

hw1-r-utkarsha

Home Work Assignment 1

Checking version of the R

```
R.version
```

```
platform      x86_64-pc-linux-gnu
arch          x86_64
os            linux-gnu
system        x86_64, linux-gnu
status
major         4
minor         3.1
year          2023
month         06
day           16
svn rev       84548
language      R
version.string R version 4.3.1 (2023-06-16)
nickname      Beagle Scouts
```

Packages in R

Packages are **collections of functions, data sets, and documentation bundled together to extend the functionality** of the base R language. Packages are essential for adding specific capabilities to R

DMwR2 : Data Mining with R

Package focuses on **methods and tools for dealing with imbalanced data sets**, which are datasets where one class (or group) of observations significantly outnumbers the other

```
install.packages("DMwR2")
```

Installing DMwR2 Package :

```
install.packages("DMwR2")
```

the **help()** function is used to access documentation and information about functions, data sets, packages, and other objects in the R environment.

```
help(package="DMwR2")
```

To see the list of pre-loaded data in base R packages, type the function **data()**:

```
data() #loads the information of the datasets
```

Using dataset provided in DMwR2 by referencing its name directly

```
algae <- data(algae) # load algae dataset
```

Warning in data(algae): data set 'algae' not found

```
algae#viewing the data
```

```
[1] "algae"
```

To view the first few rows of your dataset, you can use the **head()** function.

```
head(algae) #displays first 6 rows of the data
```

```
[1] "algae"
```

Similarly, you can use the **tail()** function to view the last few rows of your dataset.

```
tail(algae) # displays last 6 rows of the data
```

```
[1] "algae"
```

```
tail(algae, n = 10) # Display the last 10 rows
```

```
[1] "algae"
```

To get summary statistics for your dataset, including mean, median, minimum, maximum, and quartiles for numeric columns, you can use the **summary()** function.

```
summary(algae)
```

```
Length      Class      Mode
      1 character character
```

```
{#{r message=FALSE, warning=FALSE} manyNAs(algae) # find rows with too many
NAs
```

Libraries and packages in R:

library() function without any arguments to list all the packages that are currently loaded in your R session.

```
library()
```

(.packages()) function is used to list the names packages whose functions and objects are readily available for use in your R code.

```
(.packages())
```

```
[1] "stats"      "graphics"  "grDevices" "utils"      "datasets"  "methods"
[7] "base"
```

library(packagename) function loads and attaches a specific package, making its functions and datasets available for use in your R session

```
library(ggplot2) #attaching the required package in the current session
(.packages())
```

```
[1] "ggplot2"    "stats"      "graphics"   "grDevices"  "utils"      "datasets"
[7] "methods"   "base"
```

Detaching a package means that its functions and objects are no longer accessible in your current R session. This can be **useful when you want to clean up your workspace or avoid conflicts between functions or objects with the same name in different packages.**

```
detach("package:ggplot2", unload=TRUE)
(.packages())
```

```
[1] "stats"      "graphics"   "grDevices"  "utils"      "datasets"   "methods"
[7] "base"
```

installed.packages() function is used to retrieve a list of all packages that are currently installed on your system.

```
installed.packages()
```

	Package	LibPath
abind	"abind"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
askpass	"askpass"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
base64enc	"base64enc"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
bit	"bit"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
bit64	"bit64"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
bitops	"bitops"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
bslib	"bslib"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
cachem	"cachem"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
caTools	"caTools"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
cli	"cli"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
clipr	"clipr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
colorspace	"colorspace"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
cpp11	"cpp11"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
crayon	"crayon"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
credentials	"credentials"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"

curl	"curl"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
DBI	"DBI"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
desc	"desc"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
digest	"digest"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
DMwR2	"DMwR2"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
dplyr	"dplyr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
ellipsis	"ellipsis"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
evaluate	"evaluate"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
fansi	"fansi"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
farver	"farver"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
fastmap	"fastmap"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
fontawesome	"fontawesome"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
fs	"fs"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
generics	"generics"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
gert	"gert"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
ggplot2	"ggplot2"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
gh	"gh"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
gitcreds	"gitcreds"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
glue	"glue"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
gplots	"gplots"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
gtable	"gtable"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
gtools	"gtools"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
highr	"highr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
hms	"hms"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
htmltools	"htmltools"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
httr2	"httr2"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
ini	"ini"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
isoband	"isoband"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
jquerylib	"jquerylib"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
jsonlite	"jsonlite"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
KernSmooth	"KernSmooth"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
knitr	"knitr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
labeling	"labeling"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
lifecycle	"lifecycle"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
magrittr	"magrittr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
Matrix	"Matrix"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
memoise	"memoise"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
mgcv	"mgcv"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
mime	"mime"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
munsell	"munsell"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
nlme	"nlme"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
openssl	"openssl"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
palmerpenguins	"palmerpenguins"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"

pillar	"pillar"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
pkgconfig	"pkgconfig"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
prettyunits	"prettyunits"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
progress	"progress"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
purrr	"purrr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
quantmod	"quantmod"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
R6	"R6"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
rappdirs	"rappdirs"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
RColorBrewer	"RColorBrewer"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
readr	"readr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
rlang	"rlang"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
rmarkdown	"rmarkdown"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
ROCR	"ROCR"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
rprojroot	"rprojroot"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
rstudioapi	"rstudioapi"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
sass	"sass"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
scales	"scales"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
spatial	"spatial"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
stringi	"stringi"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
stringr	"stringr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
survival	"survival"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
sys	"sys"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
tibble	"tibble"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
tidyselect	"tidyselect"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
tinytex	"tinytex"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
TTR	"TTR"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
tzdb	"tzdb"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
usethis	"usethis"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
utf8	"utf8"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
vctrs	"vctrs"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
viridisLite	"viridisLite"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
vroom	"vroom"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
whisker	"whisker"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
withr	"withr"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
xfun	"xfun"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
xts	"xts"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
yaml	"yaml"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
zip	"zip"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
zoo	"zoo"	"/cloud/lib/x86_64-pc-linux-gnu-library/4.3"
base	"base"	"/opt/R/4.3.1/lib/R/library"
boot	"boot"	"/opt/R/4.3.1/lib/R/library"
class	"class"	"/opt/R/4.3.1/lib/R/library"
cluster	"cluster"	"/opt/R/4.3.1/lib/R/library"

codetools	"codetools"	"/opt/R/4.3.1/lib/R/library"
compiler	"compiler"	"/opt/R/4.3.1/lib/R/library"
datasets	"datasets"	"/opt/R/4.3.1/lib/R/library"
foreign	"foreign"	"/opt/R/4.3.1/lib/R/library"
graphics	"graphics"	"/opt/R/4.3.1/lib/R/library"
grDevices	"grDevices"	"/opt/R/4.3.1/lib/R/library"
grid	"grid"	"/opt/R/4.3.1/lib/R/library"
KernSmooth	"KernSmooth"	"/opt/R/4.3.1/lib/R/library"
lattice	"lattice"	"/opt/R/4.3.1/lib/R/library"
MASS	"MASS"	"/opt/R/4.3.1/lib/R/library"
Matrix	"Matrix"	"/opt/R/4.3.1/lib/R/library"
methods	"methods"	"/opt/R/4.3.1/lib/R/library"
mgcv	"mgcv"	"/opt/R/4.3.1/lib/R/library"
nlme	"nlme"	"/opt/R/4.3.1/lib/R/library"
nnet	"nnet"	"/opt/R/4.3.1/lib/R/library"
parallel	"parallel"	"/opt/R/4.3.1/lib/R/library"
rpart	"rpart"	"/opt/R/4.3.1/lib/R/library"
spatial	"spatial"	"/opt/R/4.3.1/lib/R/library"
splines	"splines"	"/opt/R/4.3.1/lib/R/library"
stats	"stats"	"/opt/R/4.3.1/lib/R/library"
stats4	"stats4"	"/opt/R/4.3.1/lib/R/library"
survival	"survival"	"/opt/R/4.3.1/lib/R/library"
tcltk	"tcltk"	"/opt/R/4.3.1/lib/R/library"
tools	"tools"	"/opt/R/4.3.1/lib/R/library"
utils	"utils"	"/opt/R/4.3.1/lib/R/library"
	Version	Priority
abind	"1.4-5"	NA
askpass	"1.1"	NA
base64enc	"0.1-3"	NA
bit	"4.0.5"	NA
bit64	"4.0.5"	NA
bitops	"1.0-7"	NA
bslib	"0.5.1"	NA
cachem	"1.0.8"	NA
caTools	"1.18.2"	NA
cli	"3.6.1"	NA
clipr	"0.8.0"	NA
colorspace	"2.1-0"	NA
cpp11	"0.4.6"	NA
crayon	"1.5.2"	NA
credentials	"1.3.2"	NA
curl	"5.0.2"	NA
DBI	"1.1.3"	NA

desc	"1.4.2"	NA
digest	"0.6.33"	NA
DMwR2	"0.0.2"	NA
dplyr	"1.1.2"	NA
ellipsis	"0.3.2"	NA
evaluate	"0.21"	NA
fansi	"1.0.4"	NA
farver	"2.1.1"	NA
fastmap	"1.1.1"	NA
fontawesome	"0.5.2"	NA
fs	"1.6.3"	NA
generics	"0.1.3"	NA
gert	"1.9.3"	NA
ggplot2	"3.4.3"	NA
gh	"1.4.0"	NA
gitcreds	"0.1.2"	NA
glue	"1.6.2"	NA
gplots	"3.1.3"	NA
gtable	"0.3.4"	NA
gtools	"3.9.4"	NA
highr	"0.10"	NA
hms	"1.1.3"	NA
htmltools	"0.5.6"	NA
httr2	"0.2.3"	NA
ini	"0.3.1"	NA
isoband	"0.2.7"	NA
jquerylib	"0.1.4"	NA
jsonlite	"1.8.7"	NA
KernSmooth	"2.23-22"	"recommended"
knitr	"1.43"	NA
labeling	"0.4.3"	NA
lifecycle	"1.0.3"	NA
magrittr	"2.0.3"	NA
Matrix	"1.6-1"	"recommended"
memoise	"2.0.1"	NA
mgcv	"1.9-0"	"recommended"
mime	"0.12"	NA
munSELL	"0.5.0"	NA
nlme	"3.1-163"	"recommended"
openssl	"2.1.0"	NA
palmerpenguins	"0.1.1"	NA
pillar	"1.9.0"	NA
pkgconfig	"2.0.3"	NA

prettyunits	"1.1.1"	NA
progress	"1.2.2"	NA
purrr	"1.0.2"	NA
quantmod	"0.4.25"	NA
R6	"2.5.1"	NA
rappdirs	"0.3.3"	NA
RColorBrewer	"1.1-3"	NA
readr	"2.1.4"	NA
rlang	"1.1.1"	NA
rmarkdown	"2.24"	NA
ROCR	"1.0-11"	NA
rprojroot	"2.0.3"	NA
rstudioapi	"0.15.0"	NA
sass	"0.4.7"	NA
scales	"1.2.1"	NA
spatial	"7.3-17"	"recommended"
stringi	"1.7.12"	NA
stringr	"1.5.0"	NA
survival	"3.5-7"	"recommended"
sys	"3.4.2"	NA
tibble	"3.2.1"	NA
tidyselect	"1.2.0"	NA
tinytex	"0.46"	NA
TTR	"0.24.3"	NA
tzdb	"0.4.0"	NA
usethis	"2.2.2"	NA
utf8	"1.2.3"	NA
vctrs	"0.6.3"	NA
viridisLite	"0.4.2"	NA
vroom	"1.6.3"	NA
whisker	"0.4.1"	NA
withr	"2.5.0"	NA
xfun	"0.40"	NA
xts	"0.13.1"	NA
yaml	"2.3.7"	NA
zip	"2.3.0"	NA
zoo	"1.8-12"	NA
base	"4.3.1"	"base"
boot	"1.3-28.1"	"recommended"
class	"7.3-22"	"recommended"
cluster	"2.1.4"	"recommended"
codetools	"0.2-19"	"recommended"
compiler	"4.3.1"	"base"

datasets	"4.3.1"	"base"
foreign	"0.8-84"	"recommended"
graphics	"4.3.1"	"base"
grDevices	"4.3.1"	"base"
grid	"4.3.1"	"base"
KernSmooth	"2.23-21"	"recommended"
lattice	"0.21-8"	"recommended"
MASS	"7.3-60"	"recommended"
Matrix	"1.5-4.1"	"recommended"
methods	"4.3.1"	"base"
mgcv	"1.8-42"	"recommended"
nlme	"3.1-162"	"recommended"
nnet	"7.3-19"	"recommended"
parallel	"4.3.1"	"base"
rpart	"4.1.19"	"recommended"
spatial	"7.3-16"	"recommended"
splines	"4.3.1"	"base"
stats	"4.3.1"	"base"
stats4	"4.3.1"	"base"
survival	"3.5-5"	"recommended"
tcltk	"4.3.1"	"base"
tools	"4.3.1"	"base"
utils	"4.3.1"	"base"
Depends		
abind	"R (>= 1.5.0)"	
askpass	NA	
base64enc	"R (>= 2.9.0)"	
bit	"R (>= 2.9.2)"	
bit64	"R (>= 3.0.1), bit (>= 4.0.0), utils, methods, stats"	
bitops	NA	
bslib	"R (>= 2.10)"	
cachem	NA	
caTools	"R (>= 3.6.0)"	
cli	"R (>= 3.4)"	
clipr	NA	
colorspace	"R (>= 3.0.0), methods"	
cpp11	"R (>= 3.5.0)"	
crayon	NA	
credentials	NA	
curl	"R (>= 3.0.0)"	
DBI	"methods, R (>= 3.0.0)"	
desc	"R (>= 3.4)"	
digest	"R (>= 3.3.0)"	

DMwR2	"R(>= 3.0), methods"
dplyr	"R (>= 3.5.0)"
ellipsis	"R (>= 3.2)"
evaluate	"R (>= 3.0.2)"
fansi	"R (>= 3.1.0)"
farver	NA
fastmap	NA
fontawesome	"R (>= 3.3.0)"
fs	"R (>= 3.4)"
generics	"R (>= 3.2)"
gert	NA
ggplot2	"R (>= 3.3)"
gh	"R (>= 3.4)"
gitcreds	"R (>= 3.4)"
glue	"R (>= 3.4)"
gplots	"R (>= 3.0)"
gtable	"R (>= 3.5)"
gtools	"methods, stats, utils"
highr	"R (>= 3.3.0)"
hms	NA
htmltools	"R (>= 2.14.1)"
httr2	"R (>= 3.4)"
ini	NA
isoband	NA
jquerylib	NA
jsonlite	"methods"
KernSmooth	"R (>= 2.5.0), stats"
knitr	"R (>= 3.3.0)"
labeling	NA
lifecycle	"R (>= 3.4)"
magrittr	"R (>= 3.4.0)"
Matrix	"R (>= 3.5.0), methods"
memoise	NA
mgcv	"R (>= 3.6.0), nlme (>= 3.1-64)"
mime	NA
munsell	NA
nlme	"R (>= 3.5.0)"
openssl	NA
palmerpenguins	"R (>= 2.10)"
pillar	NA
pkgconfig	NA
prettyunits	NA
progress	NA

purrr	"R (>= 3.5.0)"
quantmod	"R (>= 3.2.0), xts(>= 0.9-0), zoo, TTR(>= 0.2), methods"
R6	"R (>= 3.0)"
rappdirs	"R (>= 3.2)"
RColorBrewer	"R (>= 2.0.0)"
readr	"R (>= 3.5)"
rlang	"R (>= 3.5.0)"
rmarkdown	"R (>= 3.0)"
ROCR	"R (>= 3.6)"
rprojroot	"R (>= 3.0.0)"
rstudioapi	NA
sass	NA
scales	"R (>= 3.2)"
spatial	"R (>= 3.0.0), graphics, stats, utils"
stringi	"R (>= 3.1)"
stringr	"R (>= 3.3)"
survival	"R (>= 3.5.0)"
sys	NA
tibble	"R (>= 3.4.0)"
tidyselect	"R (>= 3.4)"
tinytex	NA
TTR	NA
tzdb	"R (>= 3.5.0)"
usethis	"R (>= 3.6)"
utf8	"R (>= 2.10)"
vctrs	"R (>= 3.5.0)"
viridisLite	"R (>= 2.10)"
vroom	"R (>= 3.4)"
whisker	NA
withr	"R (>= 3.2.0)"
xfun	NA
xts	"R (>= 3.6.0), zoo (>= 1.7-12)"
yaml	NA
zip	NA
zoo	"R (>= 3.1.0), stats"
base	NA
boot	"R (>= 3.0.0), graphics, stats"
class	"R (>= 3.0.0), stats, utils"
cluster	"R (>= 3.5.0)"
codetools	"R (>= 2.1)"
compiler	NA
datasets	NA
foreign	"R (>= 4.0.0)"

graphics	NA
grDevices	NA
grid	NA
KernSmooth	"R (>= 2.5.0), stats"
lattice	"R (>= 4.0.0)"
MASS	"R (>= 4.0), grDevices, graphics, stats, utils"
Matrix	"R (>= 3.5.0), methods"
methods	NA
mgcv	"R (>= 3.6.0), nlme (>= 3.1-64)"
nlme	"R (>= 3.5.0)"
nnet	"R (>= 3.0.0), stats, utils"
parallel	NA
rpart	"R (>= 2.15.0), graphics, stats, grDevices"
spatial	"R (>= 3.0.0), graphics, stats, utils"
splines	NA
stats	NA
stats4	NA
survival	"R (>= 3.5.0)"
tcltk	NA
tools	NA
utils	NA
	Imports
abind	"methods, utils"
askpass	"sys (>= 2.1)"
base64enc	NA
bit	NA
bit64	NA
bitops	NA
bslib	"base64enc, cachem, grDevices, htmltools (>= 0.5.4), jquerylib\n(>= 0.1.3), j
cachem	"rlang, fastmap (>= 1.1.1)"
caTools	"bitops"
cli	"utils"
clipr	"utils"
colorspace	"graphics, grDevices, stats"
cpp11	NA
crayon	"grDevices, methods, utils"
credentials	"openssl (>= 1.3), sys (>= 2.1), curl, jsonlite, askpass"
curl	NA
DBI	NA
desc	"cli, R6, rprojroot, utils"
digest	"utils"
DMwR2	"xts (>= 0.9-7), zoo (>= 1.7-10), class (>= 7.3-14), rpart (>= 4.1-10), quant
dplyr	"cli (>= 3.4.0), generics, glue (>= 1.3.2), lifecycle (>= 1.0.3), magrittr (>= 2.0.3)"

ellipsis	"rlang (>= 0.3.0)"
evaluate	"methods"
fansi	"grDevices, utils"
farver	NA
fastmap	NA
fontawesome	"rlang (>= 1.0.6), htmltools (>= 0.5.1.1)"
fs	"methods"
generics	"methods"
gert	"askpass, credentials (>= 1.2.1), openssl (>= 2.0.3), \nrstudioapi (>= 0.11), s"
ggplot2	"cli, glue, grDevices, grid, gtable (>= 0.1.1), isoband, \nlifecycle (> 1.0.1)"
gh	"cli (>= 3.0.1), gitcreds, httr2, ini, jsonlite, rlang (>= \n1.0.0)"
gitcreds	NA
glue	"methods"
gplots	"gtools, stats, caTools, KernSmooth, methods"
gtable	"cli, glue, grid, lifecycle, rlang (>= 1.1.0)"
gtools	NA
highr	"xfun (>= 0.18)"
hms	"lifecycle, methods, pkgconfig, rlang (>= 1.0.2), vctrs (>= \n0.3.8)"
htmltools	"utils, digest, grDevices, base64enc, rlang (>= 0.4.12), \nfastmap (>= 1.1.0), "
httr2	"cli (>= 3.0.0), curl, glue, magrittr, openssl, R6, rappdirs, \nrlang (>= 1.0.0)"
ini	NA
isoband	"grid, utils"
jquerylib	"htmltools"
jsonlite	NA
KernSmooth	NA
knitr	"evaluate (>= 0.15), highr, methods, tools, xfun (>= 0.39), \nyaml (>= 2.1.19)"
labeling	"stats, graphics"
lifecycle	"cli (>= 3.4.0), glue, rlang (>= 1.0.6)"
magrittr	NA
Matrix	"grDevices, graphics, grid, lattice, stats, utils"
memoise	"rlang (>= 0.4.10), cachem"
mgcv	"methods, stats, graphics, Matrix, splines, utils"
mime	"tools"
munSELL	"colorspace, methods"
nlme	"graphics, stats, utils, lattice"
openssl	"askpass"
palmerpenguins	NA
pillar	"cli (>= 2.3.0), fansi, glue, lifecycle, rlang (>= 1.0.2), utf8 \n (>= 1.1.0), v"
pkgconfig	"utils"
prettyunits	NA
progress	"hms, prettyunits, R6, crayon"
purrr	"cli (>= 3.6.1), lifecycle (>= 1.0.3), magrittr (>= 1.5.0), \nrlang (>= 1.1.1)"
quantmod	"curl, jsonlite (>= 1.1)"

R6	NA
rappdirs	NA
RColorBrewer	NA
readr	"cli (>= 3.2.0), clipr, crayon, hms (>= 0.4.1), lifecycle (>=\n0.2.0), methods"
rlang	"utils"
rmarkdown	"bslib (>= 0.2.5.1), evaluate (>= 0.13), fontawesome (>=\n0.5.0), htmltools"
ROCR	"methods, graphics, grDevices, gplots, stats"
rprojroot	NA
rstudioapi	NA
sass	"fs (>= 1.2.4), rlang (>= 0.4.10), htmltools (>= 0.5.1), R6,\nrappdirs"
scales	"farver (>= 2.0.3), labeling, lifecycle, munsell (>= 0.5), R6,\nRColorBrewer,
spatial	NA
stringi	"tools, utils, stats"
stringr	"cli, glue (>= 1.6.1), lifecycle (>= 1.0.3), magrittr, rlang\n(>= 1.0.0), str"
survival	"graphics, Matrix, methods, splines, stats, utils"
sys	NA
tibble	"fansi (>= 0.4.0), lifecycle (>= 1.0.0), magrittr, methods,\npillar (>= 1.8.1)
tidyselect	"cli (>= 3.3.0), glue (>= 1.3.0), lifecycle (>= 1.0.3), rlang\n(>= 1.0.4), vc"
tinytex	"xfun (>= 0.29)"
TTR	"xts (>= 0.10-0), zoo, curl"
tzdb	NA
usethis	"cli (>= 3.0.1), clipr (>= 0.3.0), crayon, curl (>= 2.7), desc\n(>= 1.4.2), f"
utf8	NA
vctrs	"cli (>= 3.4.0), glue, lifecycle (>= 1.0.3), rlang (>= 1.1.0)"
viridisLite	NA
vroom	"bit64, cli (>= 3.2.0), crayon, glue, hms, lifecycle (>=\n1.0.3), methods, rl"
whisker	NA
withr	"graphics, grDevices, stats"
xfun	"stats, tools"
xts	"methods"
yaml	NA
zip	NA
zoo	"utils, graphics, grDevices, lattice (>= 0.20-27)"
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
utils	NA
	LinkingTo
abind	NA
askpass	NA
base64enc	NA
bit	NA
bit64	NA
bitops	NA
bslib	NA
cachem	NA
caTools	NA
cli	NA
clipr	NA
colorspace	NA
cpp11	NA
crayon	NA
credentials	NA
curl	NA
DBI	NA
desc	NA
digest	NA
DMwR2	NA
dplyr	NA
ellipsis	NA
evaluate	NA

fansi	NA
farver	NA
fastmap	NA
fontawesome	NA
fs	NA
generics	NA
gert	NA
ggplot2	NA
gh	NA
gitcreds	NA
glue	NA
gplots	NA
gtable	NA
gtools	NA
highr	NA
hms	NA
htmltools	NA
httr2	NA
ini	NA
isoband	NA
jquerylib	NA
jsonlite	NA
KernSmooth	NA
knitr	NA
labeling	NA
lifecycle	NA
magrittr	NA
Matrix	NA
memoise	NA
mgcv	NA
mime	NA
munsell	NA
nlme	NA
openssl	NA
palmerpenguins	NA
pillar	NA
pkgconfig	NA
prettyunits	NA
progress	NA
purrr	"cli"
quantmod	NA
R6	NA
rappdirs	NA

RColorBrewer	NA
readr	"cpp11, tzdb (>= 0.1.1)"
rlang	NA
rmarkdown	NA
ROCR	NA
rprojroot	NA
rstudioapi	NA
sass	NA
scales	NA
spatial	NA
stringi	NA
stringr	NA
survival	NA
sys	NA
tibble	NA
tidyselect	NA
tinytex	NA
TTR	"xts"
tzdb	"cpp11 (>= 0.4.2)"
usethis	NA
utf8	NA
vctrs	NA
viridisLite	NA
vroom	"cpp11 (>= 0.2.0), progress (>= 1.2.1), tzdb (>= 0.1.1)"
whisker	NA
withr	NA
xfun	NA
xts	"zoo"
yaml	NA
zip	NA
zoo	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
utils	NA
	Suggests
abind	NA
askpass	"testthat"
base64enc	NA
bit	"testthat (>= 0.11.0), roxygen2, knitr, rmarkdown,\nmicrobenchmark, bit64 (>=
bit64	NA
bitops	NA
bslib	"bsicons, curl, fontawesome, ggplot2, knitr, magrittr,\nrappdirs, rmarkdown (>=
cachem	"testthat"
caTools	"MASS, rpart"
cli	"callr, covr, crayon, digest, glue (>= 1.6.0), grDevices,\nhtmltools, htmlwid
clipr	"covr, knitr, rmarkdown, rstudioapi (>= 0.5), testthat (>= \n2.0.0)"
colorspace	"datasets, utils, KernSmooth, MASS, kernlab, mvtnorm, vcd,\nntcltk, shiny, shin
cpp11	"bench, brio, callr, cli, covr, decor, desc, ggplot2, glue,\nknitr, lobstr, m
crayon	"mockery, rstudioapi, testthat, withr"
credentials	"testthat, knitr, rmarkdown"
curl	"spelling, testthat (>= 1.0.0), knitr, jsonlite, rmarkdown,\nmagrittr, httpuv
DBI	"blob, covr, DBItest, dbplyr, downlit, dplyr, glue, hms,\nknitr, magrittr, RM
desc	"callr, covr, gh, spelling, testthat, whoami, withr"
digest	"tinytest, simplermardown"
DMwR2	NA
dplyr	"bench, broom, callr, covr, DBI, dbplyr (>= 2.2.1), ggplot2,\nknitr, Lahman, I
ellipsis	"covr, testthat"
evaluate	"covr, ggplot2, lattice, rlang, testthat (>= 3.0.0), withr"
fansi	"unitizer, knitr, rmarkdown"
farver	"covr, testthat (>= 3.0.0)"

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,\ntestthat (>= 3.0.0)"
generics	"covr, pkgload, testthat (>= 3.0.0), tibble, withr"
gert	"spelling, knitr, rmarkdown, testthat"
ggplot2	"covr, dplyr, ggplot2movies, hexbin, Hmisc, knitr, lattice,\nmapproj, maps, mapproj"
gh	"covr, knitr, mockery, rmarkdown, rprojroot, spelling,\ntestthat (>= 3.0.0), vctrs"
gitcreds	"codetools, covr, knitr, mockery, oskeyring, rmarkdown,\ntestthat (>= 3.0.0), withr"
glue	"covr, crayon, DBI, dplyr, forcats, ggplot2, knitr, magrittr,\nmicrobenchmark, rmarkdown, testthat"
gplots	"grid, MASS, knitr, r2d2"
gtable	"covr, ggplot2, knitr, profvis, rmarkdown, testthat (>= 3.0.0)"
gtools	"car, gplots, knitr, rstudioapi, SGP, taxize"
highr	"knitr, markdown, testit"
hms	"crayon, lubridate, pillar (>= 1.1.0), testthat (>= 3.0.0)"
htmltools	"markdown, testthat, withr, Cairo, ragg, shiny"
httr2	"askpass, bench, clipr, covr, docopt, httpuv, jose, jsonlite,\nnknitr, purrr, rlang, rmarkdown, testthat"
ini	"testthat"
isoband	"covr, ggplot2, knitr, magick, microbenchmark, rmarkdown, sf,\ntestthat, xml2"
jquerylib	"testthat"
jsonlite	"httr, vctrs, testthat, knitr, rmarkdown, R.rsp, sf"
KernSmooth	"MASS, carData"
knitr	"bslib, codetools, DBI (>= 0.4-1), digest, formatR, gifski,\ngridSVG, htmlwidg"
labeling	NA
lifecycle	"covr, crayon, knitr, lintr, rmarkdown, testthat (>= 3.0.1),\ntibble, tidyver"
magrittr	"covr, knitr, rlang, rmarkdown, testthat"
Matrix	"MASS, datasets, sfsmisc"
memoise	"digest, aws.s3, covr, googleAuthR, googleCloudStorageR, httr,\ntestthat"
mgcv	"parallel, survival, MASS"
mime	NA
munSELL	"ggplot2, testthat"
nlme	"Hmisc, MASS, SASmixed"
openssl	"curl, testthat (>= 2.1.0), digest, knitr, rmarkdown,\nnjsonlite, jose, sodium"
palmerpenguins	"knitr, rmarkdown, tibble, ggplot2, dplyr, tidyr, recipes"
pillar	"bit64, DBI, debugme, DiagrammeR, dplyr, formattable, ggplot2,\nnknitr, lubridate"
pkgconfig	"covr, testthat, disposables (>= 1.0.3)"
prettyunits	"codetools, covr, testthat"
progress	"Rcpp, testthat, withr"
purrr	"covr, dplyr (>= 0.7.8), httr, knitr, lubridate, rmarkdown,\ntestthat (>= 3.0.0)"
quantmod	"DBI,RMySQL,RSQLite,timeSeries,xml2,downloader"
R6	"testthat, pryr"
rappdirs	"roxygen2, testthat (>= 3.0.0), covr, withr"
RColorBrewer	NA
readr	"covr, curl, datasets, knitr, rmarkdown, spelling, stringi,\ntestthat (>= 3.1.0)"

rlang	"cli (>= 3.1.0), covr, crayon, fs, glue, knitr, magrittr,\nmetho
rmarkdown	"digest, dygraphs, fs, rsconnect, downlit (>= 0.4.0), katex\n(>= 1.4.0), sass
ROCR	"testthat, knitr, rmarkdown"
rprojroot	"covr, knitr, lifecycle, mockr, rmarkdown, testthat (>=\n3.0.0), withr"
rstudioapi	"testthat, knitr, rmarkdown, clipr, covr"
sass	"testthat, knitr, rmarkdown, withr, shiny, curl"
scales	"bit64, covr, dichromat, ggplot2, hms (>= 0.5.0), stringi,\ntestthat (>= 3.0.0)
spatial	"MASS"
stringi	NA
stringr	"covr, htmltools, htmlwidgets, knitr, rmarkdown, testthat (>=\n3.0.0)"
survival	NA
sys	"unix (>= 1.4), spelling, testthat"
tibble	"bench, bit64, blob, brio, callr, cli, covr, crayon (>=\n1.3.4), DiagrammeR, c
tidyselect	"covr, crayon, dplyr, knitr, magrittr, rmarkdown, stringr,\ntestthat (>= 3.1.0)
tinytex	"testit, rstudioapi"
TTR	"RUnit"
tzdb	"covr, testthat (>= 3.0.0)"
usethis	"covr, knitr, magick, pkgload, rmarkdown, roxygen2 (>= 7.1.2),\nspelling (>= 1.2.0), testthat (>= 3.0.0), withr"
utf8	"cli, covr, knitr, rlang, rmarkdown, testthat (>= 3.0.0),\nwit
vctrs	"bit64, covr, crayon, dplyr (>= 0.8.5), generics, knitr,\npillar (>= 1.4.4), p
viridisLite	"hexbin (>= 1.27.0), ggplot2 (>= 1.0.1), testthat, covr"
vroom	"archive, bench (>= 1.1.0), covr, curl, dplyr, forcats, fs,\nnggplot2, knitr, p
whisker	"markdown"
withr	"callr, covr, DBI, knitr, lattice, methods, rlang, rmarkdown\n(>= 2.12), RSQL
xfun	"testit, parallel, codetools, rstudioapi, tinytex (>= 0.30),\nmime, markdown
xts	"timeSeries, timeDate, tseries, chron, tinytest"
yaml	"RUnit"
zip	"covr, processx, R6, testthat, withr"
zoo	"AER, coda, chron, ggplot2 (>= 3.0.0), mondate, scales,\nstinepack, strucchang
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	"MASS, expm"
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, mathjaxr, V8"
utils	"methods, xml2, commonmark, knitr"
	Enhances
abind	NA
askpass	NA
base64enc	"png"
bit	NA
bit64	NA
bitops	NA
bslib	NA
cachem	NA
caTools	NA
cli	NA
clipr	NA
colorspace	NA
cpp11	NA
crayon	NA
credentials	NA
curl	NA
DBI	NA
desc	NA
digest	NA
DMwR2	NA
dplyr	NA
ellipsis	NA
evaluate	NA
fansi	NA
farver	NA
fastmap	NA
fontawesome	NA

fs	NA
generics	NA
gert	NA
ggplot2	"sp"
gh	NA
gitcreds	NA
glue	NA
gplots	NA
gtable	NA
gtools	NA
highr	NA
hms	NA
htmltools	"knitr"
httr2	NA
ini	NA
isoband	NA
jquerylib	NA
jsonlite	NA
KernSmooth	NA
knitr	NA
labeling	NA
lifecycle	NA
magrittr	NA
Matrix	"SparseM, graph"
memoise	NA
mgcv	NA
mime	NA
munsell	NA
nlme	NA
openssl	NA
palmerpenguins	NA
pillar	NA
pkgconfig	NA
prettyunits	NA
progress	NA
purrr	NA
quantmod	NA
R6	NA
rappdirs	NA
RColorBrewer	NA
readr	NA
rlang	"winch"
rmarkdown	NA

ROCR	NA
rprojroot	NA
rstudioapi	NA
sass	NA
scales	NA
spatial	NA
stringi	NA
stringr	NA
survival	NA
sys	NA
tibble	NA
tidyselect	NA
tinytex	NA
TTR	"quantmod"
tzdb	NA
usethis	NA
utf8	NA
vctrs	NA
viridisLite	NA
vroom	NA
whisker	NA
withr	NA
xfun	NA
xts	NA
yaml	NA
zip	NA
zoo	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, SparseM, graph, igraph, maptools, sfsmisc, 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	
utils	NA	
	License	License_is_FOSS
abind	"LGPL (>= 2)"	NA
askpass	"MIT + file LICENSE"	NA
base64enc	"GPL-2 GPL-3"	NA
bit	"GPL-2 GPL-3"	NA
bit64	"GPL-2 GPL-3"	NA
bitops	"GPL (>= 2)"	NA
bslib	"MIT + file LICENSE"	NA
cachem	"MIT + file LICENSE"	NA
caTools	"GPL-3"	NA
cli	"MIT + file LICENSE"	NA
clipr	"GPL-3"	NA
colorspace	"BSD_3_clause + file LICENSE"	NA
cpp11	"MIT + file LICENSE"	NA
crayon	"MIT + file LICENSE"	NA
credentials	"MIT + file LICENSE"	NA
curl	"MIT + file LICENSE"	NA
DBI	"LGPL (>= 2.1)"	NA
desc	"MIT + file LICENSE"	NA
digest	"GPL (>= 2)"	NA
DMwR2	"GPL (>= 2)"	NA
dplyr	"MIT + file LICENSE"	NA
ellipsis	"MIT + file LICENSE"	NA
evaluate	"MIT + file LICENSE"	NA
fansi	"GPL-2 GPL-3"	NA
farver	"MIT + file LICENSE"	NA
fastmap	"MIT + file LICENSE"	NA
fontawesome	"MIT + file LICENSE"	NA
fs	"MIT + file LICENSE"	NA
generics	"MIT + file LICENSE"	NA

gert	"MIT + file LICENSE"	NA
ggplot2	"MIT + file LICENSE"	NA
gh	"MIT + file LICENSE"	NA
gitcreds	"MIT + file LICENSE"	NA
glue	"MIT + file LICENSE"	NA
gplots	"GPL-2"	NA
gtable	"MIT + file LICENSE"	NA
gtools	"GPL-2"	NA
highr	"GPL"	NA
hms	"MIT + file LICENSE"	NA
htmltools	"GPL (>= 2)"	NA
httr2	"MIT + file LICENSE"	NA
ini	"GPL-3"	NA
isoband	"MIT + file LICENSE"	NA
jquerylib	"MIT + file LICENSE"	NA
jsonlite	"MIT + file LICENSE"	NA
KernSmooth	"Unlimited"	NA
knitr	"GPL"	NA
labeling	"MIT + file LICENSE Unlimited"	NA
lifecycle	"MIT + file LICENSE"	NA
magrittr	"MIT + file LICENSE"	NA
Matrix	"GPL (>= 2) file LICENSE"	NA
memoise	"MIT + file LICENSE"	NA
mgcv	"GPL (>= 2)"	NA
mime	"GPL"	NA
munSELL	"MIT + file LICENSE"	NA
nlme	"GPL (>= 2)"	NA
openssl	"MIT + file LICENSE"	NA
palmerpenguins	"CC0"	NA
pillar	"MIT + file LICENSE"	NA
pkgconfig	"MIT + file LICENSE"	NA
prettyunits	"MIT + file LICENSE"	NA
progress	"MIT + file LICENSE"	NA
purrr	"MIT + file LICENSE"	NA
quantmod	"GPL-3"	NA
R6	"MIT + file LICENSE"	NA
rappdirs	"MIT + file LICENSE"	NA
RColorBrewer	"Apache License 2.0"	NA
readr	"MIT + file LICENSE"	NA
rlang	"MIT + file LICENSE"	NA
rmarkdown	"GPL-3"	NA
ROCR	"GPL (>= 2)"	NA
rprojroot	"MIT + file LICENSE"	NA

rstudioapi	"MIT + file LICENSE"	NA
sass	"MIT + file LICENSE"	NA
scales	"MIT + file LICENSE"	NA
spatial	"GPL-2 GPL-3"	NA
stringi	"file LICENSE"	"yes"
stringr	"MIT + file LICENSE"	NA
survival	"LGPL (>= 2)"	NA
sys	"MIT + file LICENSE"	NA
tibble	"MIT + file LICENSE"	NA
tidyselect	"MIT + file LICENSE"	NA
tinytex	"MIT + file LICENSE"	NA
TTR	"GPL (>= 2)"	NA
tzdb	"MIT + file LICENSE"	NA
usethis	"MIT + file LICENSE"	NA
utf8	"Apache License (== 2.0) file LICENSE"	NA
vctrs	"MIT + file LICENSE"	NA
viridisLite	"MIT + file LICENSE"	NA
vroom	"MIT + file LICENSE"	NA
whisker	"GPL-3"	NA
withr	"MIT + file LICENSE"	NA
xfun	"MIT + file LICENSE"	NA
xts	"GPL (>= 2)"	NA
yaml	"BSD_3_clause + file LICENSE"	NA
zip	"MIT + file LICENSE"	NA
zoo	"GPL-2 GPL-3"	NA
base	"Part of R 4.3.1"	NA
boot	"Unlimited"	NA
class	"GPL-2 GPL-3"	NA
cluster	"GPL (>= 2)"	NA
codetools	"GPL"	NA
compiler	"Part of R 4.3.1"	NA
datasets	"Part of R 4.3.1"	NA
foreign	"GPL (>= 2)"	NA
graphics	"Part of R 4.3.1"	NA
grDevices	"Part of R 4.3.1"	NA
grid	"Part of R 4.3.1"	NA
KernSmooth	"Unlimited"	NA
lattice	"GPL (>= 2)"	NA
MASS	"GPL-2 GPL-3"	NA
Matrix	"GPL (>= 2) file LICENCE"	NA
methods	"Part of R 4.3.1"	NA
mgcv	"GPL (>= 2)"	NA
nlme	"GPL (>= 2)"	NA

nnet	"GPL-2 GPL-3"			NA	
parallel	"Part of R 4.3.1"			NA	
rpart	"GPL-2 GPL-3"			NA	
spatial	"GPL-2 GPL-3"			NA	
splines	"Part of R 4.3.1"			NA	
stats	"Part of R 4.3.1"			NA	
stats4	"Part of R 4.3.1"			NA	
survival	"LGPL (>= 2)"			NA	
tcltk	"Part of R 4.3.1"			NA	
tools	"Part of R 4.3.1"			NA	
utils	"Part of R 4.3.1"			NA	
	License_restricts_use	OS_type	MD5sum	NeedsCompilation	Built
abind	NA	NA	NA	"no"	"4.3.0"
askpass	NA	NA	NA	"yes"	"4.3.0"
base64enc	NA	NA	NA	"yes"	"4.3.0"
bit	NA	NA	NA	"yes"	"4.3.0"
bit64	NA	NA	NA	"yes"	"4.3.0"
bitops	NA	NA	NA	"yes"	"4.3.0"
bslib	NA	NA	NA	"no"	"4.3.0"
cachem	NA	NA	NA	"yes"	"4.3.0"
caTools	NA	NA	NA	"yes"	"4.3.0"
cli	NA	NA	NA	"yes"	"4.3.0"
clipr	NA	NA	NA	"no"	"4.3.0"
colorspace	NA	NA	NA	"yes"	"4.3.0"
cpp11	NA	NA	NA	"no"	"4.3.0"
crayon	NA	NA	NA	"no"	"4.3.0"
credentials	NA	NA	NA	"no"	"4.3.0"
curl	NA	NA	NA	"yes"	"4.3.0"
DBI	NA	NA	NA	"no"	"4.3.0"
desc	NA	NA	NA	"no"	"4.3.0"
digest	NA	NA	NA	"yes"	"4.3.0"
DMwR2	NA	NA	NA	"no"	"4.3.0"
dplyr	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"
farver	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"
generics	NA	NA	NA	"no"	"4.3.0"
gert	NA	NA	NA	"yes"	"4.3.0"
ggplot2	NA	NA	NA	"no"	"4.3.0"

gh	NA	NA	NA	"no"	"4.3.0"
gitcreds	NA	NA	NA	"no"	"4.3.0"
glue	NA	NA	NA	"yes"	"4.3.0"
gplots	NA	NA	NA	"no"	"4.3.0"
gtable	NA	NA	NA	"no"	"4.3.0"
gtools	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"
httr2	NA	NA	NA	"no"	"4.3.0"
ini	NA	NA	NA	"no"	"4.3.0"
isoband	NA	NA	NA	"yes"	"4.3.0"
jquerylib	NA	NA	NA	"no"	"4.3.0"
jsonlite	NA	NA	NA	"yes"	"4.3.0"
KernSmooth	NA	NA	NA	"yes"	"4.3.0"
knitr	NA	NA	NA	"no"	"4.3.0"
labeling	NA	NA	NA	"no"	"4.3.0"
lifecycle	NA	NA	NA	"no"	"4.3.0"
magrittr	NA	NA	NA	"yes"	"4.3.0"
Matrix	NA	NA	NA	"yes"	"4.3.0"
memoise	NA	NA	NA	"no"	"4.3.0"
mgcv	NA	NA	NA	"yes"	"4.3.0"
mime	NA	NA	NA	"yes"	"4.3.0"
munsell	NA	NA	NA	"no"	"4.3.0"
nlme	NA	NA	NA	"yes"	"4.3.0"
openssl	NA	NA	NA	"yes"	"4.3.0"
palmerpenguins	NA	NA	NA	"no"	"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"
purrr	NA	NA	NA	"yes"	"4.3.0"
quantmod	NA	NA	NA	"no"	"4.3.0"
R6	NA	NA	NA	"no"	"4.3.0"
rappdirs	NA	NA	NA	"yes"	"4.3.0"
RColorBrewer	NA	NA	NA	"no"	"4.3.0"
readr	NA	NA	NA	"yes"	"4.3.0"
rlang	NA	NA	NA	"yes"	"4.3.0"
rmarkdown	NA	NA	NA	"no"	"4.3.0"
ROCR	NA	NA	NA	"no"	"4.3.0"
rprojroot	NA	NA	NA	"no"	"4.3.0"
rstudioapi	NA	NA	NA	"no"	"4.3.0"
sass	NA	NA	NA	"yes"	"4.3.0"

scales	NA	NA	NA	"no"	"4.3.0"
spatial	NA	NA	NA	"yes"	"4.3.0"
stringi	NA	NA	NA	"yes"	"4.3.0"
stringr	NA	NA	NA	"no"	"4.3.0"
survival	NA	NA	NA	"yes"	"4.3.0"
sys	NA	NA	NA	"yes"	"4.3.0"
tibble	NA	NA	NA	"yes"	"4.3.0"
tidyselect	NA	NA	NA	"no"	"4.3.0"
tinytex	NA	NA	NA	"no"	"4.3.0"
TTR	NA	NA	NA	"yes"	"4.3.0"
tzdb	NA	NA	NA	"yes"	"4.3.0"
usethis	NA	NA	NA	"no"	"4.3.0"
utf8	NA	NA	NA	"yes"	"4.3.0"
vctrs	NA	NA	NA	"yes"	"4.3.0"
viridisLite	NA	NA	NA	"no"	"4.3.0"
vroom	NA	NA	NA	"yes"	"4.3.0"
whisker	NA	NA	NA	"no"	"4.3.0"
withr	NA	NA	NA	"no"	"4.3.0"
xfun	NA	NA	NA	"yes"	"4.3.0"
xts	NA	NA	NA	"yes"	"4.3.0"
yaml	NA	NA	NA	"yes"	"4.3.0"
zip	NA	NA	NA	"yes"	"4.3.0"
zoo	NA	NA	NA	"yes"	"4.3.0"
base	NA	NA	NA	NA	"4.3.1"
boot	NA	NA	NA	"no"	"4.3.1"
class	NA	NA	NA	"yes"	"4.3.1"
cluster	NA	NA	NA	"yes"	"4.3.1"
codetools	NA	NA	NA	"no"	"4.3.1"
compiler	NA	NA	NA	NA	"4.3.1"
datasets	NA	NA	NA	NA	"4.3.1"
foreign	NA	NA	NA	"yes"	"4.3.1"
graphics	NA	NA	NA	"yes"	"4.3.1"
grDevices	NA	NA	NA	"yes"	"4.3.1"
grid	NA	NA	NA	"yes"	"4.3.1"
KernSmooth	NA	NA	NA	"yes"	"4.3.1"
lattice	NA	NA	NA	"yes"	"4.3.1"
MASS	NA	NA	NA	"yes"	"4.3.1"
Matrix	NA	NA	NA	"yes"	"4.3.1"
methods	NA	NA	NA	"yes"	"4.3.1"
mgcv	NA	NA	NA	"yes"	"4.3.1"
nlme	NA	NA	NA	"yes"	"4.3.1"
nnet	NA	NA	NA	"yes"	"4.3.1"
parallel	NA	NA	NA	"yes"	"4.3.1"

rpart	NA	NA	NA	"yes"	"4.3.1"
spatial	NA	NA	NA	"yes"	"4.3.1"
splines	NA	NA	NA	"yes"	"4.3.1"
stats	NA	NA	NA	"yes"	"4.3.1"
stats4	NA	NA	NA	NA	"4.3.1"
survival	NA	NA	NA	"yes"	"4.3.1"
tcltk	NA	NA	NA	"yes"	"4.3.1"
tools	NA	NA	NA	"yes"	"4.3.1"
utils	NA	NA	NA	"yes"	"4.3.1"

old.packages() function is used to check for and list packages that have newer versions available on CRAN . This function is **helpful for keeping your packages up-to-date**.

```
old.packages()
```

	Package	LibPath	Installed	Built
KernSmooth	"KernSmooth"	"/opt/R/4.3.1/lib/R/library"	"2.23-21"	"4.3.1"
Matrix	"Matrix"	"/opt/R/4.3.1/lib/R/library"	"1.5-4.1"	"4.3.1"
mgcv	"mgcv"	"/opt/R/4.3.1/lib/R/library"	"1.8-42"	"4.3.1"
nlme	"nlme"	"/opt/R/4.3.1/lib/R/library"	"3.1-162"	"4.3.1"
spatial	"spatial"	"/opt/R/4.3.1/lib/R/library"	"7.3-16"	"4.3.1"
survival	"survival"	"/opt/R/4.3.1/lib/R/library"	"3.5-5"	"4.3.1"
	ReposVer	Repository		
KernSmooth	"2.23-22"	"http://rspm/default/__linux__/focal/latest/src/contrib"		
Matrix	"1.6-1"	"http://rspm/default/__linux__/focal/latest/src/contrib"		
mgcv	"1.9-0"	"http://rspm/default/__linux__/focal/latest/src/contrib"		
nlme	"3.1-163"	"http://rspm/default/__linux__/focal/latest/src/contrib"		
spatial	"7.3-17"	"http://rspm/default/__linux__/focal/latest/src/contrib"		
survival	"3.5-7"	"http://rspm/default/__linux__/focal/latest/src/contrib"		

update.packages() function is used to update one or more packages to their latest versions available on CRAN

```
update.packages() #ask will update the package without asking for confirmation & checkBuild
```

You can use the **::** operator followed by the function name to see the package namespace it comes from. For example, to find out which package the mean function belongs to, you can do:

```
mean
```

```
function (x, ...)
UseMethod("mean")
<bytecode: 0x555a2c57af38>
<environment: namespace:base>
```

help() function is used to access documentation and information about functions, datasets & packages.

```
help(mean)
```

An alternative way to access help is by using a question mark ?

```
?mean
```

RsiteSearch() function allows you to search for specific terms, keywords, or phrases within the vast collection of R packages, functions, and documentation hosted on CRAN.

```
RSiteSearch('neural networks')
```

A search query has been submitted to <https://search.r-project.org>
The results page should open in your browser shortly

R objects and Variables

Variables are assigned values, which can be numeric, character, logical, or other data types.

```
vat <- 0.2
vat
```

```
[1] 0.2
```

Enclosing a statement or expression in `()` will have values printed directly to the console.

```
(vat <- 0.2)
```

```
[1] 0.2
```

examples :


```
x <- 5
y <- vat * x
y
```

```
[1] 1
```

```
z <- (y/2)^2
y
```

```
[1] 1
```

```
z
```

```
[1] 0.25
```

`ls()` or `objects()` function to list the names of objects (variables, functions, datasets, etc.) that are currently present in your workspace or environment.

```
ls()
```

```
[1] "algae"          "has_annotatons" "vat"           "x"
[5] "y"              "z"
```

```
objects()
```

```
[1] "algae"          "has_annotatons" "vat"           "x"
[5] "y"              "z"
```

`rm()` function is used to remove or delete objects from environment. Deleting objects means they will no longer be available for use, and **there is no undo operation**.

```
rm(vat)
ls()
```

```
[1] "algae"          "has_annotatons" "x"             "y"
[5] "z"
```

R Functions

functions are blocks of reusable code that perform specific tasks or computations.

```
max(4, 5, 6, 12, -4)
```

```
[1] 12
```

```
mean(4, 5, 6, 12, -4)
```

```
[1] 4
```

sample() function is used to generate random samples or permutations of elements from a given vector or set.

```
max(sample(1:100, 30)) # 1:100 is the vector set and 30 is random number of samples
```

```
[1] 95
```

```
mean(sample(1:100, 30))
```

```
[1] 54.03333
```

set.seed()

Setting the seed allows you to reproduce random results in your code. When you use random functions or generate random numbers without specifying a seed, the results will be different each time you run the code. **runif()**, **rnorm()**, or **sample()**, will produce the same random results as long as you use the same seed value.

```
set.seed(1)  
rnorm(1)
```

```
[1] -0.6264538
```

```
set.seed(2)
rnorm(1)
```

```
[1] -0.8969145
```

```
rnorm(1)
```

```
[1] 0.1848492
```

To create a new function, `se` (standard error of means), first test if `se` exists in our current environment.

```
exists("se")
```

```
[1] FALSE
```

No object named `se` exists, now create the function that computes the standard error of a sample:

```
se <- function(x){
  variance <- var(x)
  n <- length(x)
  return (sqrt(variance/n))
}
```

We can check if object exists or not using `exists()` function

```
exists("se")
```

```
[1] TRUE
```

Function with multiple arguments :

```
convMeters <- function (x, to="inch"){
  factor = switch(to, inch=39.3701, foot=3.28084, yard=1.09361, mile=0.000621371, NA)
  if(is.na(factor)) stop ("unknown target unit")
}
```

```

    else return (x*factor)
  }
  convMeters(23, "foot") #calling the function with parameters

```

```
[1] 75.45932
```

```
convMeters(40) #inch is used as default if 2nd argument is not used
```

```
[1] 1574.804
```

```
convMeters(to="yard", 56.2) #arguements can be provided in different order also
```

```
[1] 61.46088
```

Factors

Factors are a data type used to represent categorical or nominal data.

To create a factor with specific levels using the **factor()** function in R, you can specify the levels using the **levels** argument.

```

g <-c('f', 'm', 'f', 'f', 'f', 'm', 'm', 'f')
g #returs the values

```

```
[1] "f" "m" "f" "f" "f" "m" "m" "f"
```

```

g <- factor(g) #returns the levels of the factor
g

```

```

[1] f m f f f m m f
Levels: f m

```

More compact way to creating a factor with known levels, f and m:

```
other.g <-factor(c('m', 'm', 'm', 'm'), levels= c('f', 'm'))
other.g
```

```
[1] m m m m
Levels: f m
```

table() function to create a contingency table when you have two categorical variables.

```
g <- factor(c('f', 'm', 'f', 'f', 'f', 'm', 'm', 'f'))
table(g)
```

```
g
f m
5 3
```

```
a <- factor(c('adult', 'juvenile','adult', 'juvenile','adult', 'juvenile','juvenile', 'juv
table(a, g)
```

```
      g
a      f m
adult  3 0
juvenile 2 3
```

R assumes the values at the same index in the two factors are associated with the same entity.

```
# a <- factor(c('adult', 'juvenile','adult', 'juvenile','adult', 'juvenile','juvenile'))
# table(a, g) # will give an error as number of arguments should be same
```

```
a <- factor(c('adult', 'juvenile','adult', 'juvenile','adult', 'juvenile','juvenile', 'juv
t <- table(a, g)
t
```

```
      g
a      f m
adult  3 0
juvenile 2 3
```

Marginal frequencies for a factor:

```
margin.table(t, 1)#1 refers to the first factor, a (age)
```

```
a
  adult juvenile
    3         5
```

```
margin.table(t, 2)# now find the marginal freq of the second factor g
```

```
g
f m
5 3
```

```
prop.table(t, 1) #use the margin generated for the 1st factor a
```

```
      g
a      f  m
adult  1.0 0.0
juvenile 0.4 0.6
```

```
prop.table(t, 2)
```

```
      g
a      f  m
adult  0.6 0.0
juvenile 0.4 1.0
```

```
prop.table(t) #overall
```

```
      g
a      f  m
adult  0.375 0.000
juvenile 0.250 0.375
```

```
prop.table(t) * 100
```

	g		
a	f	m	
adult	37.5	0.0	
juvenile	25.0	37.5	

R structures

Vectors

Data structure used to store and manipulate a sequence of values.

```
v <- c(2, 5, 3, 4) #creating the vector  
length(v) #returns the lenght of the vector
```

```
[1] 4
```

```
mode(v) #returns the data type of vector
```

```
[1] "numeric"
```

```
v <- c(2, 5, 3, 4, NA) #NA will represent the missing value  
mode(v)
```

```
[1] "numeric"
```

Boolean vector

```
b <- c(TRUE, FALSE, NA, TRUE)  
mode(b)
```

```
[1] "logical"
```

```
b[3] #returns the 3rd element from vector
```

```
[1] NA
```

```
b[3] <- TRUE #update the value of 3rd element  
b
```

```
[1] TRUE FALSE TRUE TRUE
```

```
e <-vector()  
mode(e)
```

```
[1] "logical"
```

```
e <- c()  
mode(e)
```

```
[1] "NULL"
```

```
length(e) # returns the length as 0 as vector is empty
```

```
[1] 0
```

```
b2 <-c(b[1], b[3], b[5]) #using vector elements to create other vector  
b2
```

```
[1] TRUE TRUE NA
```

```
sqrt(v) # finding square root of all elements in the vector
```

```
[1] 1.414214 2.236068 1.732051 2.000000 NA
```

Vector arithmetic


```
v1 <- c(3, 6, 9)
v2 <- c(1, 4, 8)
v1+v2 #addition
```

```
[1] 4 10 17
```

```
v1*v2 #dot product
```

```
[1] 3 24 72
```

```
v1-v2 #subtraction
```

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

```
v1/v2 #divsion
```

```
[1] 3.000 1.500 1.125
```

```
v3 <- c(1, 4)
v1+v3 # Dynamically matches the length of longer vector making v3(1,4,1)
```

Warning in v1 + v3: longer object length is not a multiple of shorter object length

```
[1] 4 10 10
```

```
#vector for 'for' loop:
mysum <- function (x){
  sum <- 0
  for(i in 1:length(x)){
    sum <- sum + x[i]
  }
  return (sum)
```

```
}  
  
(mysum (c(1, 2, 3)))
```

```
[1] 6
```

Generating vectors:

```
(x <-1:10)
```

```
[1] 1 2 3 4 5 6 7 8 9 10
```

```
(x <-10:1)
```

```
[1] 10 9 8 7 6 5 4 3 2 1
```

```
10:15-1 #priority of the : operator is more than arithmetic operators
```

```
[1] 9 10 11 12 13 14
```

```
10:(15-1)
```

```
[1] 10 11 12 13 14
```

seq() to generate sequence with real numbers:

```
(seq(from=1, to=5, length=4)) # 4 values between 1 and 5 inclusive, even intervals/steps
```

```
[1] 1.000000 2.333333 3.666667 5.000000
```

```
(seq(length=10, from=-2, by=0.5)) #10 values, starting from 2, interval/step = 0.5
```

```
[1] -2.0 -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5 2.0 2.5
```

```
(rep(5, 10)) #rep(a, b) is used for repeating the a, b number of times
```

```
[1] 5 5 5 5 5 5 5 5 5 5
```

```
(rep("hi", 3))
```

```
[1] "hi" "hi" "hi"
```

```
(rep(1:2, 3)) #repeating the multiple values
```

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

```
(rep(TRUE:FALSE, 3)) #repeating the boolean values
```

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

```
(rep(1:2, each=3)) #repeating multiple values separately
```

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

gl() function is used to generate factor levels for creating factors with specific patterns, such as repeated or nested factors.

```
gl(3, 5) #three levels, each repeat 5 times
```

```
[1] 1 1 1 1 1 2 2 2 2 2 3 3 3 3 3  
Levels: 1 2 3
```

```
gl(2, 5, labels= c('female', 'male'))#two levels, each level repeat 5 times
```

```
[1] female female female female female male   male   male   male   male  
Levels: female male
```

```
#first argument 2 says two levels.
#second argument 1 says repeat once
#third argument 20 says generate 20 values
gl(2, 1, 20, labels=c('female', 'male'))#10 alternating female and male pairs, a total of
```

```
[1] female male   female male   female male   female male   female male
[11] female male   female male   female male   female male   female male
Levels: female male
```

```
n <- rep(1:2, each=3)
(n <- factor(n,
             levels = c(1, 2),
             labels = c('female', 'male')
             ))
```

```
[1] female female female male   male   male
Levels: female male
```

Generate 10 values following a normal distribution with mean = 10 and standard deviation = 3

```
(rnorm(10, mean=10, sd=3))
```

```
[1] 14.763536  6.608873  9.759245 10.397261 12.123864  9.280906 15.953422
[8]  9.583639 11.252952 12.945258
```

Exercise

Generate a random sample of normally distributed data of size 100, with a mean of 20 and standard deviation 4

```
sam <- (rnorm(100, mean=20, sd=4))
sam
```

```
[1] 18.42922 15.84132 27.12892 10.75572 23.51442 20.14323 24.05131 21.72906
[9] 28.36328 15.20030 26.35855 27.81861 20.01975 10.19317 21.90895 17.61377
[17] 23.16881 21.15855 22.95575 21.27584 24.30466 18.86337 16.89330 17.61736
[25] 13.09608 16.38966 17.76375 19.01395 18.46566 12.16359 16.63318 27.61419
[33] 22.48998 27.96368 18.77807 19.63662 19.26335 15.20493 16.64685 28.26521
[41] 17.75101 25.10286 15.80971 12.13649 18.70812 23.74345 24.55692 26.68648
[49] 12.84703 28.12497 17.18742 20.63266 22.02494 16.72002 12.00461 18.08283
[57] 20.33672 16.41805 16.31490 21.32180 19.43336 21.73939 19.78511 16.37156
[65] 25.21405 23.08716 24.21010 14.35985 23.98394 13.21694 17.86651 14.51092
[73] 11.16832 27.28849 17.38643 18.86128 18.45220 21.54678 26.40156 26.72462
[81] 15.26557 14.56617 13.94932 14.98758 27.83743 20.03058 16.62954 17.59536
[89] 24.29784 21.04239 18.74291 17.00148 16.55121 28.19216 23.75968 28.03475
[97] 18.31451 18.59666 15.89048 18.99792
```

```
mean_sam <- mean(sam) # Calculate the mean

sd_value <- sd(sam) # Calculate the standard deviation

sample_size <- length(sam) # Calculate the sample size

se <- sd_value / sqrt(sample_size) ## Compute the standard error
se
```

```
[1] 0.4688534
```

Sub-setting

Subsetting is the process of selecting a subset of elements or rows from data.

```
x <- c(0, -3, 4, -1, 45, 90, -5)
#select all elements that is greater than 0
(gtzero <- x[x>0])
```

```
[1] 4 45 90
```

```
x <- c(0, -3, 4, -1, 45, 90, -5)
(x[x<=-2 | x>5]) #Using OR operator
```

```
[1] -3 45 90 -5
```

```
(x[x>40 & x<100]) #using AND operator
```

```
[1] 45 90
```

```
x <- c(0, -3, 4, -1, 45, 90, -5)
(x[c(4, 6)]) #using vector index to selective elements
```

```
[1] -1 90
```

```
(y<-c(4,6)) #same as above
```

```
[1] 4 6
```

```
(x[y])
```

```
[1] -1 90
```

```
(x[1:3]) #select the 1st to the 3rd elements in the vector
```

```
[1] 0 -3 4
```

```
x <- c(0, -3, 4, -1, 45, 90, -5)
(x[-1]) #using negative index to exclude the element
```

```
[1] -3 4 -1 45 90 -5
```

```
(x[-c(4, 6)]) #using negative index to remove multiple elements
```

```
[1] 0 -3 4 45 -5
```

```
(x[-(1:3)]) #using negative index to remove range of elements
```

```
[1] -1 45 90 -5
```

Named elements

Named elements allow you to associate names or labels with individual components, making it easier to reference and work with your data.

```
x <- c(0, -3, 4, -1, 45, 90, -5)
names(x) <- c('s1', 's2', 's3', 's4', 's5', 's6', 's7')
x
```

```
s1 s2 s3 s4 s5 s6 s7
0 -3 4 -1 45 90 -5
```

```
(pH <- c(area1=4.5, area2=5.7, area3=9.8, mud=7.2)) #naming elements while creating the vector
```

```
area1 area2 area3 mud
4.5 5.7 9.8 7.2
```

```
pH['mud'] #selecting element using the label.
```

```
mud
7.2
```

```
pH[c('area1', 'mud')]
```

```
area1 mud
4.5 7.2
```

```
# x[-s1] #cannot use negative label to exclude the element
#x[-"s1"]
```

```
#x[s1:s7] #invalid argument
#x[c('s1':'s7')]
```

```
pH[] #empty index returns all the elements of the vector
```

```
area1 area2 area3 mud
4.5    5.7    9.8  7.2
```

```
pH
```

```
area1 area2 area3 mud
4.5    5.7    9.8  7.2
```

```
pH[] <- 0 #resetting the value of vector to zero
pH
```

```
area1 area2 area3 mud
0      0      0      0
```

Matrices and Arrays

Matrices and arrays are data structures used to store and manipulate multi-dimensional data.

```
m <- c(45, 23, 66, 77, 33, 44, 56, 12, 78, 23)
is.vector(m) #checking if vector or not
```

```
[1] TRUE
```

```
is.matrix(m) #checking if it's a matrix or not
```

```
[1] FALSE
```



```
is.array(m) #checking of it's a array
```

```
[1] FALSE
```

```
dim(m) <-c(2, 5)#organizing the vector as matrix of 2*5 dimension  
m
```

```
      [,1] [,2] [,3] [,4] [,5]  
[1,]   45   66   33   56   78  
[2,]   23   77   44   12   23
```

```
is.vector(m) #checking if vector or not
```

```
[1] FALSE
```

```
is.matrix(m) #checking of it's a matrix or not
```

```
[1] TRUE
```

```
is.array(m) #checking of it's a array
```

```
[1] TRUE
```

```
(m <- matrix(c(45, 23, 66, 77, 33, 44, 56, 12, 78, 23), 2, 5, byrow = TRUE)) #If byrow = T
```

```
      [,1] [,2] [,3] [,4] [,5]  
[1,]   45   23   66   77   33  
[2,]   44   56   12   78   23
```

Exercise:

Create a matrix with two columns:

First columns hold age data for a group of students 11, 11, 12, 13, 14, 9, 8, and second columns hold grades 5, 5, 6, 7, 8, 4, 3.

```
test <-matrix(c(11, 11, 12, 13, 14, 9, 8, 5, 5, 6, 7, 8, 4, 3), 7, 2)
test
```

```
      [,1] [,2]
[1,]   11   5
[2,]   11   5
[3,]   12   6
[4,]   13   7
[5,]   14   8
[6,]    9   4
[7,]    8   3
```

```
m <- c(45, 23, 66, 77, 33, 44, 56, 12, 78, 23)
#then 'organize' the vector as a matrix
dim(m) <- c(2, 5)#make the vector a 2 by 5 matrix, 2x5 must = lenght of the vector
m
```

```
      [,1] [,2] [,3] [,4] [,5]
[1,]   45   66   33   56   78
[2,]   23   77   44   12   23
```

```
m[2, 3]#the element at row 2 and column 3
```

```
[1] 44
```

```
(s<- m[2, 1]) # select one value
```

```
[1] 23
```

```
(m<- m [c(1,2), -c(3, 5)]) #select 1st row and 1st, 2nd, and 4th columns: result is a vect
```

```
      [,1] [,2] [,3]
[1,]   45   66   56
[2,]   23   77   12
```

```
(m [1, ]) #select complete row or column: 1st row, result is a vector
```

```
[1] 45 66 56
```

```
(v<-m [, 1]) # 1st column, result is a vector
```

```
[1] 45 23
```

```
is.vector(m) #checking if vector or not
```

```
[1] FALSE
```

```
is.matrix(m) #checking of it's a matrix or not
```

```
[1] TRUE
```

```
is.vector(s) #checking if vector or not
```

```
[1] TRUE
```

```
is.vector(v) #checking if vector or not
```

```
[1] TRUE
```

```
is.matrix(v)
```

```
[1] FALSE
```

if the result of subsetting a matrix is a single row or a single column, it remains as a matrix with one row or one column when `drop = FALSE`.

```
m <- matrix(c(45, 23, 66, 77, 33, 44, 56, 12, 78, 23), 2, 5)
(m<-m [, 2, drop = FALSE])
```

```
      [,1]
[1,]    66
[2,]    77
```

```
is.matrix(m)
```

```
[1] TRUE
```

```
is.vector(m)
```

```
[1] FALSE
```

`cbind()` and `rbind()`: join together two or more vectors or matrices, by column, or by row, respectively:

```
cbind (c(1,2,3), c(4, 5, 6))
```

```
      [,1] [,2]
[1,]     1     4
[2,]     2     5
[3,]     3     6
```

```
rbind (c(1,2,3), c(4, 5, 6))
```

```
      [,1] [,2] [,3]
[1,]     1     2     3
[2,]     4     5     6
```

```
m <- matrix(c(45, 23, 66, 77, 33, 44, 56, 12, 78, 23), 2, 5)
(a <- rbind (c(1,2,3,4,5), m))
```

	[,1]	[,2]	[,3]	[,4]	[,5]
[1,]	1	2	3	4	5
[2,]	45	66	33	56	78
[3,]	23	77	44	12	23

```
is.array(a)
```

```
[1] TRUE
```

```
is.matrix(a)
```

```
[1] TRUE
```

Exercise:

What will m1-m4 look like?

```
m1 <- matrix(rep(10, 9), 3, 3)
m2 <- cbind(c(1,2,3), c(4, 5, 6))
m3 <- cbind(m1[,1], m2[,2])
m4 <- cbind(m1[,1], m2[,2])
# m5 <- m1-m4 # will give an error as arrays that do not have compatible dimensions
```

Named rows and columns in matrix:

```
sales <- matrix(c(10, 30, 40, 50, 43, 56, 21, 30), 2, 4, byrow=TRUE)
colnames(sales) <- c('1qrt', '2qrt', '3qrt', '4qrt')
rownames(sales) <- c('store1', 'store2')
sales
```

	1qrt	2qrt	3qrt	4qrt
store1	10	30	40	50
store2	43	56	21	30

Exercise:

Find store1 1qrt sale. 2. List store2's 1st and 4th quarter sales:

```
sales['store1', '1qrt']
```

```
[1] 10
```

```
sales['store2', c('1qrt', '4qrt')]
```

```
1qrt 4qrt  
43   30
```

Arrays

Arrays are similar to matrices, but arrays can have more than 2 dimensions

```
a <- array(1:48, dim= c(4, 3, 2))  
a
```

```
, , 1
```

	[,1]	[,2]	[,3]
[1,]	1	5	9
[2,]	2	6	10
[3,]	3	7	11
[4,]	4	8	12

```
, , 2
```

	[,1]	[,2]	[,3]
[1,]	13	17	21
[2,]	14	18	22
[3,]	15	19	23
[4,]	16	20	24

```
a [1, 3, 2]
```

```
[1] 21
```

```
a [1, , 2]
```

```
[1] 13 17 21
```

```
a [1, , 2, drop=FALSE]
```

```
, , 1
```

```
      [,1] [,2] [,3]  
[1,]    13    17    21
```

```
a [4, 3, ]
```

```
[1] 12 24
```

```
a [c(2, 3), , -2]
```

```
      [,1] [,2] [,3]  
[1,]     2     6    10  
[2,]     3     7    11
```

```
dimnames(a)[[1]] <-c("1qrt", "2qrt", "3qrt", "4qrt")  
dimnames(a)[[2]] <-c("store1", "store2", "store3")  
dimnames(a)[[3]] <-c("2017", "2018")  
a #using list() to specify names
```

```
, , 2017
```

```
      store1 store2 store3  
1qrt      1      5      9  
2qrt      2      6     10  
3qrt      3      7     11  
4qrt      4      8     12
```

, , 2018

	store1	store2	store3
1qrt	13	17	21
2qrt	14	18	22
3qrt	15	19	23
4qrt	16	20	24

```
ar <- array(data      = 1:27,  
            dim       = c(3, 3, 3),  
            dimnames  = list(c("a", "b", "c"), c("d", "e", "f"), c("g", "h", "i"))  
ar
```

, , g

	d	e	f
a	1	4	7
b	2	5	8
c	3	6	9

, , h

	d	e	f
a	10	13	16
b	11	14	17
c	12	15	18

, , i

	d	e	f
a	19	22	25
b	20	23	26
c	21	24	27

Split array into matrices

```
matrix1 <- ar[,,g]  
matrix1 <- ar[,, 'g']  
matrix1
```



```
  d e f
a 1 4 7
b 2 5 8
c 3 6 9
```

```
matrix2 <- ar[,, 'h']
matrix2
```

```
  d e f
a 10 13 16
b 11 14 17
c 12 15 18
```

```
sum <- matrix1 + matrix2
sum
```

```
  d e f
a 11 17 23
b 13 19 25
c 15 21 27
```

```
matrix1*3 #multiplying all matrix elements by 3
```

```
  d e f
a 3 12 21
b 6 15 24
c 9 18 27
```

```
matrix1*c(2, 3)
```

Warning in matrix1 * c(2, 3): longer object length is not a multiple of shorter object length

```
  d e f
a 2 12 14
b 6 10 24
c 6 18 18
```

```
matrix1*c(2,3,2,3,2,3,2,3,2) #multiplying matrix with an vector
```

```
      d  e  f
a 2 12 14
b 6 10 24
c 6 18 18
```

```
matrix1*c(1, 2, 3)
```

```
      d  e  f
a 1  4  7
b 4 10 16
c 9 18 27
```

```
matrix1/c(1, 2, 3)
```

```
      d    e  f
a 1 4.0 7
b 1 2.5 4
c 1 2.0 3
```

```
matrix1/c(1, 2, 3, 1, 2, 3, 1, 2, 3)
```

```
      d    e  f
a 1 4.0 7
b 1 2.5 4
c 1 2.0 3
```

Lists:

a list is a versatile and flexible data structure that can hold elements of different data types, including vectors, matrices, data frames, functions, and even other lists.

you can access a component of a list by using the **\$** operator, followed by the name of the component you want to access.

```
mylist <- list(stud.id=34453,  
              stud.name="John",  
              stud.marks= c(13, 3, 12, 15, 19)  
              )
```

```
mylist$stud.id
```

```
[1] 34453
```

```
mylist[1] #accessing 1 st element
```

```
$stud.id  
[1] 34453
```

```
mylist[[1]]
```

```
[1] 34453
```

```
mylist["stud.id"]
```

```
$stud.id  
[1] 34453
```

```
handle <- "stud.id" #renaming  
mylist[handle]
```

```
$stud.id  
[1] 34453
```

```
mylist[["stud.id"]] # extracting single elements from a list
```

```
[1] 34453
```

```
mylist <- list(stud.id=34453,  
              stud.name="John",  
              stud.marks= c(13, 3, 12, 15, 19)  
              )  
mylist$stud.marks
```

```
[1] 13  3 12 15 19
```

```
mylist$stud.marks[2]
```

```
[1] 3
```

```
names(mylist)
```

```
[1] "stud.id"    "stud.name"  "stud.marks"
```

```
names(mylist) <- c('id','name','marks') #changing names
```

```
names(mylist)
```

```
[1] "id"    "name"  "marks"
```

```
mylist
```

```
$id
```

```
[1] 34453
```

```
$name
```

```
[1] "John"
```

```
$marks
```

```
[1] 13  3 12 15 19
```

```
mylist$parents.names <- c('Ana', "Mike")
mylist
```

```
$id
[1] 34453
```

```
$name
[1] "John"
```

```
$marks
[1] 13  3 12 15 19
```

```
$parents.names
[1] "Ana"  "Mike"
```

```
newlist <- list(age=19, sex="male");
expandedlist <-c(mylist, newlist) #concatating 2 lists
expandedlist
```

```
$id
[1] 34453
```

```
$name
[1] "John"
```

```
$marks
[1] 13  3 12 15 19
```

```
$parents.names
[1] "Ana"  "Mike"
```

```
$age
[1] 19
```

```
$sex
[1] "male"
```

```
length(expandedlist) #length of the new list
```

```
[1] 6
```

Exercise:

Starting with the expanded list given above, what will be the result of the following statement?
Consider the statement one by one.

```
expandedlist <- expandedlist[-5]
expandedlist <- expandedlist[c(-1,-5)]
expandedlist$parents.names <- NULL
expandedlist[['marks']] <- NULL
```

```
mylist
```

```
$id
```

```
[1] 34453
```

```
$name
```

```
[1] "John"
```

```
$marks
```

```
[1] 13  3 12 15 19
```

```
$parents.names
```

```
[1] "Ana" "Mike"
```

```
unlist(mylist) # convert to a one-dimensional vector
```

id	name	marks1	marks2	marks3
"34453"	"John"	"13"	"3"	"12"
marks4	marks5	parents.names1	parents.names2	
"15"	"19"	"Ana"	"Mike"	

```
mode(mylist) #finding the type
```

```
[1] "list"
```

```
mode(unlist(mylist))
```

```
[1] "character"
```

```
is.vector(unlist(mylist)) #atomic list with names
```

```
[1] TRUE
```

```
is.list(mylist)
```

```
[1] TRUE
```

```
is.atomic(mylist)
```

```
[1] FALSE
```

```
is.list(unlist(mylist))
```

```
[1] FALSE
```

Data Frames

A data frame is a widely used data structure for organizing and manipulating tabular data.

```
my.dataframe <- data.frame(site=c('A', 'B', 'A','A', 'B'),  
                           season=c('winter', 'summer', 'summer', 'spring', 'fall'),  
my.dataframe
```

	site	season	ph
1	A	winter	7.4
2	B	summer	6.3
3	A	summer	8.6
4	A	spring	7.2
5	B	fall	8.9

Exercise:

Given 'my.dataframes', what values will the following statements access?

```
my.dataframe <- data.frame(site=c('A', 'B', 'A','A', 'B'),  
                           season=c('winter', 'summer', 'summer', 'spring', 'fall'),  
my.dataframe[3, 2]
```

```
[1] "summer"
```

```
my.dataframe[['site']]
```

```
[1] "A" "B" "A" "A" "B"
```

```
my.dataframe['site']
```

```
site  
1    A  
2    B  
3    A  
4    A  
5    B
```

```
my.dataframe[my.dataframe$ph>7, ]
```

```
site season  ph  
1    A winter 7.4  
3    A summer 8.6  
4    A spring 7.2  
5    B  fall  8.9
```

```
my.dataframe[my.dataframe$ph>7, 'site']
```

```
[1] "A" "A" "A" "B"
```



```
my.dataframe[my.dataframe$ph>7, c('site', 'ph')]
```

```

site ph
1    A 7.4
3    A 8.6
4    A 7.2
5    B 8.9

```

subset() function in R is used to create a subset of a data frame or matrix based on specified conditions or criteria.

```
subset(my.dataframe, ph>7)
```

```

site season ph
1    A winter 7.4
3    A summer 8.6
4    A spring 7.2
5    B   fall 8.9

```

```
subset(my.dataframe, ph>7, c("site", "ph"))
```

```

site ph
1    A 7.4
3    A 8.6
4    A 7.2
5    B 8.9

```

```
subset(my.dataframe[1:2,], ph>7, c(site, ph))
```

```

site ph
1    A 7.4

```

```
my.dataframe[my.dataframe$season=='summer', 'ph'] <- my.dataframe[my.dataframe$season=='summer', 'ph']
my.dataframe[my.dataframe$season=='summer', 'ph']
```

```
[1] 7.3 9.6
```

```
my.dataframe[my.dataframe$season=='summer' & my.dataframe$ph>8, 'ph'] <- my.dataframe[my.d

my.dataframe[my.dataframe$season=='summer', 'ph']
```

```
[1] 7.3 10.6
```

```
my.dataframe$N03 <- c(234.5, 123.4, 456.7, 567.8, 789.0)
my.dataframe
```

```
  site season  ph  N03
1    A winter  7.4 234.5
2    B summer  7.3 123.4
3    A summer 10.6 456.7
4    A spring  7.2 567.8
5    B  fall  8.9 789.0
```

```
#my.dataframe$N03<-NULL
my.dataframe <- my.dataframe[, -4]
my.dataframe
```

```
  site season  ph
1    A winter  7.4
2    B summer  7.3
3    A summer 10.6
4    A spring  7.2
5    B  fall  8.9
```

```
str(my.dataframe)
```

```
'data.frame': 5 obs. of 3 variables:
 $ site : chr  "A" "B" "A" "A" ...
 $ season: chr  "winter" "summer" "summer" "spring" ...
 $ ph : num  7.4 7.3 10.6 7.2 8.9
```

```
nrow(my.dataframe) # number of rows in dataframe
```

```
[1] 5
```

```
ncol(my.dataframe) #number of columns in dataframes
```

```
[1] 3
```

```
dim(my.dataframe)
```

```
[1] 5 3
```

```
#edit(my.dataframe) #this brings up a data editor
```

```
#View(my.dataframe) #this brings up a uneditable tab that display the data for you to view
```

```
names(my.dataframe)
```

```
[1] "site"    "season" "ph"
```

```
names(my.dataframe) <- c('area', 'season', 'P.h.')
my.dataframe
```

```
area season P.h.
1    A winter  7.4
2    B summer  7.3
3    A summer 10.6
4    A spring  7.2
5    B  fall   8.9
```

```
names(my.dataframe)[3] <- 'ph'
my.dataframe
```

```
area season  ph
1    A winter  7.4
2    B summer  7.3
3    A summer 10.6
4    A spring  7.2
5    B  fall   8.9
```

Tibbles

Tibbles are designed to make data manipulation and analysis more intuitive and less error-prone.

```
install.packages("tibble")
```

Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
(as 'lib' is unspecified)

```
library(tibble)
```

```
my.tibble <- tibble(TempCels = sample(-10:40, size=100, replace=TRUE),  
                    TempFahr = TempCels*9/5+32,  
                    Location = rep(letters[1:2], each=50))  
my.tibble
```

```
# A tibble: 100 x 3  
  TempCels TempFahr Location  
    <int>    <dbl> <chr>  
1      24     75.2 a  
2       7     44.6 a  
3      32     89.6 a  
4      -9     15.8 a  
5       3     37.4 a  
6      23     73.4 a  
7      10     50 a  
8      19     66.2 a  
9       1     33.8 a  
10     16     60.8 a  
# i 90 more rows
```

```
install.packages("palmerpenguins")
```

Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
(as 'lib' is unspecified)

```
library(palmerpenguins)
data(penguins)
dim(penguins)
```

```
[1] 344    8
```

```
class(penguins)
```

```
[1] "tbl_df"      "tbl"        "data.frame"
```

```
penguins
```

```
# A tibble: 344 x 8
  species island bill_length_mm bill_depth_mm flipper_length_mm body_mass_g
  <fct>   <fct>         <dbl>         <dbl>           <int>         <int>
1 Adelie Torgersen     39.1           18.7             181          3750
2 Adelie Torgersen     39.5           17.4             186          3800
3 Adelie Torgersen     40.3           18              195          3250
4 Adelie Torgersen     NA              NA              NA           NA
5 Adelie Torgersen     36.7           19.3             193          3450
6 Adelie Torgersen     39.3           20.6             190          3650
7 Adelie Torgersen     38.9           17.8             181          3625
8 Adelie Torgersen     39.2           19.6             195          4675
9 Adelie Torgersen     34.1           18.1             193          3475
10 Adelie Torgersen     42            20.2             190          4250
# i 334 more rows
# i 2 more variables: sex <fct>, year <int>
```

```
pe <-as_tibble(penguins)
class(pe)
```

```
[1] "tbl_df"      "tbl"        "data.frame"
```

```
pe
```

```
# A tibble: 344 x 8
  species island  bill_length_mm bill_depth_mm flipper_length_mm body_mass_g
  <fct>   <fct>         <dbl>         <dbl>         <int>         <int>
1 Adelie  Torgersen      39.1          18.7          181          3750
2 Adelie  Torgersen      39.5          17.4          186          3800
3 Adelie  Torgersen      40.3           18          195          3250
4 Adelie  Torgersen      NA            NA            NA            NA
5 Adelie  Torgersen      36.7          19.3          193          3450
6 Adelie  Torgersen      39.3          20.6          190          3650
7 Adelie  Torgersen      38.9          17.8          181          3625
8 Adelie  Torgersen      39.2          19.6          195          4675
9 Adelie  Torgersen      34.1          18.1          193          3475
10 Adelie Torgersen      42            20.2          190          4250
# i 334 more rows
# i 2 more variables: sex <fct>, year <int>
```

```
x <- 1:16
mode(x) #finding type of frame
```

```
[1] "numeric"
```

```
dim(x) <- c(4,4)
class(x)
```

```
[1] "matrix" "array"
```

```
is.numeric(x)
```

```
[1] TRUE
```

```
mode(x) <- "character"
mode(x)
```

```
[1] "character"
```

```
class(x)
```

```
[1] "matrix" "array"
```

```
x <- factor(x)  
class(x)
```

```
[1] "factor"
```

```
mode(x)
```

```
[1] "numeric"
```

```
is.array(x)
```

```
[1] FALSE
```

```
is.data.frame(x)
```

```
[1] FALSE
```

```
is.matrix(x)
```

```
[1] FALSE
```

```
is_tibble(x)
```

```
[1] FALSE
```

```
is.vector(x)
```

```
[1] FALSE
```

```
typeof(x)
```

```
[1] "integer"
```

```
class(pe[1:15, c("bill_length_mm", "bill_depth_mm")]) #subsetting tibble in smaller one
```

```
[1] "tbl_df"      "tbl"        "data.frame"
```

```
class(penguins[1:15, c("bill_length_mm", "bill_depth_mm")])
```

```
[1] "tbl_df"      "tbl"        "data.frame"
```

```
class(pe[1:15, c("bill_length_mm")])
```

```
[1] "tbl_df"      "tbl"        "data.frame"
```

```
class(penguins[1:15, c("bill_length_mm")])
```

```
[1] "tbl_df"      "tbl"        "data.frame"
```

dplyr

Provides a set of functions and a consistent, user-friendly grammar for working with data frames or tibbles.

```
install.packages("dplyr")
```

Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
(as 'lib' is unspecified)

```
library(dplyr)
```


Attaching package: 'dplyr'

The following objects are masked from 'package:stats':

filter, lag

The following objects are masked from 'package:base':

intersect, setdiff, setequal, union

```
select(filter(pe, species=="Adelie"), bill_length_mm, bill_depth_mm) #Select bill lengths
```

A tibble: 152 x 2

	bill_length_mm	bill_depth_mm
	<dbl>	<dbl>
1	39.1	18.7
2	39.5	17.4
3	40.3	18
4	NA	NA
5	36.7	19.3
6	39.3	20.6
7	38.9	17.8
8	39.2	19.6
9	34.1	18.1
10	42	20.2

i 142 more rows

```
filter(select(pe, bill_length_mm, bill_depth_mm, species), species=="Adelie")
```

A tibble: 152 x 3

	bill_length_mm	bill_depth_mm	species
	<dbl>	<dbl>	<fct>
1	39.1	18.7	Adelie
2	39.5	17.4	Adelie
3	40.3	18	Adelie
4	NA	NA	Adelie
5	36.7	19.3	Adelie

```

6          39.3          20.6 Adelie
7          38.9          17.8 Adelie
8          39.2          19.6 Adelie
9          34.1          18.1 Adelie
10         42           20.2 Adelie
# i 142 more rows

```

Exercise

How would you achieve the same result as the above but use tibble subsetting?

```
pe
```

```

# A tibble: 344 x 8
  species island bill_length_mm bill_depth_mm flipper_length_mm body_mass_g
  <fct>   <fct>         <dbl>         <dbl>         <int>         <int>
1 Adelie Torgersen     39.1           18.7           181          3750
2 Adelie Torgersen     39.5           17.4           186          3800
3 Adelie Torgersen     40.3           18            195          3250
4 Adelie Torgersen     NA            NA            NA            NA
5 Adelie Torgersen     36.7           19.3           193          3450
6 Adelie Torgersen     39.3           20.6           190          3650
7 Adelie Torgersen     38.9           17.8           181          3625
8 Adelie Torgersen     39.2           19.6           195          4675
9 Adelie Torgersen     34.1           18.1           193          3475
10 Adelie Torgersen     42            20.2           190          4250
# i 334 more rows
# i 2 more variables: sex <fct>, year <int>

```

```
pe[pe$species=='Adelie', c("bill_length_mm", "bill_depth_mm")]
```

```

# A tibble: 152 x 2
  bill_length_mm bill_depth_mm
  <dbl>         <dbl>
1          39.1           18.7
2          39.5           17.4
3          40.3           18
4           NA            NA
5          36.7           19.3
6          39.3           20.6

```

```

7          38.9          17.8
8          39.2          19.6
9          34.1          18.1
10         42           20.2
# i 142 more rows

```

```
subset(pe, pe$species=='Adelie', c("bill_length_mm", "bill_depth_mm"))
```

```

# A tibble: 152 x 2
  bill_length_mm bill_depth_mm
      <dbl>         <dbl>
1          39.1          18.7
2          39.5          17.4
3          40.3          18
4           NA          NA
5          36.7          19.3
6          39.3          20.6
7          38.9          17.8
8          39.2          19.6
9          34.1          18.1
10         42           20.2
# i 142 more rows

```

```
select(pe, bill_length_mm, bill_depth_mm, species) |> filter(species=="Adelie")
```

```

# A tibble: 152 x 3
  bill_length_mm bill_depth_mm species
      <dbl>         <dbl> <fct>
1          39.1          18.7 Adelie
2          39.5          17.4 Adelie
3          40.3          18   Adelie
4           NA          NA   Adelie
5          36.7          19.3 Adelie
6          39.3          20.6 Adelie
7          38.9          17.8 Adelie
8          39.2          19.6 Adelie
9          34.1          18.1 Adelie
10         42           20.2 Adelie
# i 142 more rows

```

```
filter(pe, species=="Adelie") |> select(bill_length_mm, bill_depth_mm, species)
```

```
# A tibble: 152 x 3
  bill_length_mm bill_depth_mm species
      <dbl>         <dbl> <fct>
1         39.1         18.7 Adelie
2         39.5         17.4 Adelie
3         40.3          18  Adelie
4          NA          NA  Adelie
5         36.7         19.3 Adelie
6         39.3         20.6 Adelie
7         38.9         17.8 Adelie
8         39.2         19.6 Adelie
9         34.1         18.1 Adelie
10        42          20.2 Adelie
# i 142 more rows
```

Exercise

Create a data object to hold student names (Judy, Max, Dan) and their grades (78,85,99)
Convert number grades to letter grades:90-100:A;80-89:B;70-79:C; \<70:F'

```
students <- list(names=c("Judy", "Max", "Dan"),
                  grades=c(78, 85, 99))
print ("before:")
```

```
[1] "before:"
```

```
students
```

```
$names
[1] "Judy" "Max"  "Dan"
```

```
$grades
[1] 78 85 99
```

```

gradeConvertor<- function (grade){
  grade = as.numeric(grade)
  if(grade > 100 | grade < 0) print ("grade out of the range")
  else if(grade >= 90 & grade <= 100) return ("A")
  else if(grade >= 80 & grade < 90) return ("B")
  else if(grade >= 70 & grade < 80) return ("C")
  else return ("F")
} # providing different conditions

#students$grades <-sapply(students$grades, gradeConvertor)

for(i in 1:length(students$grades)){
  students$grades[i] = gradeConvertor(students$grades[i])
}

print ("after:")

```

```
[1] "after:"
```

```
students
```

```
$names
```

```
[1] "Judy" "Max"  "Dan"
```

```
$grades
```

```
[1] "C" "B" "A"
```