

R basics

SIT22003: Introduction to Data Science
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Conditionals - If Statement

```
if (condition) {  
  expr  
}
```

```
# Variables related to your last day of recordings  
medium <- "LinkedIn"  
num_views <- 14
```

```
      true  
if (medium == "LinkedIn") {  
  print("Showing LinkedIn information")  
}
```

```
## [1] "Showing LinkedIn information"
```

```
      false  
if (num_views > 15) {  
  print("You're popular!")  
}
```

No output

Conditionals - If Statement

Variables related to your last day of recordings

```
medium <- "LinkedIn"
```

```
num_views <- 14
```

true

```
if (medium == "LinkedIn") {  
  print("Showing LinkedIn information") <-  
} else {  
  print("Unknown medium")  
}
```

```
## [1] "Showing LinkedIn information"
```

```
if (num_views > 15) {  
  print("You're popular!")  
} else {  
  print("Try to be more visible!") <-  
}
```

```
## [1] "Try to be more visible!"
```

```
if (condition) {  
  expr1  
} else {  
  expr2  
}
```

For Loop

```
for(var in seq) {  
    expr  
}
```

```
cities <- c("New York", "Paris",  
            "London", "Tokyo",  
            "Rio de Janeiro", "Cape  
Town")  
for(city in cities) {  
    print(city)  
}  
  
## [1] "New York"  
## [1] "Paris"  
## [1] "London"  
## [1] "Tokyo"  
## [1] "Rio de Janeiro"  
## [1] "Cape Town"
```

Vectorized Operation

```
numbers_vector
```

```
## [1] 1 3 4 2 6 8 7 5
```

```
numbers_even_odd
```

```
## [1] "odd" "odd" "even" "even" "even" "even" "odd" "odd"
```

How could you **numbers_even_odd** vector from **numbers_vector**?

Maybe for-loop with if command?

Not a good idea in R

Vectorized Operation

```
numbers_vector <- c(1,3,4,2,6,8,7,5)

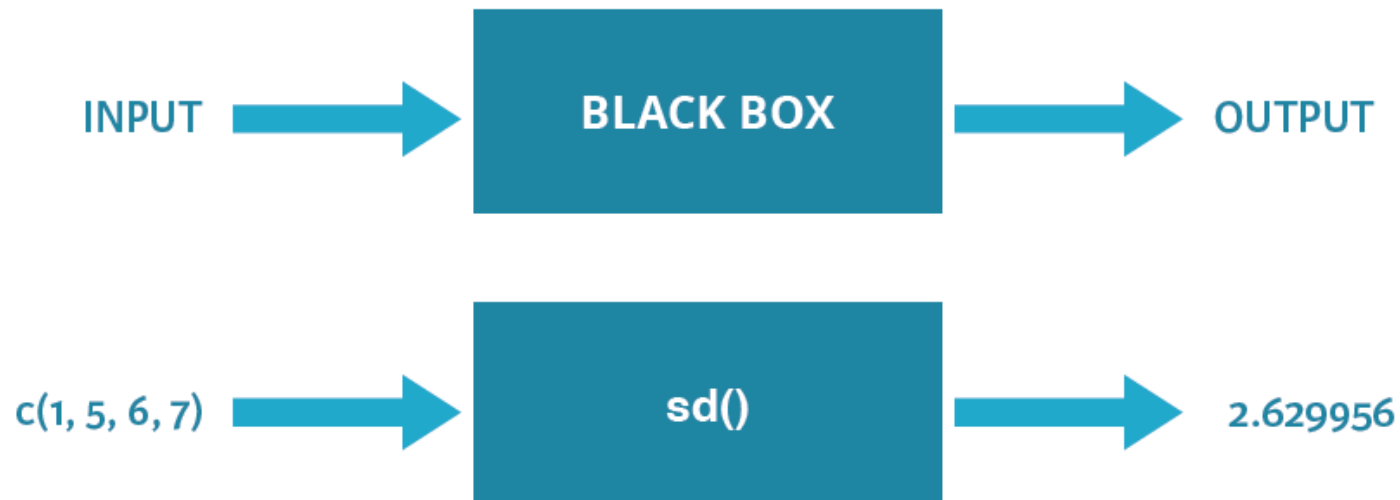
numbers_even_odd <- ifelse(numbers_vector %% 2 == 0, 'even', 'odd')

numbers_even_odd
## [1] "odd"  "odd"  "even" "even" "even" "even" "odd"  "odd"
table(numbers_even_odd)
## numbers_even_odd
## even  odd
##    4    4
```

Easier! and Much More Efficient!

Function

- Function is an automated routine of task
 - Print, mean, max, min are all functions



User-defined Functions

```
cube <- function(n) {  
  return(n*n*n)  
}
```

```
cube(10)
```

```
## [1] 1000
```

```
cube(1:5)
```

```
## [1] 1 8 27 64 125
```

User-defined Functions

```
is.even.number <- function(n) {  
  n %% 2 == 0  
}
```

```
is.even.number(10)
```

```
## [1] TRUE
```

```
is.even.number(5)
```

```
## [1] FALSE
```

```
is.even.number(c(1,2,5,6,7,9,15))
```

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

```
numbers_vector <- c(1,3,4,2,6,8,7,5)
```

```
ifelse(is.even.number(numbers_vector), 'even', 'odd')
```

```
## [1] "odd" "odd" "even" "even" "even" "even" "odd" "odd"
```

User-defined Functions

```
is.even.number <- function(n) {  
  n %% 2 == 0  
}
```

```
is.even.number(10)
```

```
## [1] TRUE
```

```
is.even.number(5)
```

```
## [1] FALSE
```

```
is.even.number(c(1,2,5,6,7,9,15))
```

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

```
numbers_vector <- c(1,3,4,2,6,8,7,5)
```

```
ifelse(is.even.number(numbers_vector), 'even', 'odd')
```

```
## [1] "odd" "odd" "even" "even" "even" "even" "odd" "odd"
```

User-defined Functions

```
diff.max.min <- function(...) {  
  a <- c(...)  
  largest <- max(a)  
  smallest <- min(a)  
  
  largest - smallest  
}
```

```
diff.max.min(6,5,6,23,4,25)
```

```
## [1] 21
```

```
diff.max.min(-55, 100, 23, -7)
```

```
## [1] 155
```

Vectorized operations

- Most of R operations can be applied to both a vector and scalar

```
my.vector <- c(1,3,5,8,13)
my.vector * 2
## [1]  2  6 10 16 26
my.vector >= 5
## [1] FALSE FALSE  TRUE  TRUE  TRUE
my.vector < 10
## [1]  TRUE  TRUE  TRUE  TRUE FALSE
my.vector >= 5 & my.vector < 10
## [1] FALSE FALSE  TRUE  TRUE FALSE
```

Vectorized operations

- Most of R operations can be applied to both a vector and scalar

**try not to use “for-loop”
use vectorized operations instead!**

```
my.vector
## [1] 1 3 5 8 13
sum(my.vector)
## [1] 30
mean(my.vector)
## [1] 6
median(my.vector)
## [1] 5
min(my.vector)
## [1] 1
max(my.vector)
## [1] 13
```

```
summary(my.vector)
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##         1         3         5         6         8        13

ifelse(my.vector %% 2 == 0, 'even', 'odd')
## [1] "odd"  "odd"  "odd"  "even" "odd"
```

Importing Data in to R

- What to import?

- 5 Types

- Flat files



- Data from Excel



- Databases




- Web



- Statistical software



Flat Files

 states.csv

Comma Separated Values

state,capital,pop_mill,area_sqm
South Dakota,Pierre,0.853,77116
New York,Albany,19.746,54555
Oregon,Salem,3.970,98381
Vermont,Montpelier,0.627,9616
Hawaii,Honolulu,1.420,10931

← Field names

> wanted_df

	state	capital	pop_mill	area_sqm
1	South Dakota	Pierre	0.853	77116
2	New York	Albany	19.746	54555
3	Oregon	Salem	3.970	98381
4	Vermont	Montpelier	0.627	9616
5	Hawaii	Honolulu	1.420	10931



read.csv

The `utils` package, which is automatically loaded in your R session on startup, can import CSV files with the `read.csv()` function.

“`swimming_pools.csv`” contains data on swimming pools in Brisbane, Australia (Source: **data.gov.au**). The file contains the column names in the first row. It uses a comma to separate values within rows.

```
# Import swimming_pools.csv: pools
pools <- read.csv('swimming_pools.csv')

# Print the structure of pools
str(pools)

## 'data.frame':    20 obs. of  4 variables:
##  $ Name      : Factor w/ 20 levels "Acacia Ridge Leisure Centre",...: 1 2 3 4 5 6 19
##  $ Address   : Factor w/ 20 levels "1 Fairlead Crescent, Manly",...: 5 20 18 10 9 11
##  $ Latitude  : num  -27.6 -27.6 -27.6 -27.5 -27.4 ...
##  $ Longitude : num  153 153 153 153 153 ...
```

read.csv stringsAsFactor

With `stringsAsFactors`, you can tell R whether it should convert strings in the flat file to factors.

```
# Import swimming_pools.csv correctly: pools
pools <- read.csv('swimming_pools.csv', stringsAsFactors = FALSE)

# Check the structure of pools
str(pools)

## 'data.frame':    20 obs. of  4 variables:
##  $ Name      : chr  "Acacia Ridge Leisure Centre" "Bellbowrie Pool" "Carole Park"
##              "Centenary Pool (inner City)" ...
##  $ Address   : chr  "1391 Beaudesert Road, Acacia Ridge" "Sugarwood Street, Bellbowrie"
##              "Cnr Boundary Road and Waterford Road Wacol" "400 Gregory Terrace, Spring Hill" ...
##  $ Latitude  : num  -27.6 -27.6 -27.6 -27.5 -27.4 ...
##  $ Longitude: num  153 153 153 153 153 ...
```

Converting Types Afterward

```
# Import swimming_pools.csv: pools
pools <- read.csv('swimming_pools.csv')

pools$Name <- as.character(pools$Name)
pools$Address <- as.character(pools$Address)

# Print the structure of pools
str(pools)

## 'data.frame':    20 obs. of  4 variables:
##  $ Name      : chr  "Acacia Ridge Leisure Centre" "Bellbowrie Pool" "Carole Park"
##              "Centenary Pool (inner City)" ...
##  $ Address   : chr  "1391 Beaudesert Road, Acacia Ridge" "Sugarwood Street,
##              Bellbowrie" "Cnr Boundary Road and Waterford Road Wacol" "400 Gregory Terrace, Spring
##              Hill" ...
##  $ Latitude  : num  -27.6 -27.6 -27.6 -27.5 -27.4 ...
##  $ Longitude: num  153 153 153 153 153 ...
```

read.table

- Read any tabular file as a data frame
- Number of arguments is huge

 states2.txt

```
state/capital/pop_mill/area_sqm
South Dakota/Pierre/0.853/77116
New York/Albany/19.746/54555
Oregon/Salem/3.970/98381
Vermont/Montpelier/0.627/9616
Hawaii/Honolulu/1.420/10931
```

```
> read.table("states2.txt",
             header = TRUE,    first row lists variable names (default FALSE)
             sep = "/",        field separator is a forward slash
             stringsAsFactors = FALSE)
```

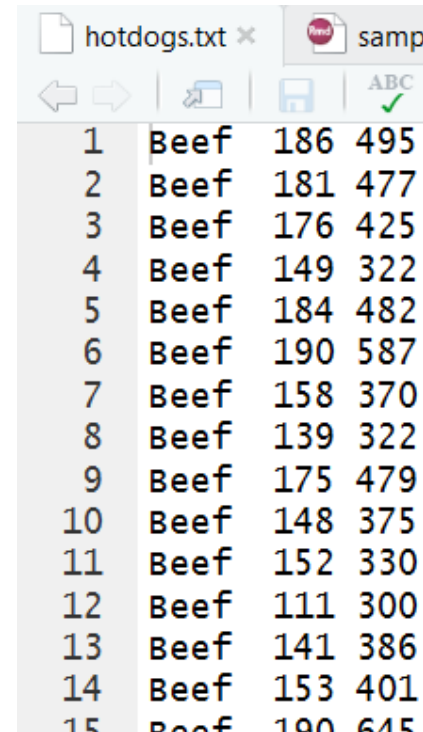
	state	capital	pop_mill	area_sqm
1	South Dakota	Pierre	0.853	77116
2	New York	Albany	19.746	54555
3	Oregon	Salem	3.970	98381
4	Vermont	Montpelier	0.627	9616
5	Hawaii	Honolulu	1.420	10931

reading hotdog.txt

[hotdogs.txt](#), contains information on sodium and calorie levels in different hotdogs (Source: **UCLA**).

The dataset has 3 variables, but the variable names are not available in the first line of the file.

The file uses [tabs](#) as field [separators](#).



1	Beef	186	495
2	Beef	181	477
3	Beef	176	425
4	Beef	149	322
5	Beef	184	482
6	Beef	190	587
7	Beef	158	370
8	Beef	139	322
9	Beef	175	479
10	Beef	148	375
11	Beef	152	330
12	Beef	111	300
13	Beef	141	386
14	Beef	153	401
15	Beef	100	615

reading hotdog.txt

```
hotdogs <- read.table('hotdogs.txt',  
                      sep = '\\t')
```

```
# Call head() on hotdogs  
head(hotdogs)
```

```
##      V1  V2  V3  
## 1 Beef 186 495  
## 2 Beef 181 477  
## 3 Beef 176 425  
## 4 Beef 149 322  
## 5 Beef 184 482  
## 6 Beef 190 587
```

reading hotdog.txt

```
hotdogs <- read.table('hotdogs.txt',  
                      sep = '\\t',  
                      col.names = c("type", "calories", "sodium"))
```

Call head() on hotdogs

```
head(hotdogs)
```

##	type	calories	sodium
## 1	Beef	186	495
## 2	Beef	181	477
## 3	Beef	176	425
## 4	Beef	149	322
## 5	Beef	184	482
## 6	Beef	190	587

Exporting Data Frame as csv file


write.csv function

```
# add new variable named "cal.type"
```

```
hotdogs$cal.type <- ifelse(hotdogs$calories >= 150, 'heavy',  
'light')
```

```
head(hotdogs)
```

```
##      type calories sodium cal.type  
## 1 Beef      186     495    heavy  
## 2 Beef      181     477    heavy  
## 3 Beef      176     425    heavy  
## 4 Beef      149     322    light  
## 5 Beef      184     482    heavy  
## 6 Beef      190     587    heavy
```



row names

```
write.csv(hotdogs, "newhotdog.csv", row.names = F)
```


Exporting Data Frame as tsv file

`write.table` function

```
write.table(hotdogs, "newhotdog.tsv",  
            row.names = F, sep = '\t')
```

"type"	"calories"	"sodium"	"cal.type"
"Beef"	186	495	"heavy"
"Beef"	181	477	"heavy"
"Beef"	176	425	"heavy"
"Beef"	149	322	"light"
"Beef"	184	482	"heavy"
"Beef"	190	587	"heavy"
"Beef"	158	370	"heavy"
"Beef"	139	322	"light"
"Beef"	175	479	"heavy"
"Beef"	148	375	"light"
"Beef"	152	330	"heavy"
"Beef"	111	300	"light"
"Beef"	141	386	"light"
"Beef"	153	401	"heavy"
"Beef"	190	645	"heavy"
"Beef"	157	440	"heavy"
"Beef"	131	317	"light"
"Beef"	149	319	"light"
"Beef"	135	298	"light"
"Beef"	132	253	"light"

Save Your Variables as a File

- Any type of R variables can be saved in RData file
- `save(var1, var2, ..., file = "myfile.RData")`

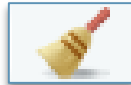
```
my.var <- 10
my.var2 <- c(1,4,6,22,3)
my.var3 <- c('John', 'Bob', 'Alice')
my.var4 <- data.frame(A = 1:3, B = 9:11)

save(my.var, my.var2, my.var3, my.var4,
      file = "myVariables.RData")
```

Clear Your Workspace

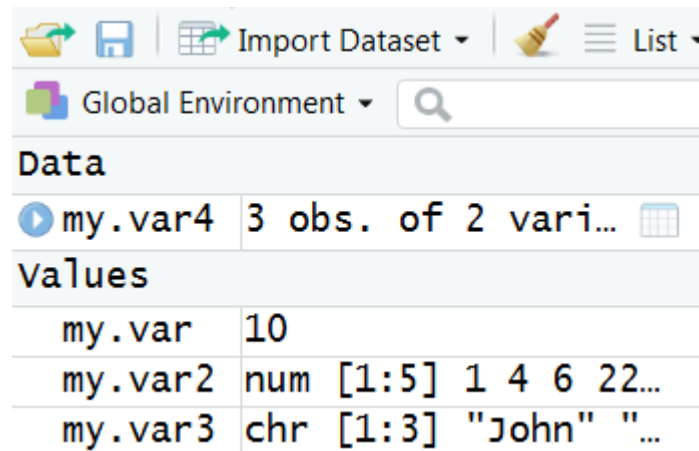
```
rm(list = ls())
```

or press sweep button



Load Your Variables Back

```
load("myVariables.RData")
```



The screenshot shows the RStudio Global Environment pane. At the top, there are icons for file operations and a menu labeled 'Import Dataset'. Below this is a search bar and the text 'Global Environment'. The main area is divided into 'Data' and 'Values' sections. In the 'Data' section, 'my.var4' is listed with '3 obs. of 2 vari...'. In the 'Values' section, three variables are shown: 'my.var' with value '10', 'my.var2' with value 'num [1:5] 1 4 6 22...', and 'my.var3' with value 'chr [1:3] "John" "...'

Data	
my.var4	3 obs. of 2 vari...
Values	
my.var	10
my.var2	num [1:5] 1 4 6 22...
my.var3	chr [1:3] "John" "...

With Rdata files,

You may share your work in R with your colleges or

You may continue to work on the same data on different PC or

You may restore your work from some unexpected errors, etc.

Saving working environment as a file

- To save everything in your working environment

```
save(list = ls(), file = "myWork.RData")
```

or press disk button



The `apply()` Family

- The `apply()` family pertains to the R base package
- It applies functions to manipulate slices of data from `matrices`, `arrays`, `lists` and `dataframes`
- `apply()`, `lapply()` , `sapply()`, `vapply()`, `mapply()`, `rapply()`, and `tapply()`

For More Information:



<https://www.datacamp.com/community/tutorials/r-tutorial-apply-family>

apply()

apply(X, MARGIN, FUN, ...)

- **X** is matrix or dataframe
- **MARGIN** is a variable defining how the function is applied:
 - MARGIN=1, it applies over rows
 - MARGIN=2, it works over columns
- **FUN** is the function that you want to apply to the data

apply()

X		apply(X ,2, sum)					
Dimension — 2							
1 	-1.7189391	-1.0863995	1.0996117	-0.55559727	-0.1792310	-0.8088577	
	-2.2542126	-1.3201873	-2.0533779	1.29055209	0.3264156	0.5412132	
	1.9874737	0.6265486	-0.3684977	1.40028967	-0.7574303	-2.3241569	
	0.2140376	0.8850445	1.4782993	-1.28177703	-0.5015628	1.1537703	
	0.9637687	1.3191502	0.8000988	0.09345943	1.4535431	1.0935720	
Result		sum					
		-0.8078717	0.4241565	0.9561342	0.9469269	0.3417346	-0.3444591

apply()

```
set.seed(2018)
myMat <- matrix(runif(12), ncol = 4)
myMat

##           [,1]      [,2]      [,3]      [,4]
## [1,] 0.33615347 0.1974336 0.6067589 0.5468495
## [2,] 0.46372327 0.4743142 0.1300121 0.3956160
## [3,] 0.06058539 0.3010486 0.9586547 0.6645386

apply(myMat, 1, mean)

## [1] 0.4217989 0.3659164 0.4962068

apply(myMat, 2, mean)

## [1] 0.2868207 0.3242655 0.5651419 0.5356680
```

apply()

alternatively, colSums, rowSums, ...

```
head(iris)
```

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1          5.1          3.5          1.4          0.2 setosa
## 2          4.9          3.0          1.4          0.2 setosa
## 3          4.7          3.2          1.3          0.2 setosa
## 4          4.6          3.1          1.5          0.2 setosa
## 5          5.0          3.6          1.4          0.2 setosa
## 6          5.4          3.9          1.7          0.4 setosa
```

```
apply(iris[, 1:4], 2, mean)
```

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width
##      5.843333      3.057333      3.758000      1.199333
```

```
apply(iris[, 1:4], 1, mean)
```

```
## [1] 2.550 2.375 2.350 2.350 2.550 2.850 2.425 2.525 2.225 2.400 2.700
## [12] 2.500 2.325 2.125 2.800 3.000 2.750 2.575 2.875 2.675 2.675 2.675
.....
```

```
colMeans(iris[, 1:4])
```

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width
##      5.843333      3.057333      3.758000      1.199333
```

lapply()

- It applies function to dataframes, **lists** or vectors
- It gives you back a **list**

```
myList <- list(num = 3.14, chr = "char", logi = TRUE)
myList
```

```
## $num
## [1] 3.14
##
## $chr
## [1] "char"
##
## $logi
## [1] TRUE
```

```
lapply(myList, typeof)
```

```
## $num
## [1] "double"
##
## $chr
## [1] "character"
##
## $logi
## [1] "logical"
```

lapply()

```
myList2 <- list(vec = 1:5, mat = matrix(runif(12), ncol = 4), df = iris)
result <- lapply(myList2, length)
result

## $vec
## [1] 5      # number of elements
##
## $mat
## [1] 12     # number of elements
##
## $df
## [1] 5      # number of columns
unlist(result) # list -> vector

## vec mat  df
##   5  12   5
```

lapply()

```
lapply(c(1,4,9,16), sqrt)
```

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

```
lapply(mtcars, max)
```

```
## $mpg
## [1] 33.9
##
## $cyl
## [1] 8
##
## $disp
## [1] 472
##
## $hp
## [1] 335
##
```

.....

```
unlist(lapply(mtcars, max))
```

```
##      mpg      cyl    disp      hp      drat      wt      qsec      vs      am
## 33.900    8.000 472.000 335.000    4.930    5.424   22.900    1.000    1.000
##      gear      carb
##    5.000    8.000
```

sapply()

- It applies function to dataframes, lists or vectors
- It gives you back a **vector or matrix**

```
sapply(iris[, 1:4], mean)
```

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width
##      5.843333      3.057333      3.758000      1.199333
```

```
sapply(iris, is.numeric)
```

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##           TRUE           TRUE           TRUE           TRUE      FALSE
```

```
sapply(c(1,3,5,7,9), function(x) {x**2})
```

```
## [1]  1  9 25 49 81
```

sapply()

```
myMat <- matrix(1:12, ncol = 4)
```

```
myMat
```

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

```
sapply(myMat, function(x) {x/2})
```

```
## [1] 0.5 1.0 1.5 2.0 2.5 3.0 3.5 4.0 4.5 5.0 5.5 6.0
```

```
sapply(pools, typeof)
```

```
##      Name      Address      Latitude      Longitude  
## "character" "character" "double"  "double"
```

apply()

```
x <- apply(iris[,1:4], function(x) { x > 3 })  
head(x)
```

##		Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
##	[1,]	TRUE	TRUE	FALSE	FALSE
##	[2,]	TRUE	FALSE	FALSE	FALSE
##	[3,]	TRUE	TRUE	FALSE	FALSE
##	[4,]	TRUE	TRUE	FALSE	FALSE
##	[5,]	TRUE	TRUE	FALSE	FALSE
##	[6,]	TRUE	TRUE	FALSE	FALSE

```
colSums(x)
```

##	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
##	150	67	99	0

tapply()

- **tapply(X, GRP_VAR, FUN, ...)**
 - apply **FUN** to **X** after grouping with **GRP_VAR**

```
tapply(iris$Sepal.Length, iris$Species, mean)
```

```
##      setosa versicolor  virginica  
##      5.006      5.936      6.588
```

```
tapply(mtcars$mpg, mtcars$cyl, mean)
```

```
##          4          6          8  
## 26.66364 19.74286 15.10000
```

```
tapply(mtcars$wt, mtcars$mpg>20, mean)
```

```
##      FALSE      TRUE  
## 3.838833 2.418071
```

tapply()

- **tapply(X, GRP_VAR, FUN, ...)**
 - apply **FUN** to **X** after grouping with **GRP_VAR**

```
x <- tapply(mtcars$mpg, mtcars$cyl, function(x){x>20})
x

## $`4`
## [1] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE
##
## $`6`
## [1] TRUE TRUE TRUE FALSE FALSE FALSE FALSE
##
## $`8`
## [1] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [12] FALSE FALSE FALSE

sapply(x, sum)

## 4 6 8
## 11 3 0
```

aggregate()

- **aggregate(var1 ~ var2, data = X, FUN = func, ...)**
 - Apply **func** to **var1** of **X** after grouping by **var2**
 - Alternates to **tapply**
 - Result is **data.frame**

```
aggregate(mpg ~ cyl, data = mtcars, FUN = mean)
```

```
##      cyl      mpg
## 1     4 26.66364
## 2     6 19.74286
## 3     8 15.10000
```

```
aggregate(Sepal.Length ~ Species, data = iris, FUN = mean)
```

```
##      Species Sepal.Length
## 1    setosa      5.006
## 2 versicolor      5.936
## 3 virginica      6.588
```

order() and sort()

- `order()` gives a vector of index of smallest element, second smallest, ..., the largest element
- `sort()` gives a sorted vector of numbers
- `decreasing` option to have result in descending order

```
my_vector <- c(6,12,4,89,23, 35)
order(my_vector)
```

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

```
my_vector[order(my_vector)]
```

```
## [1] 4 6 12 23 35 89
```

```
sort(my_vector)
```

```
## [1] 4 6 12 23 35 89
```

```
my_vector[order(my_vector,decreasing = T)]
```

```
## [1] 89 35 23 12 6 4
```

```
sort(my_vector, decreasing = T)
```

```
## [1] 89 35 23 12 6 4
```

Why Use `order()` instead of `sort()`

- Sorting dataframe

```
mtcars[order(mtcars$mpg, decreasing = T), ]
```

##	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
## Toyota Corolla	33.9	4	71.1	65	4.22	1.835	19.90	1	1	4	1
## Fiat 128	32.4	4	78.7	66	4.08	2.200	19.47	1	1	4	1
## Honda Civic	30.4	4	75.7	52	4.93	1.615	18.52	1	1	4	2
## Lotus Europa	30.4	4	95.1	113	3.77	1.513	16.90	1	1	5	2
## Fiat X1-9	27.3	4	79.0	66	4.08	1.935	18.90	1	1	4	1
## Porsche 914-2	26.0	4	120.3	91	4.43	2.140	16.70	0	1	5	2
## Merc 240D	24.4	4	146.7	62	3.69	3.190	20.00	1	0	4	2
## Datsun 710	22.8	4	108.0	93	3.85	2.320	18.61	1	1	4	1
## Merc 230	22.8	4	140.8	95	3.92	3.150	22.90	1	0	4	2
## Toyota Corona	21.5	4	120.1	97	3.70	2.465	20.01	1	0	3	1
## Hornet 4 Drive	21.4	6	258.0	110	3.08	3.215	19.44	1	0	3	1
## Volvo 142E	21.4	4	121.0	109	4.11	2.780	18.60	1	1	4	2
## Mazda RX4	21.0	6	160.0	110	3.90	2.620	16.46	0	1	4	4
## Mazda RX4 Wag	21.0	6	160.0	110	3.90	2.875	17.02	0	1	4	4
## Ferrari Dino	19.7	6	145.0	175	3.62	2.770	15.50	0	1	5	6
## Merc 280	19.2	6	167.6	123	3.92	3.440	18.30	1	0	4	4
## Pontiac Firebird	19.2	8	400.0	175	3.08	3.845	17.05	0	0	3	2
## Hornet Sportabout	18.7	8	360.0	175	3.15	3.440	17.02	0	0	3	2

References

- Practical Data Science with R, by Nina Zumel and John Mount
- R을 이용한 데이터 분석 실무, 서민구, 길벗