tools for checking normality
addmargins()-Add marginal sums to an existing table
prop.table()-Compute proportions from a contingency table
par()-query and edit graphical settings
power.t.test()-power calculations for one and two sample t test

```
apropos("partword")
```

```
## character(0)
```

```
installed.packages()
```

```
##      Package  
## base64enc  "base64enc"  
## bslib     "bslib"  
## cachem    "cachem"  
## cellranger "cellranger"
```

```

## cellaranger "cellaranger"
## cli "cli"
## commonmark "commonmark"
## cpp11 "cpp11"
## crayon "crayon"
## digest "digest"
## ellipsis "ellipsis"
## evaluate "evaluate"
## fansi "fansi"
## fastmap "fastmap"
## fontawesome "fontawesome"
## fs "fs"
## glue "glue"
## highr "highr"
## hms "hms"
## htmltools "htmltools"
## jquerylib "jquerylib"
## jsonlite "jsonlite"
## knitr "knitr"
## lifecycle "lifecycle"
## magrittr "magrittr"
## markdown "markdown"
## memoise "memoise"
## mime "mime"
## pillar "pillar"
## pkgconfig "pkgconfig"
## prettyunits "prettyunits"
## progress "progress"
## R6 "R6"
## rappdirs "rappdirs"
## Rcpp "Rcpp"
## readxl "readxl"
## rematch "rematch"
## rlang "rlang"
## rmarkdown "rmarkdown"
## sass "sass"
## stringi "stringi"
## stringr "stringr"
## tibble "tibble"
## tinytex "tinytex"
## utf8 "utf8"
## vctrs "vctrs"
## xfun "xfun"
## yaml "yaml"
## base "base"
## boot "boot"
## class "class"
## cluster "cluster"
## codetools "codetools"
## compiler "compiler"
## datasets "datasets"
## foreign "foreign"
## graphics "graphics"
## grDevices "grDevices"
## grid "grid"
## KernSmooth "KernSmooth"
## lattice "lattice"
## MASS "MASS"
## Matrix "Matrix"
## methods "methods"
## mgcv "mgcv"
## nlme "nlme"
## nnet "nnet"
## parallel "parallel"
## rpart "rpart"
## spatial "spatial"
## splines "splines"
## stats "stats"
## stats4 "stats4"
## survival "survival"
## tcltk "tcltk"
## tools "tools"
## translations "translations"
## utils "utils"
## LibPath Version
## base64enc "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.1-3"
## bslib "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.4.2"
## cachem "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.0.8"
## callr "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.1.0"

```

```

## cellranger "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.1.0"
## cli "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "3.6.1"
## commonmark "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.9.0"
## cpp11 "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.4.3"
## crayon "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.5.2"
## digest "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.6.31"
## ellipsis "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.3.2"
## evaluate "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.21"
## fansi "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.0.4"
## fastmap "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.1.1"
## fontawesome "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.5.1"
## fs "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.6.2"
## glue "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.6.2"
## highr "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.10"
## hms "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.1.3"
## htmltools "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.5.5"
## jquerylib "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.1.4"
## jsonlite "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.8.4"
## knitr "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.42"
## lifecycle "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.0.3"
## magrittr "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "2.0.3"
## markdown "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.6"
## memoise "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "2.0.1"
## mime "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.12"
## pillar "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.9.0"
## pkgconfig "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "2.0.3"
## prettyunits "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.1.1"
## progress "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.2.2"
## R6 "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "2.5.1"
## rappdirs "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.3.3"
## Rcpp "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.0.10"
## readxl "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.4.2"
## rematch "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.0.1"
## rlang "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.1.1"
## rmarkdown "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "2.21"
## sass "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.4.6"
## stringi "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.7.12"
## stringr "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.5.0"
## tibble "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "3.2.1"
## tinytex "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.45"
## utf8 "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "1.2.3"
## vctrs "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.6.2"
## xfun "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "0.39"
## yaml "C:/Users/jayam tulasi/AppData/Local/R/win-library/4.3" "2.3.7"
## base "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## boot "C:/Program Files/R/R-4.3.0/library" "1.3-28.1"
## class "C:/Program Files/R/R-4.3.0/library" "7.3-21"
## cluster "C:/Program Files/R/R-4.3.0/library" "2.1.4"
## codetools "C:/Program Files/R/R-4.3.0/library" "0.2-19"
## compiler "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## datasets "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## foreign "C:/Program Files/R/R-4.3.0/library" "0.8-84"
## graphics "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## grDevices "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## grid "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## KernSmooth "C:/Program Files/R/R-4.3.0/library" "2.23-20"
## lattice "C:/Program Files/R/R-4.3.0/library" "0.21-8"
## MASS "C:/Program Files/R/R-4.3.0/library" "7.3-58.4"
## Matrix "C:/Program Files/R/R-4.3.0/library" "1.5-4"
## methods "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## mgcv "C:/Program Files/R/R-4.3.0/library" "1.8-42"
## nlme "C:/Program Files/R/R-4.3.0/library" "3.1-162"
## nnet "C:/Program Files/R/R-4.3.0/library" "7.3-18"
## parallel "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## rpart "C:/Program Files/R/R-4.3.0/library" "4.1.19"
## spatial "C:/Program Files/R/R-4.3.0/library" "7.3-16"
## splines "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## stats "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## stats4 "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## survival "C:/Program Files/R/R-4.3.0/library" "3.5-5"
## tcltk "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## tools "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## translations "C:/Program Files/R/R-4.3.0/library" "4.3.0"
## utils "C:/Program Files/R/R-4.3.0/library" "4.3.0"
##
## Priority Depends
## base64enc NA "R (>= 2.9.0)"
## bslib NA "R (>= 2.10)"
## cachem NA NA
## callr NA "R (>= 3.6.2)"

```

```

## cellranger NA "R (>= 3.0.0)"
## cli NA "R (>= 3.4)"
## commonmark NA NA
## cpp11 NA NA
## crayon NA NA
## digest NA "R (>= 3.3.0)"
## ellipsis NA "R (>= 3.2)"
## evaluate NA "R (>= 3.0.2)"
## fansi NA "R (>= 3.1.0)"
## fastmap NA NA
## fontawesome NA "R (>= 3.3.0)"
## fs NA "R (>= 3.4)"
## glue NA "R (>= 3.4)"
## highr NA "R (>= 3.3.0)"
## hms NA NA
## htmltools NA "R (>= 2.14.1)"
## jquerylib NA NA
## jsonlite NA "methods"
## knitr NA "R (>= 3.3.0)"
## lifecycle NA "R (>= 3.4)"
## magrittr NA "R (>= 3.4.0)"
## markdown NA "R (>= 2.11.1)"
## memoise NA NA
## mime NA NA
## pillar NA NA
## pkgconfig NA NA
## prettyunits NA NA
## progress NA NA
## R6 NA "R (>= 3.0)"
## rappdirs NA "R (>= 3.2)"
## Rcpp NA NA
## readxl NA "R (>= 3.5)"
## rematch NA NA
## rlang NA "R (>= 3.5.0)"
## rmarkdown NA "R (>= 3.0)"
## sass NA NA
## stringi NA "R (>= 3.1)"
## stringr NA "R (>= 3.3)"
## tibble NA "R (>= 3.4.0)"
## tinytex NA NA
## utf8 NA "R (>= 2.10)"
## vctrs NA "R (>= 3.5.0)"
## xfun NA NA
## yaml NA NA
## base "base" NA
## boot "recommended" "R (>= 3.0.0), graphics, stats"
## class "recommended" "R (>= 3.0.0), stats, utils"
## cluster "recommended" "R (>= 3.5.0)"
## codetools "recommended" "R (>= 2.1)"
## compiler "base" NA
## datasets "base" NA
## foreign "recommended" "R (>= 4.0.0)"
## graphics "base" NA
## grDevices "base" NA
## grid "base" NA
## KernSmooth "recommended" "R (>= 2.5.0), stats"
## lattice "recommended" "R (>= 4.0.0)"
## MASS "recommended" "R (>= 4.3.0), grDevices, graphics, stats, utils"
## Matrix "recommended" "R (>= 3.5.0), methods"
## methods "base" NA
## mgcv "recommended" "R (>= 3.6.0), nlme (>= 3.1-64)"
## nlme "recommended" "R (>= 3.5.0)"
## nnet "recommended" "R (>= 3.0.0), stats, utils"
## parallel "base" NA
## rpart "recommended" "R (>= 2.15.0), graphics, stats, grDevices"
## spatial "recommended" "R (>= 3.0.0), graphics, stats, utils"
## splines "base" NA
## stats "base" NA
## stats4 "base" NA
## survival "recommended" "R (>= 3.5.0)"
## tcltk "base" NA
## tools "base" NA
## translations NA NA
## utils "base" NA
## Imports

## base64enc NA

```

```
## bslib      "grDevices, htmltools (>= 0.5.4), jsonlite, sass (>= 0.4.0), jquerylib (>= 0.1.3), rlang, cachem, memoise (>= 2.0.1), \nbase64enc, mime"

## cachem     "rlang, fastmap (>= 1.1.1)"

## cellranger  "rematch, tibble"

## cli        "utils"

## commonmark  NA

## cpp11       NA

## crayon     "grDevices, methods, utils"

## digest     "utils"

## ellipsis   "rlang (>= 0.3.0)"

## evaluate   "methods"

## fansi      "grDevices, utils"

## fastmap     NA

## fontawesome "rlang (>= 1.0.6), htmltools (>= 0.5.1.1)"

## fs         "methods"

## glue       "methods"

## highr      "xfun (>= 0.18)"

## hms        "lifecycle, methods, pkgconfig, rlang (>= 1.0.2), vctrs (>= 0.3.8)"

## htmltools  "utils, digest, grDevices, base64enc, rlang (>= 0.4.10), \nfastmap (>= 1.1.0), ellipsis"

## jquerylib  "htmltools"

## jsonlite   NA

## knitr      "evaluate (>= 0.15), highr, methods, yaml (>= 2.1.19), xfun (>= 0.34), tools"

## lifecycle  "cli (>= 3.4.0), glue, rlang (>= 1.0.6)"

## magrittr   NA

## markdown   "utils, commonmark (>= 1.9.0), xfun (>= 0.38)"

## memoise    "rlang (>= 0.4.10), cachem"

## mime       "tools"

## pillar     "cli (>= 2.3.0), fansi, glue, lifecycle, rlang (>= 1.0.2), utf8 (>= 1.1.0), utils, vctrs (>= 0.5.0)"

## pkgconfig  "utils"

## prettyunits NA

## progress   "hms, prettyunits, R6, crayon"

## R6         NA

## rappdirs   NA

## Rcpp       "methods, utils"

## readxl     "cellranger, tibble (>= 2.0.1), utils"

## rematch    NA

## rlang      "utils"

## rmarkdown  "bslib (>= 0.2.5.1), evaluate (>= 0.13), fontawesome (>= 0.5.0), htmltools (>= 0.5.1), jquerylib, jsonlite, knitr (>= 1.22), methods, stringr (>= 1.2.0), tinytex (>= 0.31), tools, \nutils, xfun (>= 0.36), yaml (>= 2.1.19)"
## sass       "fs, rlang (>= 0.4.10), htmltools (>= 0.5.1), R6, rappdirs"

## stringi    "tools, utils, stats"
```

```
## stringr      "cli, glue (>= 1.6.1), lifecycle (>= 1.0.3), magrittr, rlang\n(>= 1.0.0), stringi (>= 1.5.3), vctrs"

## tibble      "fansI (>= 0.4.0), lifecycle (>= 1.0.0), magrittr, methods,\npillar (>= 1.8.1), pkgconfig, rlang (>= 1.0.2), utils, vctrs\n(>= 0.4.2)"

## tinytex     "xfun (>= 0.29)"

## utf8        NA

## vctrs       "cli (>= 3.4.0), glue, lifecycle (>= 1.0.3), rlang (>= 1.1.0)"

## xfun        "stats, tools"

## yaml        NA

## base        NA

## boot        NA

## class       "MASS"

## cluster     "graphics, grDevices, stats, utils"

## codetools   NA

## compiler    NA

## datasets    NA

## foreign     "methods, utils, stats"

## graphics    "grDevices"

## grDevices   NA

## grid        "grDevices, utils"

## KernSmooth  NA

## lattice     "grid, grDevices, graphics, stats, utils"

## MASS        "methods"

## Matrix      "graphics, grid, lattice, stats, utils"

## methods     "utils, stats"

## mgcv        "methods, stats, graphics, Matrix, splines, utils"

## nlme        "graphics, stats, utils, lattice"

## nnet        NA

## parallel    "tools, compiler"

## rpart       NA

## spatial     NA

## splines     "graphics, stats"

## stats       "utils, grDevices, graphics"

## stats4      "graphics, methods, stats"

## survival    "graphics, Matrix, methods, splines, stats, utils"

## tcltk       "utils"

## tools       NA

## translations NA

## utils       NA

##             LinkingTo
## base64enc    NA
## bslib        NA
## cachem       NA
```

```
## cellranger NA
## cli NA
## commonmark NA
## cpp11 NA
## crayon NA
## digest NA
## ellipsis NA
## evaluate NA
## fansi NA
## fastmap NA
## fontawesome NA
## fs NA
## glue NA
## highr NA
## hms NA
## htmltools NA
## jquerylib NA
## jsonlite NA
## knitr NA
## lifecycle NA
## magrittr NA
## markdown NA
## memoise NA
## mime NA
## pillar NA
## pkgconfig NA
## prettyunits NA
## progress NA
## R6 NA
## rappdirs NA
## Rcpp NA
## readxl "cpp11 (>= 0.4.0), progress"
## rematch NA
## rlang NA
## rmarkdown NA
## sass NA
## stringi NA
## stringr NA
## tibble NA
## tinytex NA
## utf8 NA
## vctrs NA
## xfun NA
## yaml NA
## base NA
## boot NA
## class NA
## cluster NA
## codetools NA
## compiler NA
## datasets NA
## foreign NA
## graphics NA
## grDevices NA
## grid NA
## KernSmooth NA
## lattice NA
## MASS NA
## Matrix NA
## methods NA
## mgcv NA
## nlme NA
## nnet NA
## parallel NA
## rpart NA
## spatial NA
## splines NA
## stats NA
## stats4 NA
## survival NA
## tcltk NA
## tools NA
## translations NA
## utils NA
## Suggests

## base64enc NA
```



```

## bslib      "shiny (>= 1.6.0), rmarkdown (>= 2.7), thematic, knitr,\ntestthat, withr, rappdirs, curl, magrittr, fontawesome, bsicons"

## cachem     "testthat"

## cellranger  "covr, testthat (>= 1.0.0), knitr, rmarkdown"

## cli        "callr, covr, crayon, digest, glue (>= 1.6.0), grDevices,\nhtmltools, htmlwidgets, knitr, methods, mockery, processx, ps\n(>= 1.3.4.9000), rlang (  
>= 1.0.2.9003), rmarkdown, rprojroot,\nrstudioapi, testthat, tibble, whoami, withr"

## commonmark  "curl, testthat, xml2"

## cpp11      "bench, brio, callr, cli, covr, decor, desc, ggplot2, glue,\nknitr, lobstr, mockery, progress, rmarkdown, scales, Rcpp,\nntestthat, tibble, utils, vct  
rs, withr"
## crayon     "mockery, rstudioapi, testthat, withr"

## digest     "tinytest, simplermardown"

## ellipsis   "covr, testthat"

## evaluate   "covr, ggplot2, lattice, rlang, testthat (>= 3.0.0), withr"

## fansi      "unitizer, knitr, rmarkdown"

## fastmap    "testthat (>= 2.1.1)"

## fontawesome "covr, dplyr (>= 1.0.8), knitr (>= 1.31), testthat (>= 3.0.0),\nrsvg"

## fs         "covr, crayon, knitr, pillar (>= 1.0.0), rmarkdown, spelling,\nntestthat (>= 3.0.0), tibble (>= 1.1.0), vctrs (>= 0.3.0), withr"

## glue       "covr, crayon, DBI, dplyr, forcats, ggplot2, knitr, magrittr,\nrmicrobenchmark, R.utils, rmarkdown, rprintf, RSQLite, stringr,\nntestthat (>= 3.0.0), v  
ctrs (>= 0.3.0), waldo (>= 0.3.0), withr"
## highr      "knitr, markdown, testit"

## hms        "crayon, lubridate, pillar (>= 1.1.0), testthat (>= 3.0.0)"

## htmltools  "markdown, testthat, withr, Cairo, ragg, shiny"

## jquerylib  "testthat"

## jsonlite   "httr, vctrs, testthat, knitr, rmarkdown, R.rsp, sf"

## knitr      "markdown (>= 1.3), formatR, testit, digest, rgl (>=\n0.95.1201), codetools, rmarkdown, htmlwidgets (>= 0.7),\nwebshot, tikzDevice (>= 0.10), t  
inytex, reticulate (>= 1.4),\nJuliaCall (>= 0.11.1), magick, png, jpeg, gifski, xml2 (>=\n1.2.0), httr, DBI (>= 0.4-1), showtext, tibble, sass, bslib,\nragg, gridSV  
G, styler (>= 1.2.0), targets (>= 0.6.0)"
## lifecycle  "covr, crayon, knitr, lintr, rmarkdown, testthat (>= 3.0.1),\ntibble, tidyverse, tools, vctrs, withr"

## magrittr   "covr, knitr, rlang, rmarkdown, testthat"

## markdown   "knitr, rmarkdown (>= 2.18), yaml, RCurl"

## memoise    "digest, aws.s3, covr, googleAuthR, googleCloudStorageR, httr,\nntestthat"

## mime       NA

## pillar     "bit64, DBI, debugme, DiagrammeR, dplyr, formattable, ggplot2,\nknitr, lubridate, nanotime, nycflights13, palmerpenguins,\nrmarkdown, scale  
s, stringi, survival, testthat (>= 3.1.1),\ntibble, units (>= 0.7.2), vdiff, withr"

## pkgconfig  "covr, testthat, disposables (>= 1.0.3)"

## prettyunits "codetools, covr, testthat"

## progress   "Rcpp, testthat, withr"

## R6         "testthat, pryr"

## rappdirs   "roxygen2, testthat (>= 3.0.0), covr, withr"

## Rcpp       "tinytest, inline, rbenchmark, pkgKitten (>= 0.1.2)"

## readxl     "covr, knitr, rmarkdown, testthat (>= 3.1.6), withr"

## rematch    "covr, testthat"

## rlang      "cli (>= 3.1.0), covr, crayon, fs, glue, knitr, magrittr,\nmethods, pillar, rmarkdown, stats, testthat (>= 3.0.0), tibble,\nusenethis, vctrs (>= 0.2.3), wit  
hr"
## rmarkdown  "digest, dygraphs, fs, rsconnect, downlit (>= 0.4.0), katex\n(>= 1.4.0), sass (>= 0.4.0), shiny (>= 1.6.0), testthat (>=\n3.0.3), tibble, vctrs, wi  
thr (>= 2.4.2)"
## sass       "testthat, knitr, rmarkdown, withr, shiny, curl"

```

```
## stringi    NA

## stringr    "covr, htmltools, htmlwidgets, knitr, rmarkdown, testthat (>=\n3.0.0)"

## tibble     "bench, bit64, blob, brio, callr, cli, covr, crayon (>=\n1.3.4), DiagrammeR, dplyr, evaluate, formattable, ggplot2,\nhere, hms, htmltools, knitr, lu
brideate, mockr, nycflights13,\npkgbuild, pkgload, purrr, rmarkdown, stringi, testthat (>=\n3.0.2), tidyr, withr"

## tinytex    "testit, rstudioapi"

## utf8       "cli, covr, knitr, rlang, rmarkdown, testthat (>= 3.0.0),\nwithr"

## vctrs      "bit64, covr, crayon, dplyr (>= 0.8.5), generics, knitr,\npillar (>= 1.4.4), pkgdown (>= 2.0.1), rmarkdown, testthat (>=\n3.0.0), tibble (>= 3.1.3),
waldo (>= 0.2.0), withr, xml2,\nzeallot"
## xfun       "testit, parallel, codetools, rstudioapi, tinytex (>= 0.30),\nmime, markdown (>= 1.5), knitr (>= 1.42), htmltools, remotes,\npak, rhub, renv, curl, j
sonlite, magick, yaml, rmarkdown"
## yaml       "RUnit"

## base       "methods"

## boot       "MASS, survival"

## class      NA

## cluster    "MASS, Matrix"

## codetools  NA

## compiler   NA

## datasets   NA

## foreign    NA

## graphics   NA

## grDevices  "KernSmooth"

## grid       NA

## KernSmooth "MASS, carData"

## lattice    "KernSmooth, MASS, latticeExtra, colorspace"

## MASS       "lattice, nlme, nnet, survival"

## Matrix     "expm, MASS"

## methods    "codetools"

## mgcv       "parallel, survival, MASS"

## nlme       "Hmisc, MASS, SASmixed"

## nnet       "MASS"

## parallel   "methods"

## rpart      "survival"

## spatial    "MASS"

## splines    "Matrix, methods"

## stats      "MASS, Matrix, SuppDists, methods, stats4"

## stats4     NA

## survival   NA

## tcltk      NA

## tools      "codetools, methods, xml2, curl, commonmark, knitr, xfun,\nmathjaxr, V8"

## translations NA

## utils      "methods, xml2, commonmark, knitr"
```

```
## Enhances
## base64enc "png"
## bslib NA
## cachem NA
## cellranger NA
## cli NA
## commonmark NA
## cpp11 NA
## crayon NA
## digest NA
## ellipsis NA
## evaluate NA
## fansi NA
## fastmap NA
## fontawesome NA
## fs NA
## glue NA
## highr NA
## hms NA
## htmltools "knitr"
## jquerylib NA
## jsonlite NA
## knitr NA
## lifecycle NA
## magrittr NA
## markdown NA
## memoise NA
## mime NA
## pillar NA
## pkgconfig NA
## prettyunits NA
## progress NA
## R6 NA
## rappdirs NA
## Rcpp NA
## readxl NA
## rematch NA
## rlang "winch"
## rmarkdown NA
## sass NA
## stringi NA
## stringr NA
## tibble NA
## tinytex NA
## utf8 NA
## vctrs NA
## xfun NA
## yaml NA
## base NA
## boot NA
## class NA
## cluster NA
## codetools NA
## compiler NA
## datasets NA
## foreign NA
## graphics NA
## grDevices NA
## grid NA
## KernSmooth NA
## lattice "chron"
## MASS NA
## Matrix "MatrixModels, graph, SparseM, sfsmisc, igraph, maptools, sp,\nspdep"
## methods NA
## mgcv NA
## nlme NA
## nnet NA
## parallel "snow, Rmpi"
## rpart NA
## spatial NA
## splines NA
## stats NA
## stats4 NA
## survival NA
## tcltk NA
## tools NA
## translations NA
## utils NA
```

##	License	License_is_FOSS
## base64enc	"GPL-2 GPL-3"	NA
## bslib	"MIT + file LICENSE"	NA
## cachem	"MIT + file LICENSE"	NA
## cellranger	"MIT + file LICENSE"	NA
## cli	"MIT + file LICENSE"	NA
## commonmark	"BSD_2_clause + file LICENSE"	NA
## cpp11	"MIT + file LICENSE"	NA
## crayon	"MIT + file LICENSE"	NA
## digest	"GPL (>= 2)"	NA
## ellipsis	"MIT + file LICENSE"	NA
## evaluate	"MIT + file LICENSE"	NA
## fansi	"GPL-2 GPL-3"	NA
## fastmap	"MIT + file LICENSE"	NA
## fontawesome	"MIT + file LICENSE"	NA
## fs	"MIT + file LICENSE"	NA
## glue	"MIT + file LICENSE"	NA
## highr	"GPL"	NA
## hms	"MIT + file LICENSE"	NA
## htmltools	"GPL (>= 2)"	NA
## jquerylib	"MIT + file LICENSE"	NA
## jsonlite	"MIT + file LICENSE"	NA
## knitr	"GPL"	NA
## lifecycle	"MIT + file LICENSE"	NA
## magrittr	"MIT + file LICENSE"	NA
## markdown	"GPL-2"	NA
## memoise	"MIT + file LICENSE"	NA
## mime	"GPL"	NA
## pillar	"MIT + file LICENSE"	NA
## pkgconfig	"MIT + file LICENSE"	NA
## prettyunits	"MIT + file LICENSE"	NA
## progress	"MIT + file LICENSE"	NA
## R6	"MIT + file LICENSE"	NA
## rappdirs	"MIT + file LICENSE"	NA
## Rcpp	"GPL (>= 2)"	NA
## readxl	"MIT + file LICENSE"	NA
## rematch	"MIT + file LICENSE"	NA
## rlang	"MIT + file LICENSE"	NA
## rmarkdown	"GPL-3"	NA
## sass	"MIT + file LICENSE"	NA
## stringi	"file LICENSE"	"yes"
## stringr	"MIT + file LICENSE"	NA
## tibble	"MIT + file LICENSE"	NA
## tinytex	"MIT + file LICENSE"	NA
## utf8	"Apache License (== 2.0) file LICENSE"	NA
## vctrs	"MIT + file LICENSE"	NA
## xfun	"MIT + file LICENSE"	NA
## yaml	"BSD_3_clause + file LICENSE"	NA
## base	"Part of R 4.3.0"	NA
## boot	"Unlimited"	NA
## class	"GPL-2 GPL-3"	NA
## cluster	"GPL (>= 2)"	NA
## codetools	"GPL"	NA
## compiler	"Part of R 4.3.0"	NA
## datasets	"Part of R 4.3.0"	NA
## foreign	"GPL (>= 2)"	NA
## graphics	"Part of R 4.3.0"	NA
## grDevices	"Part of R 4.3.0"	NA
## grid	"Part of R 4.3.0"	NA
## KernSmooth	"Unlimited"	NA
## lattice	"GPL (>= 2)"	NA
## MASS	"GPL-2 GPL-3"	NA
## Matrix	"GPL (>= 2) file LICENSE"	NA
## methods	"Part of R 4.3.0"	NA
## mgcv	"GPL (>= 2)"	NA
## nlme	"GPL (>= 2)"	NA
## nnet	"GPL-2 GPL-3"	NA
## parallel	"Part of R 4.3.0"	NA
## rpart	"GPL-2 GPL-3"	NA
## spatial	"GPL-2 GPL-3"	NA
## splines	"Part of R 4.3.0"	NA
## stats	"Part of R 4.3.0"	NA
## stats4	"Part of R 4.3.0"	NA
## survival	"LGPL (>= 2)"	NA
## tcltk	"Part of R 4.3.0"	NA
## tools	"Part of R 4.3.0"	NA
## translations	"Part of R 4.3.0"	NA
## utils	"Part of R 4.3.0"	NA

##	License_restricts_use	OS_type	MD5sum	NeedsCompilation	Built
## base64enc	NA	NA	NA	"yes"	"4.3.0"
## bslib	NA	NA	NA	"no"	"4.3.0"
## cachem	NA	NA	NA	"yes"	"4.3.0"
## cellranger	NA	NA	NA	"no"	"4.3.0"
## cli	NA	NA	NA	"yes"	"4.3.0"
## commonmark	NA	NA	NA	"yes"	"4.3.0"
## cpp11	NA	NA	NA	"no"	"4.3.0"
## crayon	NA	NA	NA	"no"	"4.3.0"
## digest	NA	NA	NA	"yes"	"4.3.0"
## ellipsis	NA	NA	NA	"yes"	"4.3.0"
## evaluate	NA	NA	NA	"no"	"4.3.0"
## fansi	NA	NA	NA	"yes"	"4.3.0"
## fastmap	NA	NA	NA	"yes"	"4.3.0"
## fontawesome	NA	NA	NA	"no"	"4.3.0"
## fs	NA	NA	NA	"yes"	"4.3.0"
## glue	NA	NA	NA	"yes"	"4.3.0"
## highr	NA	NA	NA	"no"	"4.3.0"
## hms	NA	NA	NA	"no"	"4.3.0"
## htmltools	NA	NA	NA	"yes"	"4.3.0"
## jquerylib	NA	NA	NA	"no"	"4.3.0"
## jsonlite	NA	NA	NA	"yes"	"4.3.0"
## knitr	NA	NA	NA	"no"	"4.3.0"
## lifecycle	NA	NA	NA	"no"	"4.3.0"
## magrittr	NA	NA	NA	"yes"	"4.3.0"
## markdown	NA	NA	NA	"no"	"4.3.0"
## memoise	NA	NA	NA	"no"	"4.3.0"
## mime	NA	NA	NA	"yes"	"4.3.0"
## pillar	NA	NA	NA	"no"	"4.3.0"
## pkgconfig	NA	NA	NA	"no"	"4.3.0"
## prettyunits	NA	NA	NA	"no"	"4.3.0"
## progress	NA	NA	NA	"no"	"4.3.0"
## R6	NA	NA	NA	"no"	"4.3.0"
## rappdirs	NA	NA	NA	"yes"	"4.3.0"
## Rcpp	NA	NA	NA	"yes"	"4.3.0"
## readxl	NA	NA	NA	"yes"	"4.3.0"
## rematch	NA	NA	NA	"no"	"4.3.0"
## rlang	NA	NA	NA	"yes"	"4.3.0"
## rmarkdown	NA	NA	NA	"no"	"4.3.0"
## sass	NA	NA	NA	"yes"	"4.3.0"
## stringi	NA	NA	NA	"yes"	"4.3.0"
## stringr	NA	NA	NA	"no"	"4.3.0"
## tibble	NA	NA	NA	"yes"	"4.3.0"
## tinytex	NA	NA	NA	"no"	"4.3.0"
## utf8	NA	NA	NA	"yes"	"4.3.0"
## vctrs	NA	NA	NA	"yes"	"4.3.0"
## xfun	NA	NA	NA	"yes"	"4.3.0"
## yaml	NA	NA	NA	"yes"	"4.3.0"
## base	NA	NA	NA	NA	"4.3.0"
## boot	NA	NA	NA	"no"	"4.3.0"
## class	NA	NA	NA	"yes"	"4.3.0"
## cluster	NA	NA	NA	"yes"	"4.3.0"
## codetools	NA	NA	NA	"no"	"4.3.0"
## compiler	NA	NA	NA	NA	"4.3.0"
## datasets	NA	NA	NA	NA	"4.3.0"
## foreign	NA	NA	NA	"yes"	"4.3.0"
## graphics	NA	NA	NA	"yes"	"4.3.0"
## grDevices	NA	NA	NA	"yes"	"4.3.0"
## grid	NA	NA	NA	"yes"	"4.3.0"
## KernSmooth	NA	NA	NA	"yes"	"4.3.0"
## lattice	NA	NA	NA	"yes"	"4.3.0"
## MASS	NA	NA	NA	"yes"	"4.3.0"
## Matrix	NA	NA	NA	"yes"	"4.3.0"
## methods	NA	NA	NA	"yes"	"4.3.0"
## mgcv	NA	NA	NA	"yes"	"4.3.0"
## nlme	NA	NA	NA	"yes"	"4.3.0"
## nnet	NA	NA	NA	"yes"	"4.3.0"
## parallel	NA	NA	NA	"yes"	"4.3.0"
## rpart	NA	NA	NA	"yes"	"4.3.0"
## spatial	NA	NA	NA	"yes"	"4.3.0"
## splines	NA	NA	NA	"yes"	"4.3.0"
## stats	NA	NA	NA	"yes"	"4.3.0"
## stats4	NA	NA	NA	NA	"4.3.0"
## survival	NA	NA	NA	"yes"	"4.3.0"
## tcltk	NA	NA	NA	"yes"	"4.3.0"
## tools	NA	NA	NA	"yes"	"4.3.0"
## translations	NA	NA	NA	NA	"4.3.0"
## utils	NA	NA	NA	"yes"	"4.3.0"

```
search()
```

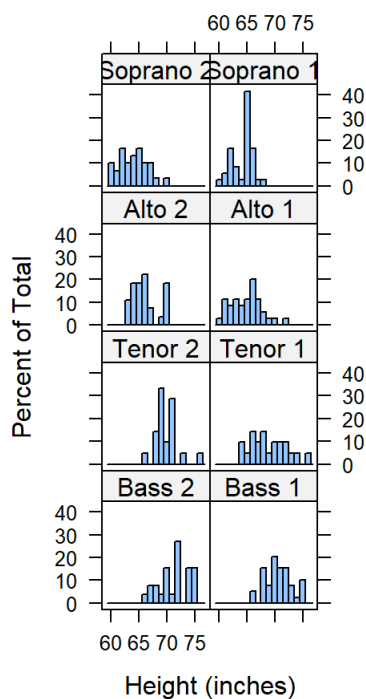
```
## [1] ".GlobalEnv"      "package:stats"    "package:graphics"
## [4] "package:grDevices" "package:utils"    "package:datasets"
## [7] "package:methods"  "Autoloads"        "package:base"
```

```
require(lattice)
```

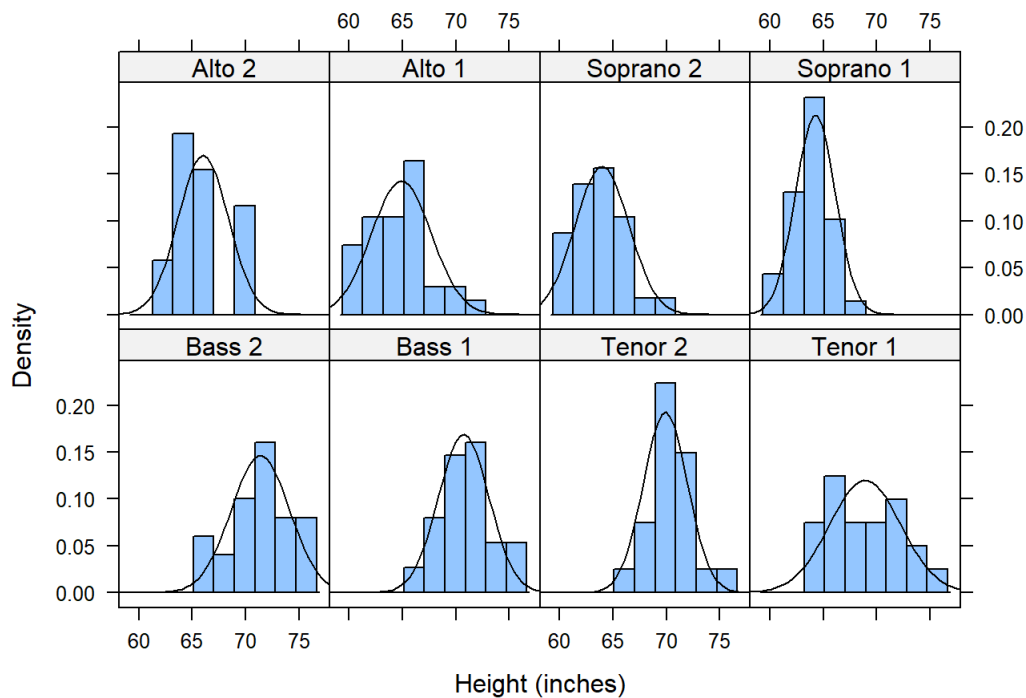
```
## Loading required package: lattice
```

```
example(histogram)
```

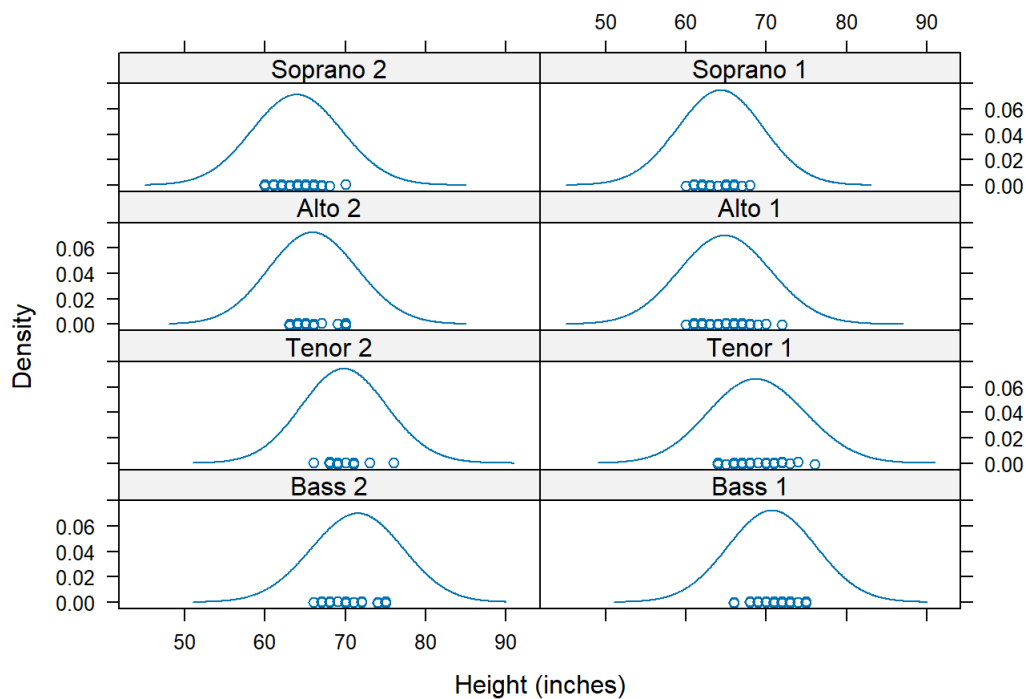
```
##
## hstgrm> require(stats)
##
## hstgrm> histogram( ~ height | voice.part, data = singer, nint = 17,
## hstgrm+           endpoints = c(59.5, 76.5), layout = c(2,4), aspect = 1,
## hstgrm+           xlab = "Height (inches)")
```



```
##
## hstgrm> histogram( ~ height | voice.part, data = singer,
## hstgrm+           xlab = "Height (inches)", type = "density",
## hstgrm+           panel = function(x, ...) {
## hstgrm+             panel.histogram(x, ...)
## hstgrm+             panel.mathdensity(dmath = dnorm, col = "black",
## hstgrm+               args = list(mean=mean(x),sd=sd(x)))
## hstgrm+           } )
```



```
##
## hstgrm> densityplot( ~ height | voice.part, data = singer, layout = c(2, 4),
## hstgrm+               xlab = "Height (inches)", bw = 5)
```



```
ls()
```

```
## character(0)
```

```
x=c(1,4,5,4,6,8)
x
```

```
## [1] 1 4 5 4 6 8
```

```
names=c("Book", "Pen", "Bat", "Biscuit", "Lake")
names
```

```
## [1] "Book" "Pen" "Bat" "Biscuit" "Lake"
```

```
example=c(rep(1,14),rep(7,22))
example
```

```
## [1] 1 1 1 1 1 1 1 1 1 1 1 1 7 7 7 7 7 7 7 7 7 7 7 7 7 7
```

```
res=seq(9,19,1)
res
```

```
## [1] 9 10 11 12 13 14 15 16 17 18 19
```

```
data(trees)
names(trees)
```

```
## [1] "Girth" "Height" "Volume"
```

```
dim(trees)
```

```
## [1] 31 3
```

```
str(trees)
```

```
## 'data.frame': 31 obs. of 3 variables:
## $ Girth : num 8.3 8.6 8.8 10.5 10.7 10.8 11 11 11.1 11.2 ...
## $ Height: num 70 65 63 72 81 83 66 75 80 75 ...
## $ Volume: num 10.3 10.3 10.2 16.4 18.8 19.7 15.6 18.2 22.6 19.9 ...
```

View(trees)

```
train=read.csv(file.choose())
```

```
names(train)
```

```
## [1] "id" "vendor_id" "passenger_count"
## [4] "pickup_longitude" "pickup_latitude" "dropoff_longitude"
## [7] "dropoff_latitude" "store_and_fwd_flag" "trip_duration"
```

```
x=train$vendor_id
mean(x,na.rm=T)
```

```
## [1] 1.534503
```

```
median(x,na.rm=T)
```

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

```
summary(x)
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1.000	1.000	2.000	1.535	2.000	2.000

```
sd(x,na.rm=T)
```

```
## [1] 0.4988084
```

```
var(x,na.rm=T)
```

```
## [1] 0.2488098
```

```
quantile(x, probs=seq(0, 1, 0.5), na.rm=T)
```



```
## 0% 50% 100%
## 1 2 2
```

```
add=train$passenger_count
sum(add)
```

```
## [1] 1745229
```

stem()

```
stem(train$pickup_latitude)
```

[illegible]

table()

```
table(train$store_and_fwd_flag)
```

```
##
##      N      Y
## 1042766  5809
```

```
table(train$store_and_fwd_flag,train$passenger_count)
```

```
##
##      0      1      2      3      4      5      6      7      8      9
## N    38 738485 150235 42840 20214 56249 34700      3      1      1
## Y      1 4647  843  193  122      3      0      0      0      0
```

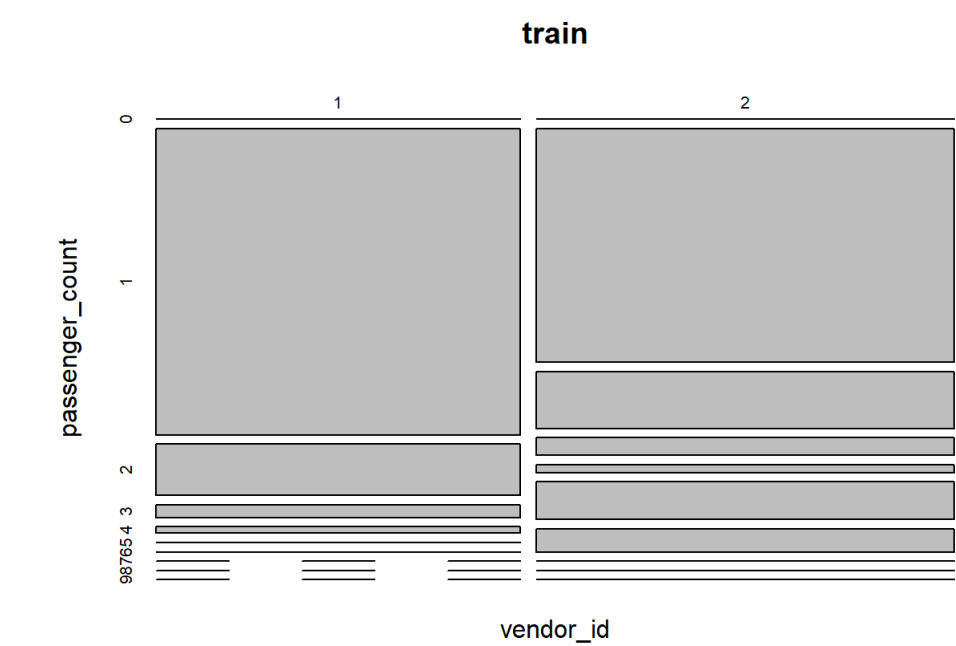
```
xtabs(~store_and_fwd_flag,data=train)
```

```
## store_and_fwd_flag
##      N      Y
## 1042766 5809
```

```
xtabs(~passenger_count+store_and_fwd_flag,data=train)
```

```
##           store_and_fwd_flag
## passenger_count    N      Y
##      0      38      1
##      1 738485 4647
##      2 150235 843
##      3 42840 193
##      4 20214 122
##      5 56249 3
##      6 34700 0
##      7 3 0
##      8 1 0
##      9 1 0
```

```
mosaicplot(~vendor_id+passenger_count,data=train)
```



```
table(train$passenger_count)
```

```
##
##  0  1  2  3  4  5  6  7  8  9
## 39 74 31 32 15 10 78 43 0 33 20 33 56 252 34 700 3 1 1
```

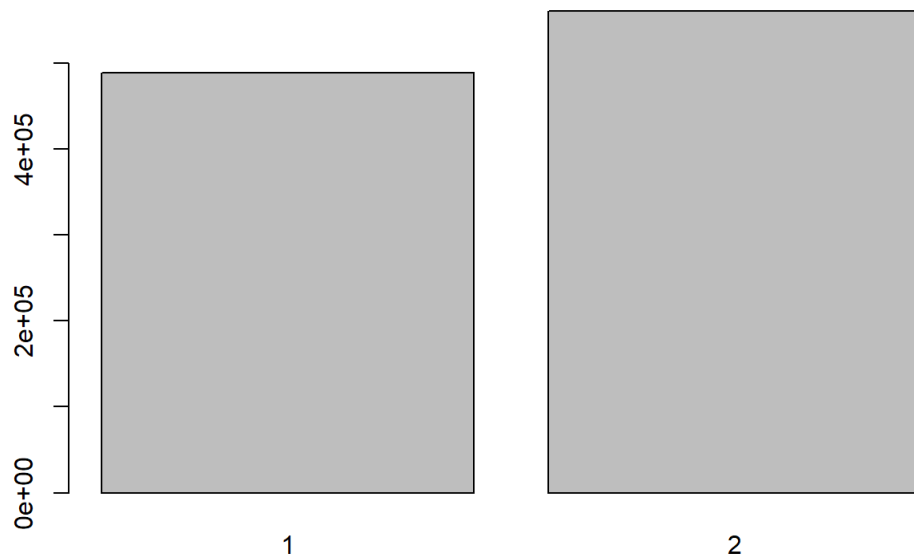
```
xtabs(~passenger_count,data=train)
```

```
## passenger_count
##  0  1  2  3  4  5  6  7  8  9
## 39 74 31 32 15 10 78 43 0 33 20 33 56 252 34 700 3 1 1
```

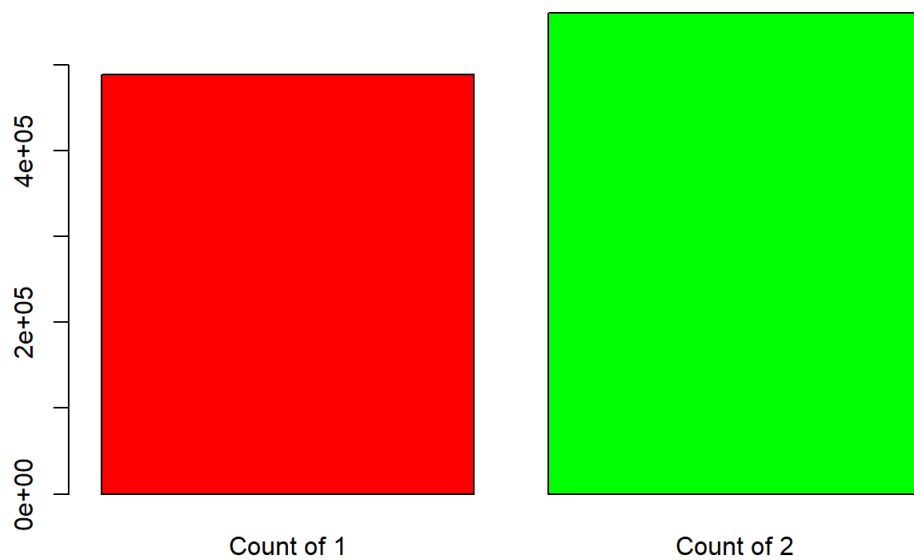
```
levels=cut(train$passenger_count,breaks=seq(0,60,10))
xtabs(~levels)
```

```
## levels
## (0,10] (10,20] (20,30] (30,40] (40,50] (50,60]
## 10 48 536 0 0 0 0 0 0
```

```
barplot()
barplot(table(train$vendor_id))
```



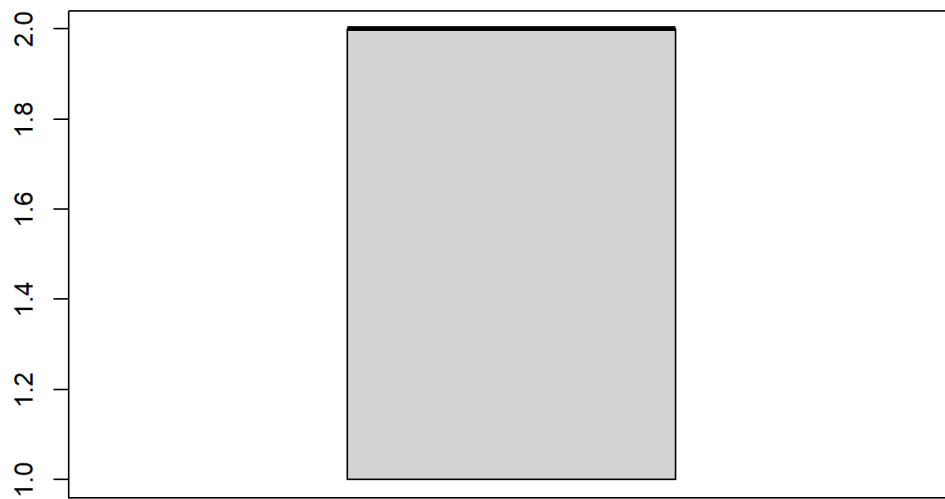
```
barplot(table(train$vendor_id),col=c("red","green"),names=c("Count of 1","Count of 2"))
```



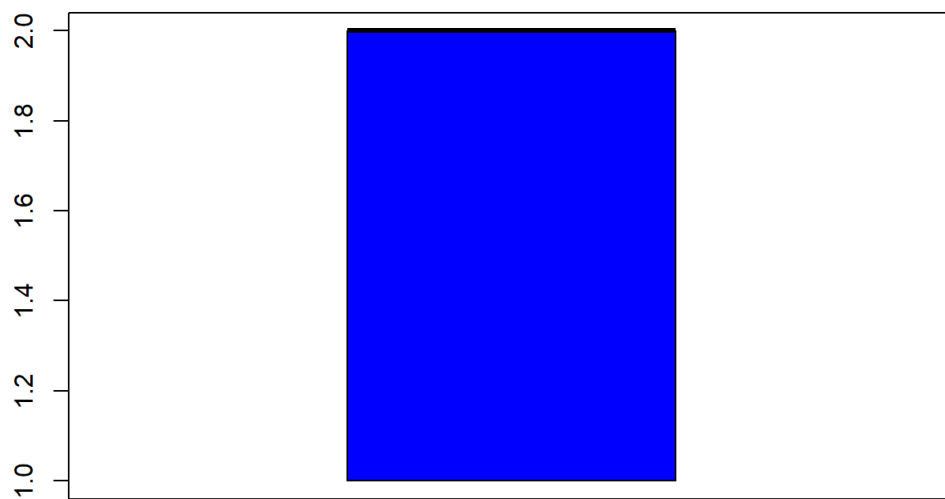
```
data(train)
```

```
## Warning in data(train): data set 'train' not found
```

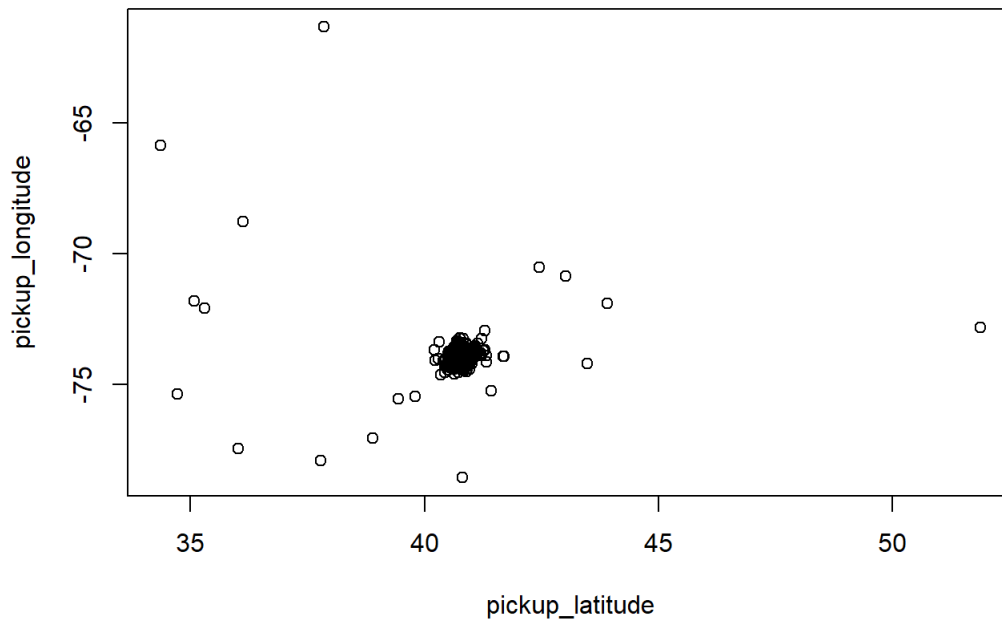
```
boxplot(train$vendor_id)
```



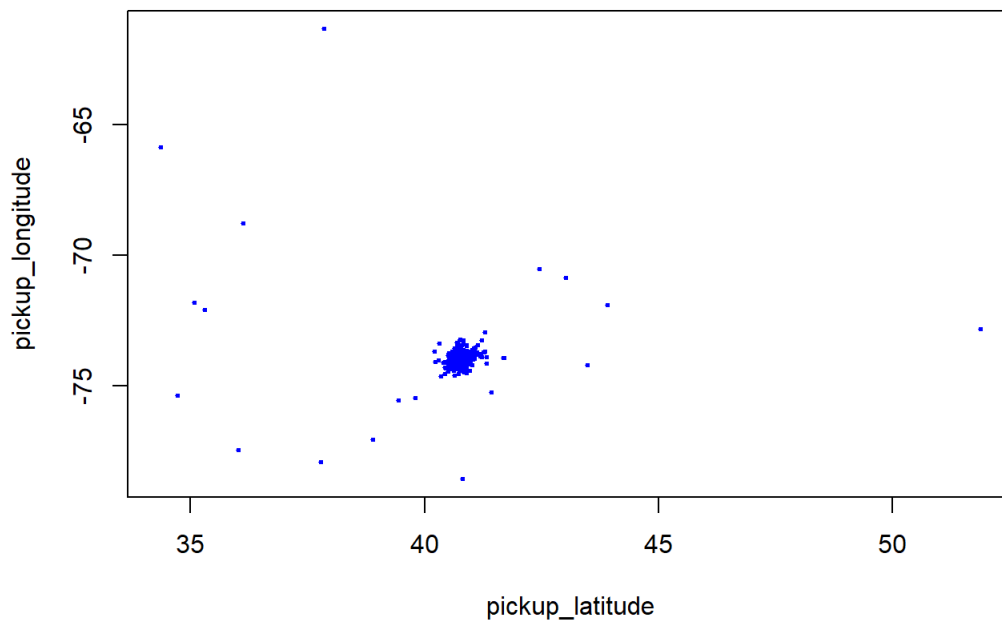
```
boxplot(train$vendor_id,col="blue")
```



```
plot(pickup_longitude~pickup_latitude,data=train)
```



```
plot(pickup_longitude~pickup_latitude,data=train,cex=0.5,pch=20,col="blue")
```



```
sample(c("rows,columns"),size=1)
```

```
## [1] "rows,columns"
```

```
sample(c("rows","columns"),size=20,replace=T)
```

```
## [1] "columns" "rows" "rows" "columns" "rows" "rows" "columns"
## [8] "columns" "columns" "rows" "rows" "columns" "rows" "rows"
## [15] "rows" "columns" "rows" "rows" "rows" "rows"
```

```
sample(c(0,1),size=20,replace=T)
```

```
## [1] 0 0 1 0 0 0 1 0 1 0 1 1 0 0 0 0 0 0 0 0
```

```
sample(c("rows","columns"),2,replace=T)
```

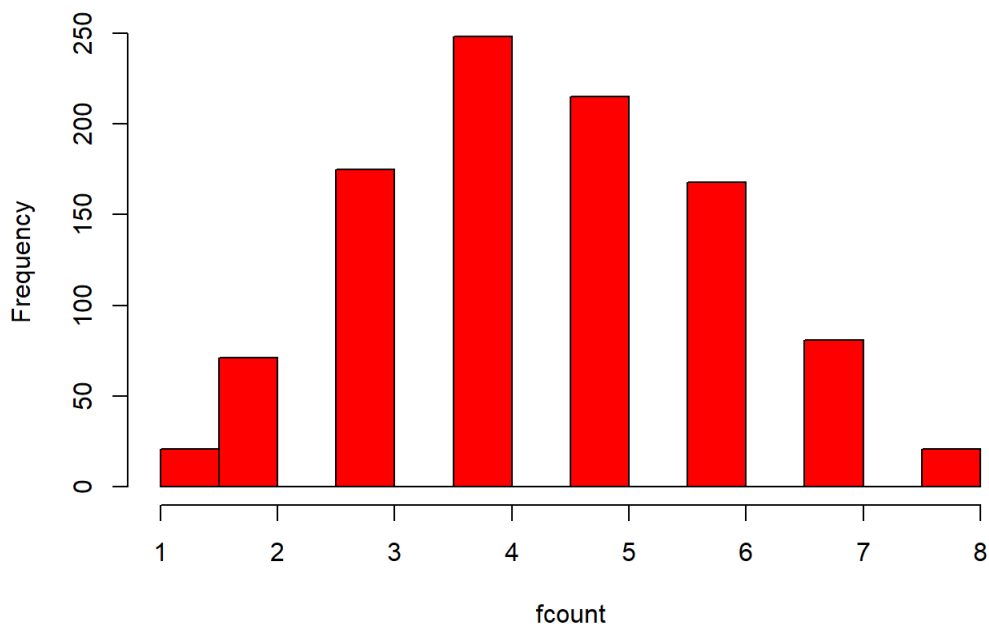
```
## [1] "columns" "columns"
```

```
replicate(10,sample(c("rows","columns"),2,replace=T))
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7]  
## [1,] "rows" "columns" "rows" "columns" "rows" "columns" "columns"  
## [2,] "rows" "rows" "columns" "rows" "columns" "rows" "columns"  
##      [,8] [,9] [,10]  
## [1,] "columns" "rows" "rows"  
## [2,] "columns" "columns" "rows"
```

```
fcount=replicate(1000,sum(sample(c(0,1),10,rep=T,prob=c(.5,.4))))  
hist(fcount,col="red")
```

Histogram of fcount



```
dbinom(1,6,0.4) #probability of 1 heads in 6 flips
```

```
## [1] 0.186624
```

```
dbinom(0:6,6,0.4)#full probability dist for 6 flips
```

```
## [1] 0.046656 0.186624 0.311040 0.276480 0.138240 0.036864 0.004096
```

```
sum(dbinom(0:3,6,0.4))#probability of 3 or fewer heads in 6 flips
```

```
## [1] 0.8208
```

```
pbinom(3,6,0.4)
```

```
## [1] 0.8208
```

```
passenger_count=replicate(1000,sum(sample(c("H","T"),5,rep=T)== "H"))  
table(passenger_count)/1000
```

```
## passenger_count  
## 0 1 2 3 4 5  
## 0.027 0.149 0.335 0.299 0.169 0.021
```

```
table(rbinom(1000,5,0.5))/1000
```

```
##
## 0 1 2 3 4 5
## 0.025 0.139 0.302 0.325 0.167 0.042
```

```
qbinom(seq(0,2,0.3),40,0.1) #quantiles in binom distirbution
```

```
## Warning in qbinom(seq(0, 2, 0.3), 40, 0.1): NaNs produced
```

```
## [1] 0 3 4 6 NaN NaN NaN
```

```
pchisq(4.13,df=4,lower.tail = F) #gives P-value associated with X-squared stat 4.13 when df=4
```

```
## [1] 0.3886981
```

```
qchisq(c(0.001,0.005,0.99),2,lower.tail = F) #gives critical values
```

```
## [1] 13.81551056 10.59663473 0.02010067
```

```
values=c(35,65,76,34,98,87)
probability=c(0.13,0.13,0.14,0.16,0.24,0.20)
chisq.test(values,p=probability) #to find the chi-square test
```

```
##
## Chi-squared test for given probabilities
##
## data: values
## X-squared = 30.992, df = 5, p-value = 9.401e-06
```

```
t=table(train$vendor_id,train$store_and_fwd_flag)#to add both row/columns
addmargins(t)
```

```
##
##      N    Y   Sum
## 1  482300 5809 488109
## 2  560466    0 560466
## Sum 1042766 5809 1048575
```

```
addmargins(t,1)#to add oly column totals
```

```
##
##      N    Y
## 1  482300 5809
## 2  560466    0
## Sum 1042766 5809
```

```
addmargins(t,2)#to add oly row totals
```

```
##
##      N    Y   Sum
## 1  482300 5809 488109
## 2  560466    0 560466
```

```
my_data <- data.frame(
  gender = c("Male", "Male", "Female", "Female", "Female", "Male", "Male"),
  age = c(25, 30, 35, 40, 45, 50, 55)
)

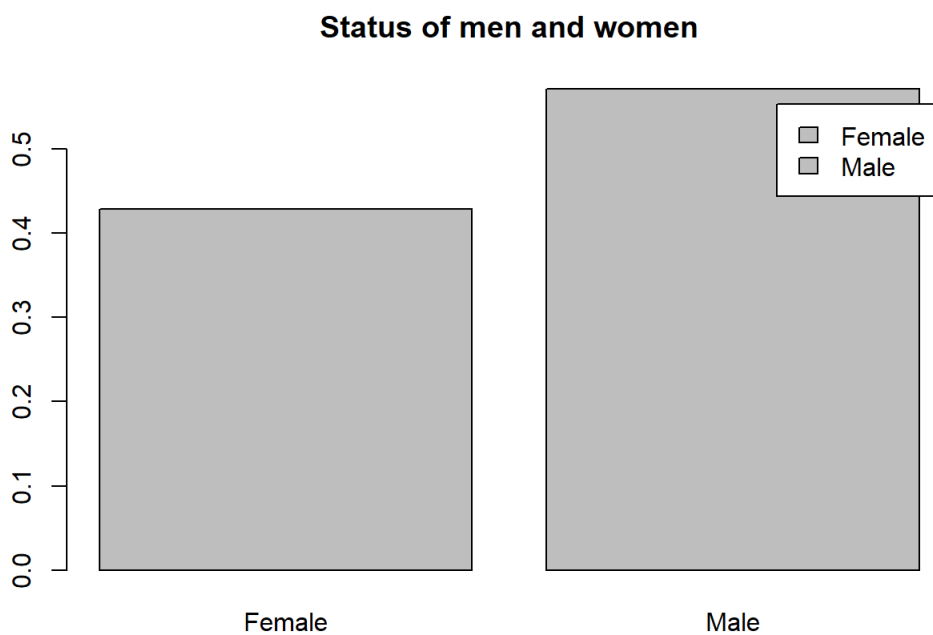
summary(my_data)
```

```
## gender age
## Length:7 Min. :25.0
## Class :character 1st Qu.:32.5
## Mode :character Median :40.0
## Mean :40.0
## 3rd Qu.:47.5
## Max. :55.0
```

```
gender_table <- table(my_data$gender)
gender_prop <- prop.table(gender_table)
gender_prop
```

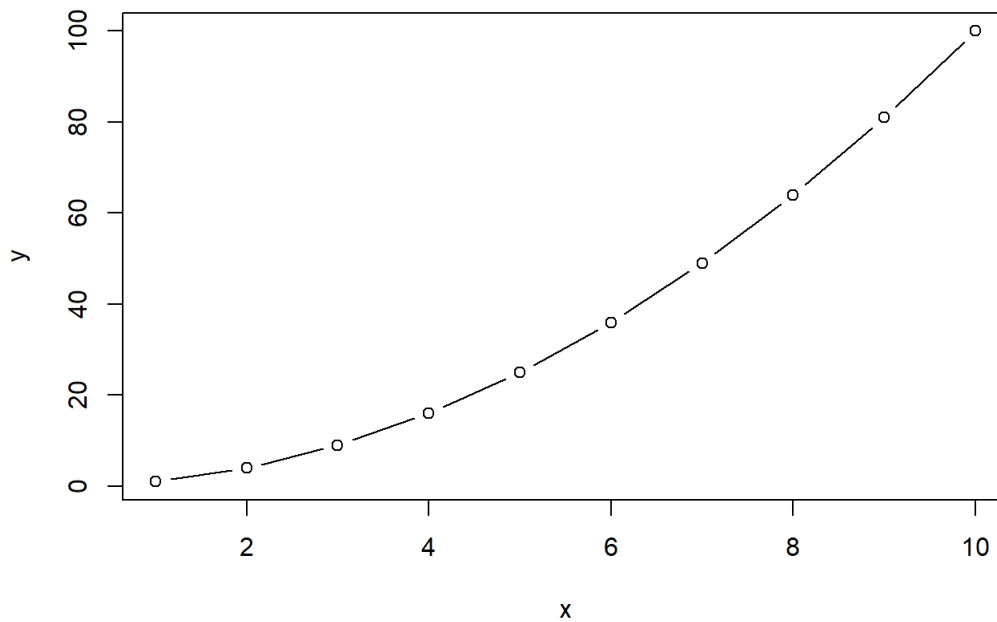
```
##
## Female Male
## 0.4285714 0.5714286
```

```
barplot(gender_prop,legend=T,beside=T,main="Status of men and women")
```



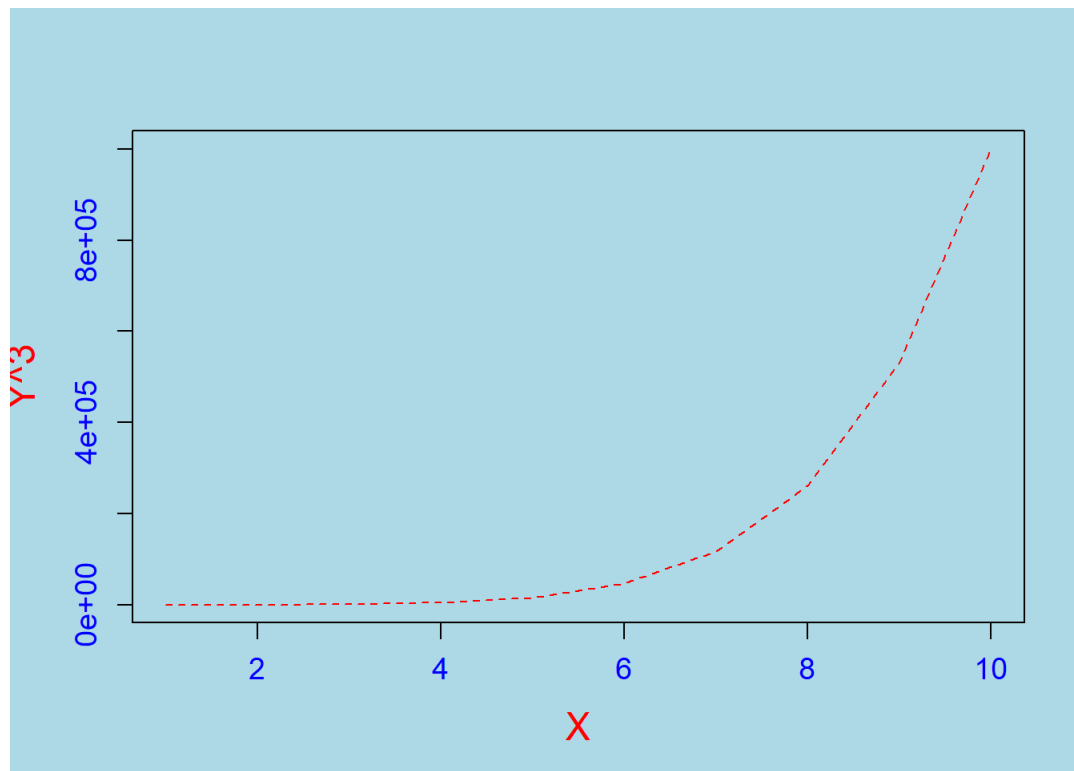
```
x <- 1:10
y <- x^2
plot(x, y, type = "b", main = "Plot")
```


Plot



```
par(
  mar = c(5, 4, 4, 2), # Set the margin size
  col.axis = "blue",    # Set the color of the axes
  col.lab = "red",       # Set the color of the labels
  cex.axis = 1.2,        # Set the size of the axis text
  cex.lab = 1.5,         # Set the size of the label text
  pch = 16,             # Set the plotting symbol
  lty = 2,              # Set the line type
  bg = "lightblue"      # Set the background color
)

plot(x, y^3, type = "l", col = "red", xlab = "X", ylab = "Y^3")
```



```
contingency_table <- matrix(c(10, 20, 30, 40), nrow = 2)
rownames(contingency_table) <- c("Group A", "Group B")
colnames(contingency_table) <- c("Success", "Failure")
contingency_table
```

```
##      Success Failure
## Group A    10     30
## Group B    20     40
```

```
result <- fisher.test(contingency_table)
```

```
result
```

```
##
## Fisher's Exact Test for Count Data
##
## data: contingency_table
## p-value = 0.5045
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
##  0.2420054 1.7643388
## sample estimates:
## odds ratio
## 0.6693434
```

```
# Calculate the PMF of the Poisson distribution
lambda <- 2 # Mean parameter for the Poisson distribution
x <- 0:10 # Values at which to evaluate the PMF

pmf <- dpois(x, lambda)

# Display the PMF
print(pmf)
```

```
## [1] 1.353353e-01 2.706706e-01 2.706706e-01 1.804470e-01 9.022352e-02
## [6] 3.608941e-02 1.202980e-02 3.437087e-03 8.592716e-04 1.909493e-04
## [11] 3.818985e-05
```

```
# Calculate the CDF of the Poisson distribution
lambda <- 2 # Mean parameter for the Poisson distribution
x <- 0:10 # Values at which to evaluate the CDF

cdf <- ppois(x, lambda)

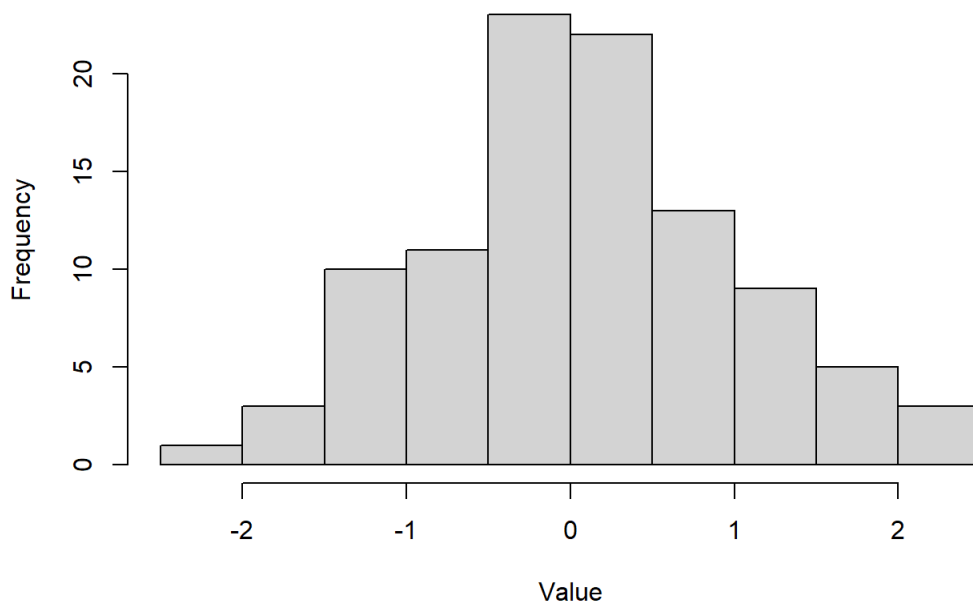
# Display the CDF
print(cdf)
```

```
## [1] 0.1353353 0.4060058 0.6766764 0.8571235 0.9473470 0.9834364 0.9954662
## [8] 0.9989033 0.9997626 0.9999535 0.9999917
```

```
# Generate a random sample from a normal distribution
set.seed(123) # Set seed for reproducibility
sample <- rnorm(100, mean = 0, sd = 1)

# Create a histogram of the sample
hist(sample, main = "Histogram of Random Sample", xlab = "Value")
```

Histogram of Random Sample



```
# Calculate the probability of values below a certain threshold
```

```
threshold <- 1.5
```

```
probability <- pnorm(threshold, mean = 0, sd = 1)
```

```
# Calculate the value corresponding to a given quantile
```

```
quantile <- 0.75
```

```
value <- qnorm(quantile, mean = 0, sd = 1)
```

```
# Calculate the probability density at a given value
```

```
density <- dnorm(value, mean = 0, sd = 1)
```

```
# Display the results
```

```
print(probability)
```

```
## [1] 0.9331928
```

```
print(value)
```

```
## [1] 0.6744898
```

```
print(density)
```

```
## [1] 0.3177766
```

```
# Generate a random sample from a normal distribution
```

```
set.seed(123) # Set seed for reproducibility
```

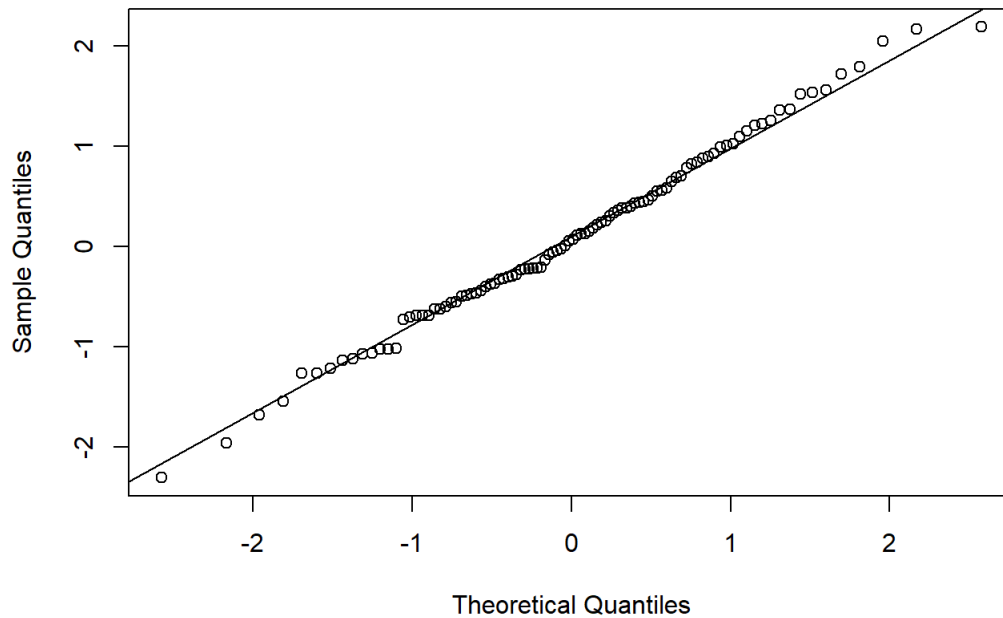
```
sample <- rnorm(100, mean = 0, sd = 1)
```

```
# Create a QQ plot
```

```
qqnorm(sample, main = "QQ Plot")
```

```
qqline(sample)
```

QQ Plot



```
# Calculate the probability of values below a certain threshold
```

```
threshold <- 1.5
```

```
probability <- pnorm(threshold, mean = 0, sd = 1)
```

```
# Calculate the value corresponding to a given quantile
```

```
quantile <- 0.75
```

```
value <- qnorm(quantile, mean = 0, sd = 1)
```

```
# Calculate the probability density at a given value
```

```
density <- dnorm(value, mean = 0, sd = 1)
```

```
# Display the results
```

```
print(probability)
```

```
## [1] 0.9331928
```

```
print(value)
```

```
## [1] 0.6744898
```

```
print(density)
```

```
## [1] 0.3177766
```

```
# Generate two independent samples
```

```
set.seed(123) # Set seed for reproducibility
```

```
sample1 <- rnorm(50, mean = 5, sd = 2)
```

```
sample2 <- rnorm(50, mean = 7, sd = 2)
```

```
# Perform a two-sample t-test
```

```
result <- t.test(sample1, sample2)
```

```
# Display the t-test result
```

```
print(result)
```

```
##
## Welch Two Sample t-test
##
## data: sample1 and sample2
## t = -6.0718, df = 97.951, p-value = 2.406e-08
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -2.950897 -1.497122
## sample estimates:
## mean of x mean of y
## 5.068807 7.292817
```

```
# Perform a power analysis for a two-sample t-test
effect_size <- 0.5 # Expected effect size
n <- 50 # Sample size per group
alpha <- 0.05 # Significance level

power <- power.t.test(n = n, delta = effect_size, sd = 2, sig.level = alpha, type = "two.sample")

# Display the power analysis result
print(power)
```

```
##
## Two-sample t test power calculation
##
## n = 50
## delta = 0.5
## sd = 2
## sig.level = 0.05
## power = 0.2350874
## alternative = two.sided
##
## NOTE: n is number in *each* group
```

```
# Create a linear regression model
set.seed(123) # Set seed for reproducibility
x <- 1:10
y <- 2*x + rnorm(10)
model <- lm(y ~ x)

# Generate new values for prediction
new_x <- 11:15

# Make predictions using the model
predictions <- predict(model, newdata = data.frame(x = new_x))

# Display the predictions
print(predictions)
```

```
## 1 2 3 4 5
## 21.62378 23.54181 25.45984 27.37787 29.29590
```

```
# Example data
x <- c(1, 2, 3, 4, 5)
y <- c(1.5, 3.5, 6, 9, 13)

# Define a nonlinear function to fit
model <- function(x, a, b, c) {
  a * x^2 + b * x + c
}

# Fit the nonlinear model using nls()
fit <- nls(y ~ model(x, a, b, c), start = list(a = 1, b = 1, c = 1))

# Print the model summary
summary(fit)
```

```
##  
## Formula: y ~ model(x, a, b, c)  
##  
## Parameters:  
##   Estimate Std. Error t value Pr(>|t|)  
## a  0.32143   0.03194  10.062  0.00973 **  
## b  0.92143   0.19535   4.717  0.04213 *  
## c  0.30000   0.25635   1.170  0.36246  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 0.1195 on 2 degrees of freedom  
##  
## Number of iterations to convergence: 1  
## Achieved convergence tolerance: 1.145e-07
```

```
# Get the residuals from the fitted model  
residuals <- residuals(fit)
```

```
# Print the residuals  
print(residuals)
```

```
## [1] -0.04285714  0.07142857  0.04285714 -0.12857143  0.05714286  
## attr(,"label")  
## [1] "Residuals"
```

```
# Example data  
x <- c(1, 2, 3, 4, 5)  
  
# Calculate the cumulative sum using cumsum()  
cumulative_sum <- cumsum(x)  
  
# Print the cumulative sum  
print(cumulative_sum)
```

```
## [1]  1  3  6 10 15
```

```
# Calculate the empirical cumulative distribution function (ECDF) using ecdf()  
ecdf_func <- ecdf(x)  
  
# Evaluate the ECDF at specific values  
probability_2 <- ecdf_func(2)  
probability_4 <- ecdf_func(4)  
  
# Print the probabilities  
print(probability_2)
```

```
## [1] 0.4
```

```
print(probability_4)
```

```
## [1] 0.8
```

```
class(train)
```

```
## [1] "data.frame"
```

```
x <- c(5, 2, 9, 1, 7)  
sorted_x <- sort(x)  
print(sorted_x)
```

```
## [1] 1 2 5 7 9
```

```
x <- c(5, 2, 9, 1, 7)  
ranked_x <- rank(x)  
print(ranked_x)
```

```
## [1] 3 2 5 1 4
```

```
# Create a data frame
```

```
df <- data.frame(x = c(1, 2, 3), y = c("A", "B", "C"))
```

```
# Display the row names
```

```
print(row.names(df))
```

```
## [1] "1" "2" "3"
```

1. be↵