

DR. ALVIN'S PUBLICATIONS

# DATA WRANGLING WITH R

---

BY DR. ALVIN ANG



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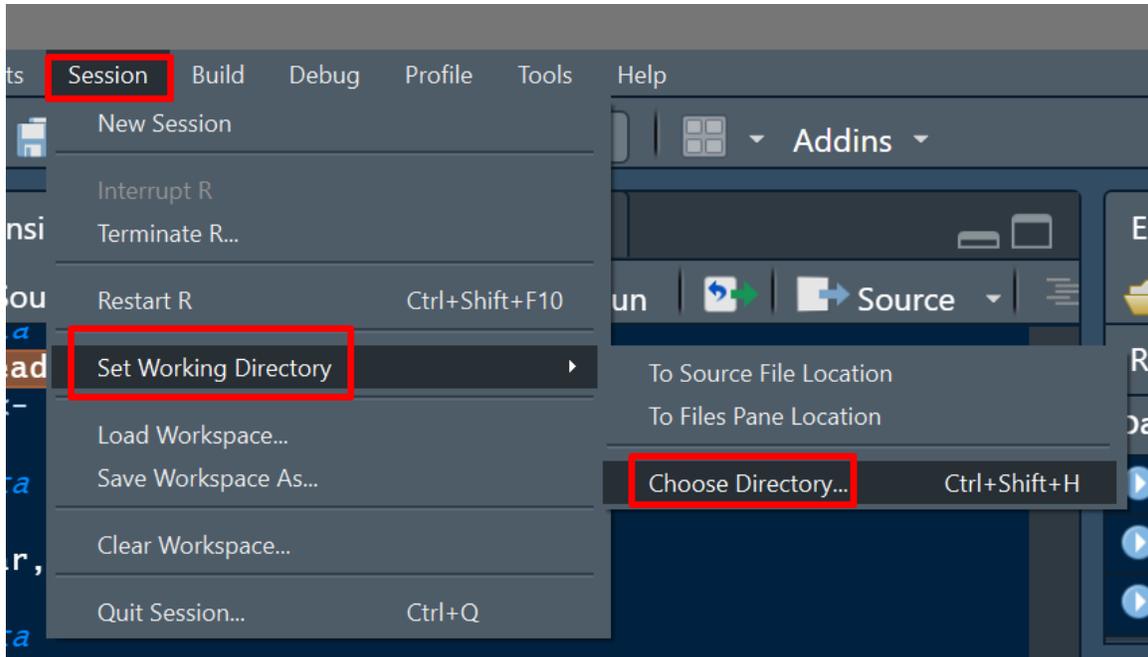
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## I. DATA WRANGLING USING CORE R

---

### A. SET YOUR WORKING DIRECTORY TO YOUR DOWNLOADS FOLDER



#### 1. CHECK YOUR CURRENT WORKING DIRECTORY

```
#-----  
# Data Wrangling using Core R by Dr. Alvin Ang  
#-----  
#1. Check Current Working Directory  
  
getwd()
```

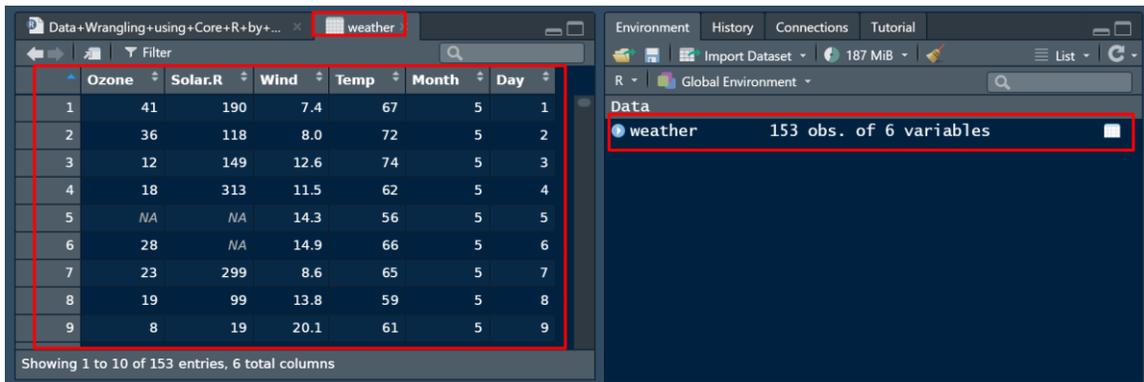
```
> getwd()  
[1] "C:/Users/User/Downloads"
```

## B. WRANGLING WEATHER.CSV

### 1. IMPORT CSV

<https://www.alvinang.sg/s/weather.csv>

```
#-----  
#2. Wrangling Weather.csv  
  
#2a. Import CSV  
#https://www.alvinang.sg/s/weather.csv  
weather = read.csv('weather.csv', header = TRUE)
```



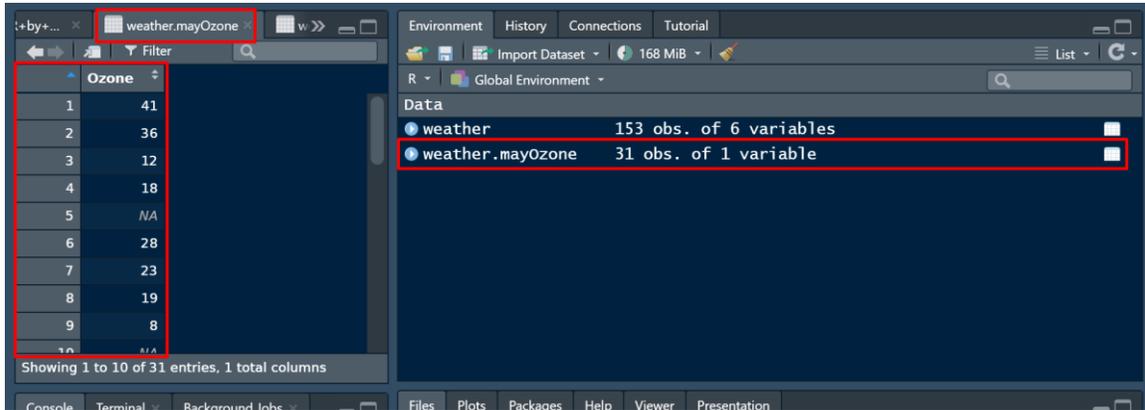
The screenshot displays the RStudio interface. The top-left pane shows the 'weather' dataset imported from a CSV file. The Environment pane on the right shows the 'weather' object with 153 observations and 6 variables. The Data Viewer on the left shows the first 10 rows of the dataset.

	Ozone	Solar.R	Wind	Temp	Month	Day
1	41	190	7.4	67	5	1
2	36	118	8.0	72	5	2
3	12	149	12.6	74	5	3
4	18	313	11.5	62	5	4
5	NA	NA	14.3	56	5	5
6	28	NA	14.9	66	5	6
7	23	299	8.6	65	5	7
8	19	99	13.8	59	5	8
9	8	19	20.1	61	5	9

## 2. SLICE OUT COLUMN USING SUBSET

```
#2b. Slice out column using Subset  
#Slice out the "Ozone" column
```

```
weather.mayOzone <-  
  subset(weather, select=Ozone, subset = Month==5)
```

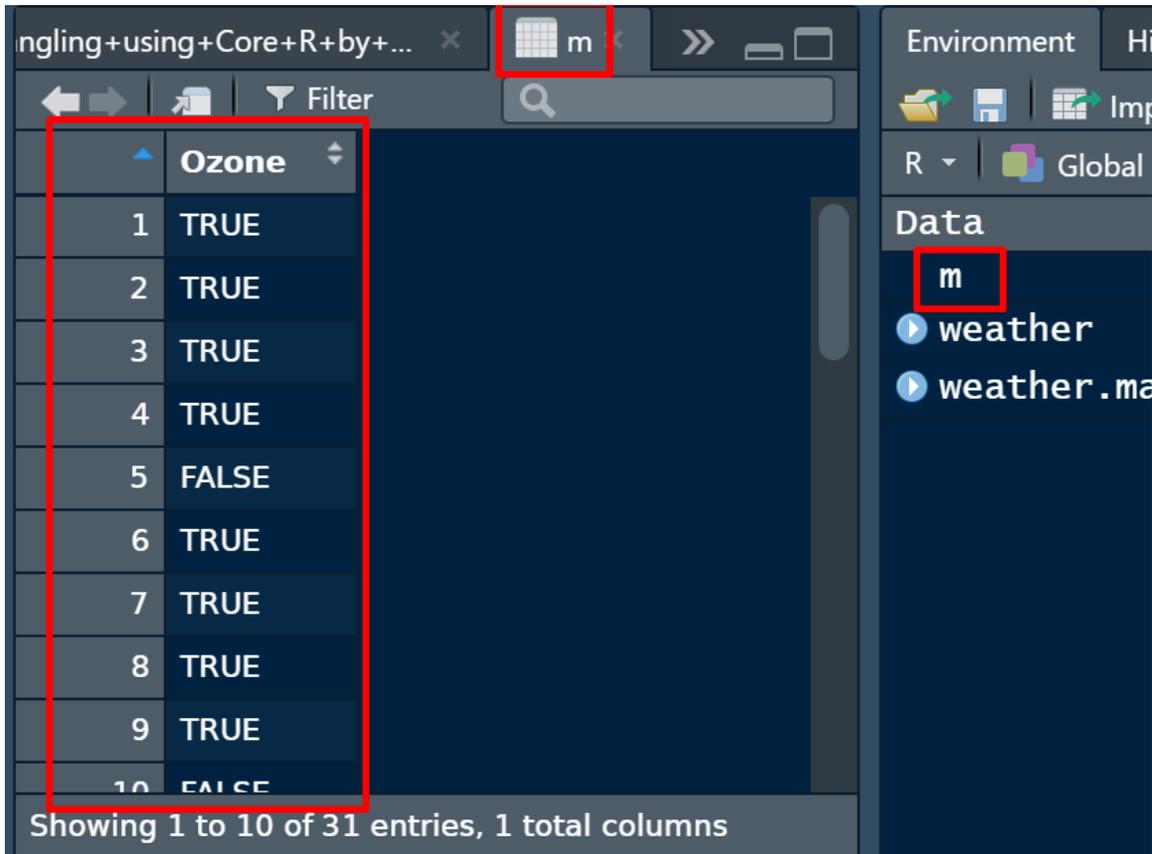


The screenshot displays the RStudio interface. The Environment pane on the right shows the object 'weather.mayOzone' with 31 observations of 1 variable. The Data viewer on the left shows the first 10 rows of the 'Ozone' column, with values ranging from 8 to 41, and a 'NA' value at row 5. The console at the bottom shows the command 'weather.mayOzone <- subset(weather, select=Ozone, subset = Month==5)'.

Row	Ozone
1	41
2	36
3	12
4	18
5	NA
6	28
7	23
8	19
9	8
10	NA

3. CHECK WHICH ROWS HAVE NAs

```
#2c. Check which Rows have NAs  
m = !is.na(weather.mayOzone)
```



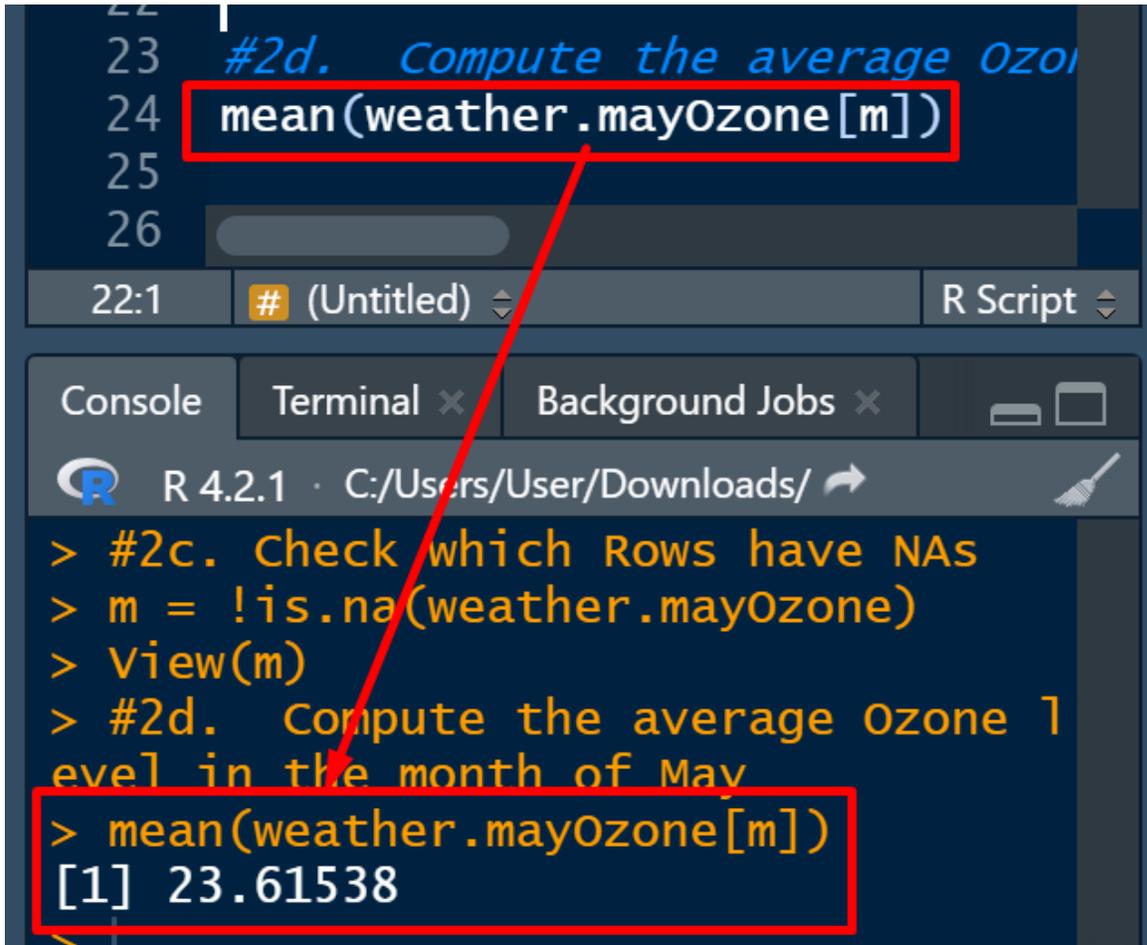
The screenshot shows the RStudio interface. The main window displays a data table with 10 rows and 1 column named 'Ozone'. The values are TRUE, TRUE, TRUE, TRUE, FALSE, TRUE, TRUE, TRUE, TRUE, FALSE. A red box highlights the first 10 rows. The Environment pane on the right shows a variable 'm' highlighted in a red box.

	Ozone
1	TRUE
2	TRUE
3	TRUE
4	TRUE
5	FALSE
6	TRUE
7	TRUE
8	TRUE
9	TRUE
10	FALSE

Showing 1 to 10 of 31 entries, 1 total columns

4. COMPUTE THE AVERAGE OZONE LEVEL IN THE MONTH OF MAY

```
#2d. Compute the average Ozone level in the month of May  
mean(weather.mayOzone[m])
```



```
23 #2d. Compute the average Ozone level in the month of May  
24 mean(weather.mayOzone[m])  
25  
26
```

22:1 # (Untitled) R Script

Console Terminal Background Jobs

R 4.2.1 · C:/Users/User/Downloads/

```
> #2c. Check which Rows have NAs  
> m = !is.na(weather.mayOzone)  
> View(m)  
> #2d. Compute the average Ozone level in the month of May  
> mean(weather.mayOzone[m])  
[1] 23.61538
```

5. FILTER OUT ALL NAs IN THE MONTH OF MAY

```
#2e. Filter out all NAs in the month of May  
a = weather.mayOzone[m]
```

Values

```
a          int [1:26] 41 36 12 18 28 23 19 8 7 16 ...
```

6. OUTPUT AS CSV

```
#2f. output as CSV  
write.csv(a, 'may_weather_data.csv')
```

may\_weather\_data.csv - LibreOffice Calc

File Edit View Insert Format Styles

Liberation Sans 10 pt

A1

	A	B	C	D
1		x		
2	1	41		
3	2	36		
4	3	12		
5	4	18		
6	5	28		
7	6	23		
8	7	19		
9	8	8		
10	9	7		
11	10	16		
12	11	11		
13	12	14		
14	13	18		
15	14	14		
16	15	34		

may\_weather\_data

## C. WRANGLING MTCARS

### 1. SLICING OUT MPG / AM / WT COLUMNS

```
#-----  
#3. Wrangling Mtcars  
  
#3a. Slicing out mpg / am / wt columns  
b = mtcars[c('mpg', 'am', 'wt')]
```

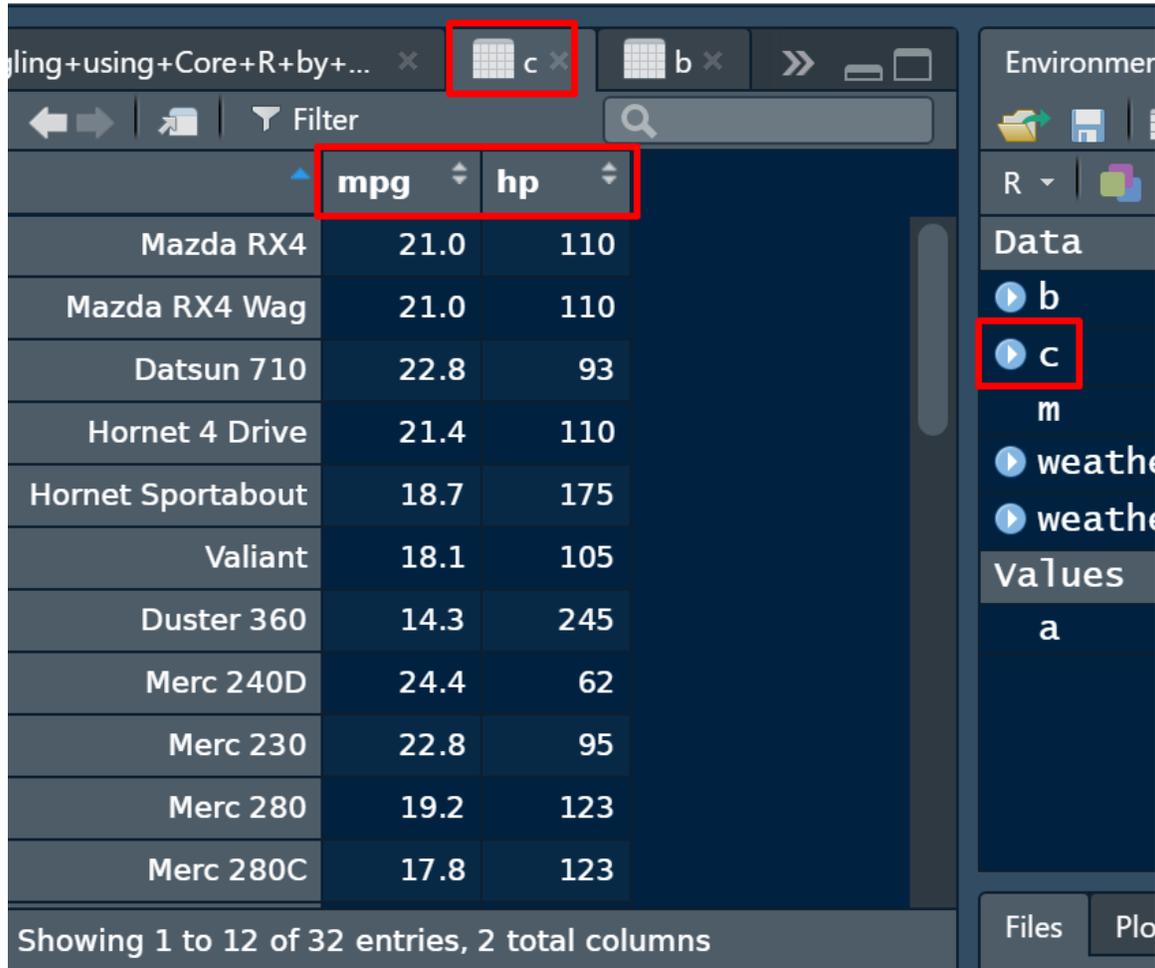
The screenshot shows the RStudio interface. The Environment pane on the right displays the variable 'b' with 32 observations of 3 variables. The Data pane on the left shows a table with columns 'mpg', 'am', and 'wt'. The table contains 11 rows of data, with the first row being Mazda RX4.

	mpg	am	wt
Mazda RX4	21.0	1	2.620
Mazda RX4 Wag	21.0	1	2.875
Datsun 710	22.8	1	2.320
Hornet 4 Drive	21.4	0	3.215
Hornet Sportabout	18.7	0	3.440
Valiant	18.1	0	3.460
Duster 360	14.3	0	3.570
Merc 240D	24.4	0	3.190
Merc 230	22.8	0	3.150
Merc 280	19.2	0	3.440



### 3. SLICING OUT MPG / HP COLUMNS

```
#3c. slicing out mpg / hp columns  
c = subset(mtcars, select=c(mpg, hp))
```

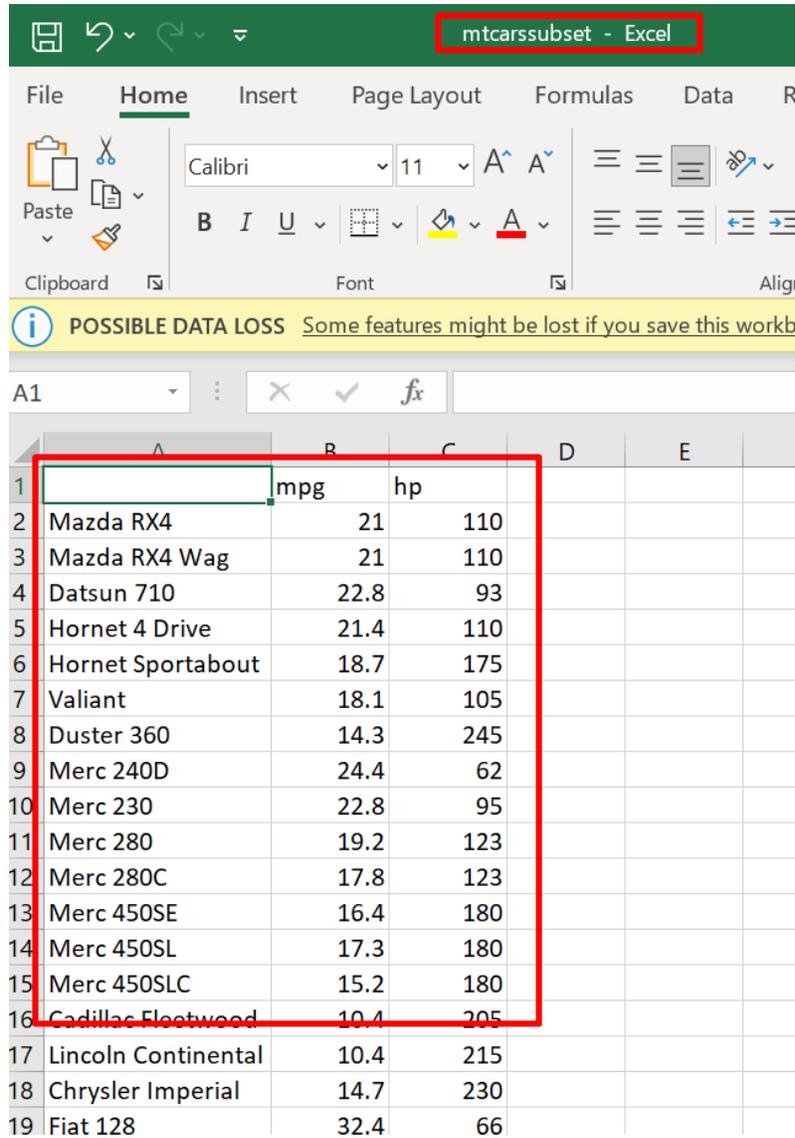


The screenshot shows the RStudio interface. The main window displays a data table with two columns, 'mpg' and 'hp', which are highlighted with a red box. The 'Data' pane on the right shows a list of objects, with 'c' highlighted in red. The status bar at the bottom indicates 'Showing 1 to 12 of 32 entries, 2 total columns'.

	mpg	hp
Mazda RX4	21.0	110
Mazda RX4 Wag	21.0	110
Datsun 710	22.8	93
Hornet 4 Drive	21.4	110
Hornet Sportabout	18.7	175
Valiant	18.1	105
Duster 360	14.3	245
Merc 240D	24.4	62
Merc 230	22.8	95
Merc 280	19.2	123
Merc 280C	17.8	123

#### 4. OUTPUT AS CSV

```
#3d. output as CSV  
write.csv(c, "mtcarssubset.csv")
```



The screenshot shows the Microsoft Excel interface with the 'mtcarssubset - Excel' window. The ribbon is set to 'Home'. A yellow warning banner at the top indicates 'POSSIBLE DATA LOSS: Some features might be lost if you save this workbook'. The active cell is A1. The following table is displayed in the spreadsheet:

	A	B	C	D	E
1		mpg	hp		
2	Mazda RX4	21	110		
3	Mazda RX4 Wag	21	110		
4	Datsun 710	22.8	93		
5	Hornet 4 Drive	21.4	110		
6	Hornet Sportabout	18.7	175		
7	Valiant	18.1	105		
8	Duster 360	14.3	245		
9	Merc 240D	24.4	62		
10	Merc 230	22.8	95		
11	Merc 280	19.2	123		
12	Merc 280C	17.8	123		
13	Merc 450SE	16.4	180		
14	Merc 450SL	17.3	180		
15	Merc 450SLC	15.2	180		
16	Cadillac Fleetwood	10.4	205		
17	Lincoln Continental	10.4	215		
18	Chrysler Imperial	14.7	230		
19	Fiat 128	32.4	66		

5. FILTER ALL THE MPG > 15 AND AM = 1

```
#3e. Filter all the mpg>15 and am=1  
d = mtcars[mtcars$mpg>15 & mtcars$am==1,]
```

The screenshot shows the RStudio interface with a data frame 'd' displayed in the viewer. The data frame contains 10 rows of car data, filtered based on the conditions 'mpg > 15' and 'am == 1'. The columns are mpg, cyl, disp, hp, drat, wt, qsec, vs, am, and gear. The 'am' column is highlighted in red, and the 'mpg' column is also highlighted in red. The 'am == 1' filter is visible in the search bar on the right side of the viewer.

	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear
Mazda RX4	21.0	6	160.0	110	3.90	2.620	16.46	0	1	
Mazda RX4 Wag	21.0	6	160.0	110	3.90	2.875	17.02	0	1	
Datsun 710	22.8	4	108.0	93	3.85	2.320	18.61	1	1	
Fiat 128	32.4	4	78.7	66	4.08	2.200	19.47	1	1	
Honda Civic	30.4	4	75.7	52	4.93	1.615	18.52	1	1	
Toyota Corolla	33.9	4	71.1	65	4.22	1.835	19.90	1	1	
Fiat X1-9	27.3	4	79.0	66	4.08	1.935	18.90	1	1	
Porsche 914-2	26.0	4	120.3	91	4.43	2.140	16.70	0	1	
Lotus Europa	30.4	4	95.1	113	3.77	1.513	16.90	1	1	
Ford Pantera L	15.8	8	351.0	264	4.22	3.170	14.50	0	1	

6. FILTER OUT ONLY MPG AND AM COLUMNS WITH AM = 1 (AUTOMATIC)

```
#3f. Filter out only mpg and am columns with am=1 (automatic)
e = subset(mtcars,
           select=c('mpg', 'am'),
           subset=am==1)
```

The screenshot shows the RStudio interface. The main window displays a data table with 12 rows and 2 columns. The columns are 'mpg' and 'am'. The rows represent different car models. The 'mpg' column contains values ranging from 15.8 to 33.9, and the 'am' column contains the value 1 for all rows. The 'e' object in the Environment pane is highlighted with a red box, and the 'mpg' and 'am' column headers in the table are also highlighted with a red box.

	mpg	am
Mazda RX4	21.0	1
Mazda RX4 Wag	21.0	1
Datsun 710	22.8	1
Fiat 128	32.4	1
Honda Civic	30.4	1
Toyota Corolla	33.9	1
Fiat X1-9	27.3	1
Porsche 914-2	26.0	1
Lotus Europa	30.4	1
Ford Pantera L	15.8	1
Ferrari Dino	19.7	1

Showing 1 to 12 of 13 entries, 2 total columns

## 7. SUMMARY OF MTCARS SUBSET

```
#3g. Summary of mtcars subset  
summary(e)
```

```
> summary(e)  
      mpg      am  
Min.   :15.00  Min.   :1  
1st Qu.:21.00  1st Qu.:1  
Median :22.80  Median :1  
Mean   :24.39  Mean   :1  
3rd Qu.:30.40  3rd Qu.:1  
Max.   :33.90  Max.   :1
```

8. CREATE A TABLE FROM MTCARS AM COLUMNS

```
#3h. Create a Table from Mtcars AM columns  
factor = factor(mtcars$am)  
table(factor)
```

```
#-----  
#THE END  
#-----
```

```
factor          Factor w/ 2 levels "0","1": 2 2 2 1 1 1 1 1 1 1 ...
```

```
> table(factor)  
factor  
 0    1  
19  13
```

---

## II. DATA WRANGLING USING TIDYVERSE

---

<https://www.alvinang.sg/s/Data-Wrangling-with-Tidyverse-by-Dr-Alvin-Ang.R>

Tons of great Data Wrangling with R here:

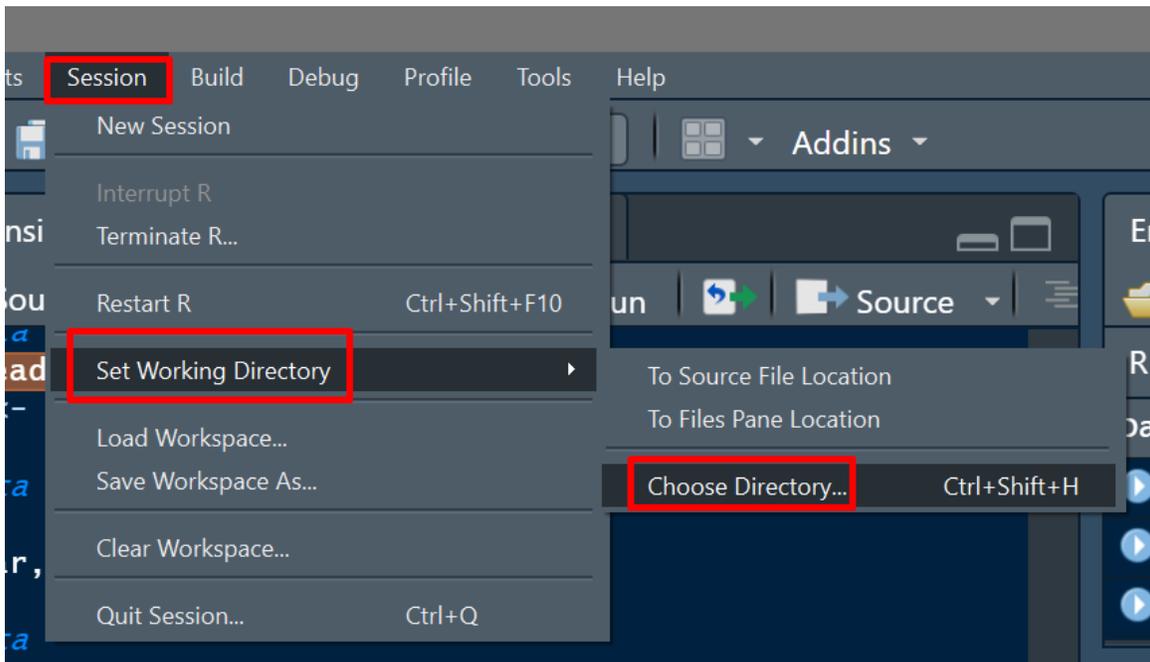
<https://www.marsja.se/how-to-rename-column-or-columns-in-r-with-dplyr/>

### A. INSTALLING TIDYVERSE PACKAGE

```
#-----  
#Data Wrangling with Tidyverse by Dr Alvin Ang  
#-----  
#1. Install Tidyverse and Load Libraries  
  
install.packages("tidyverse", dependencies = TRUE)  
  
library(tidyverse)  
library(tibble)  
library(tidyr)  
library(dplyr)  
library(readxl)  
library(ggplot2)  
library(lubridate)
```

## B. READING IN CSV

File can be found here: <https://www.alvinang.sg/s/dengue.csv>



- Do you know where you stored the dengue.csv downloaded file?
- Most probably is in your download folder.
- Make sure that you set the working directory to that folder (download folder)... so that it can import in the CSV.

```
#-----  
#2. Reading in the Dengue.csv  
dengue <- read_csv("dengue.csv")  
  
#file is here: https://www.alvinang.sg/s/dengue.csv  
  
#or if you want to read in .xls  
# dengue_xls <- read_excel("dengue.xlsx")
```

The screenshot shows the RStudio interface. The Environment pane on the right shows the 'dengue' dataset with 530 observations and 4 variables. The Data Viewer pane on the left shows a preview of the first 6 rows of the dataset.

	year	eweek	type_dengue	number
1	2014	1	Dengue	436
2	2014	1	DHF	1
3	2014	2	Dengue	479
4	2014	2	DHF	0
5	2014	3	Dengue	401
6	2014	3	DHF	0

### C. SELECTING COLUMNS

```
#-----  
#3. Selecting Columns  
  
# select 'year' and 'number' columns from dengue.csv  
a = dengue %>%  
  select(year,number)
```

The screenshot shows the RStudio interface. The 'Data' pane on the right lists two datasets: 'a' (530 obs. of 2 variables) and 'dengue' (530 obs. of 4 variables). A red box highlights the 'a' dataset. A red arrow points from this box to the 'Viewer' pane on the left, which displays a table with two columns: 'year' and 'number'. The table shows the first six rows of data.

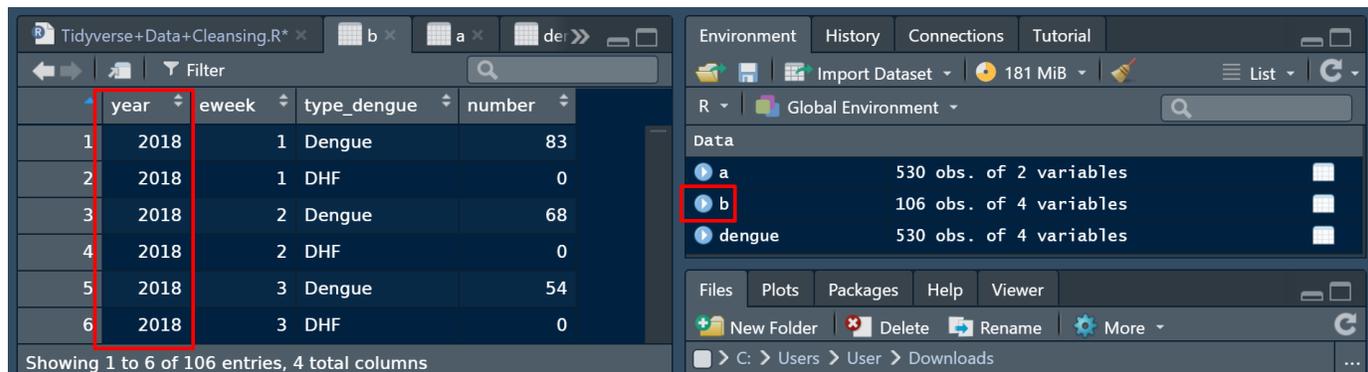
	year	number
1	2014	436
2	2014	1
3	2014	479
4	2014	0
5	2014	401
6	2014	0

Showing 1 to 6 of 530 entries, 2 total columns

## D. FILTERING DATA

### 1. FILTER OUT 'YEAR' == 2018 FROM DENGUE.CSV

```
#-----  
#4. Filter data  
  
#Filter out 'year' == 2018 from dengue.csv  
b = dengue %>%  
  filter(year==2018)
```



The screenshot displays the RStudio interface. The console window shows the following R code and its output:

```
Tidyverse+Data+Cleansing.R* x b a de>>  
Filter  
year eweek type_dengue number  
1 2018 1 Dengue 83  
2 2018 1 DHF 0  
3 2018 2 Dengue 68  
4 2018 2 DHF 0  
5 2018 3 Dengue 54  
6 2018 3 DHF 0  
Showing 1 to 6 of 106 entries, 4 total columns
```

The Environment pane on the right shows the following data frames:

- a: 530 obs. of 2 variables
- b: 106 obs. of 4 variables
- dengue: 530 obs. of 4 variables

The 'b' data frame is highlighted in red in the Environment pane, indicating it is the current active object. The Files pane shows the current directory path: C:\Users\User\Downloads.

## E. FILTER DATA BASED ON MULTIPLE CONDITIONS

### 1. FILTER OUT 2017 AND 2018

```
#-----  
#5. Filter data based on multiple conditions  
#5a. Filter out 2017 and 2018  
c = dengue %>%  
  filter(year==2017 | year==2018 )
```

The screenshot shows the RStudio interface. The main window displays a data table with the following columns: year, week, type\_dengue, and number. The rows shown are 103 to 108. A red box highlights the 'year' column, and a red arrow points from the 'year' column to the 'c' dataset in the Environment pane. The Environment pane shows three datasets: 'a' (530 obs. of 2 variables), 'b' (106 obs. of 4 variables), and 'c' (212 obs. of 4 variables). The 'c' dataset is highlighted with a red box. The Files pane shows the current directory as C:\Users\User\Downloads.

year	week	type_dengue	number
103	2017	52 Dengue	66
104	2017	52 DHF	0
105	2017	53 Dengue	NA
106	2017	53 DHF	NA
107	2018	1 Dengue	83
108	2018	1 DHF	0

## 2. FILTER OUT 2018 AND DENGUE TYPE

```
#5b. Filter out 2018 and 'Dengue' type  
d = dengue %>%  
  filter(year==2018, type_dengue=='Dengue' )
```

The screenshot shows the RStudio interface. On the left, a data table is displayed with columns: year, eweek, type\_dengue, and number. The first six rows are highlighted with a red box. On the right, the Environment pane shows four objects: c (212 obs. of 4 variables), d (53 obs. of 4 variables), dengue (530 obs. of 4 variables), and e (53 obs. of 4 variables). A red box highlights object 'd', and a red arrow points from it to the 'number' column of the first row in the table (value 68).

	year	eweek	type_dengue	number
1	2018	1	Dengue	83
2	2018	2	Dengue	68
3	2018	3	Dengue	54
4	2018	4	Dengue	45
5	2018	5	Dengue	48
6	2018	6	Dengue	50

### 3. ANOTHER WAY TO FILTER OUT 2018 AND DENGUE TYPE

```
#5c. Another way to Filter out 2018 and 'Dengue' type  
e = dengue %>%  
  filter(year==2018) %>%  
  filter(type_dengue=='Dengue')
```

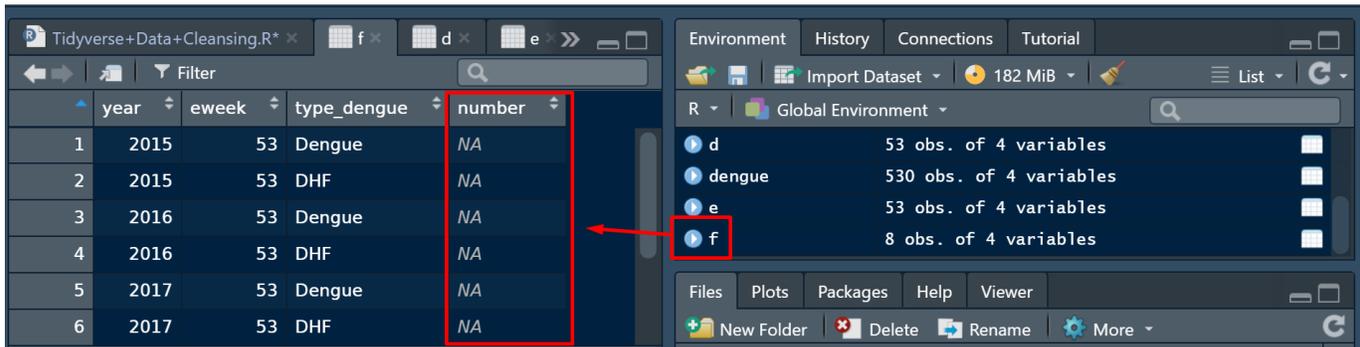
The screenshot shows the RStudio interface. On the left, a data table is displayed with columns: year, eweek, type\_dengue, and number. The table contains 6 rows of data for the year 2018, all of which are 'Dengue' type. A red box highlights the first three columns (year, eweek, type\_dengue) for all rows. On the right, the Environment pane shows the Global Environment with four objects: c (212 obs. of 4 variables), d (53 obs. of 4 variables), dengue (530 obs. of 4 variables), and e (53 obs. of 4 variables). A red box highlights the object 'e', and a red arrow points from this box to the third row of the data table.

	year	ewek	type_dengue	number
1	2018	1	Dengue	83
2	2018	2	Dengue	68
3	2018	3	Dengue	54
4	2018	4	Dengue	45
5	2018	5	Dengue	48
6	2018	6	Dengue	50

## F. HANDLING MISSING VALUES IN DENGUE.CSV DATASET

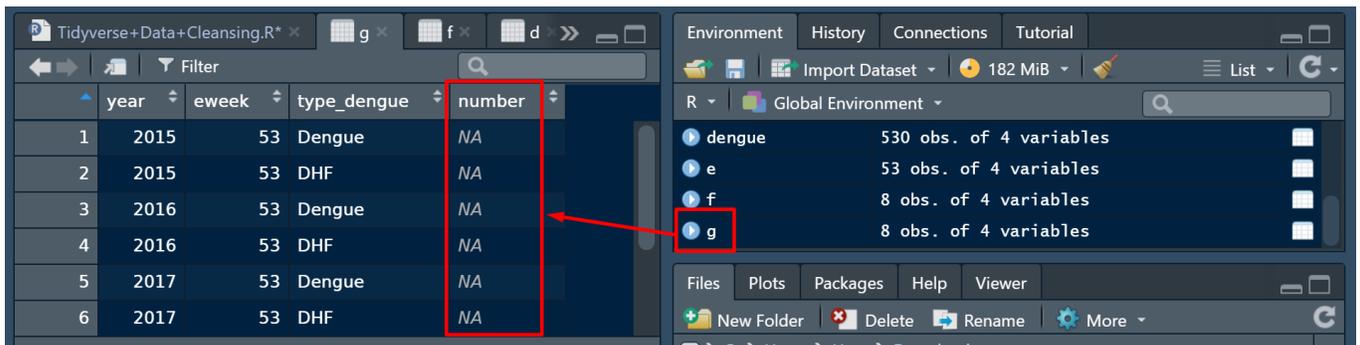
### 1. SHOW ALL NAs IN "NUMBER" COLUMN

```
#-----  
#6. Handling Missing values in Dengue.csv Dataset  
  
#6a. Show All NAs in "number" column  
f = dengue %>%  
  filter(is.na(number))
```



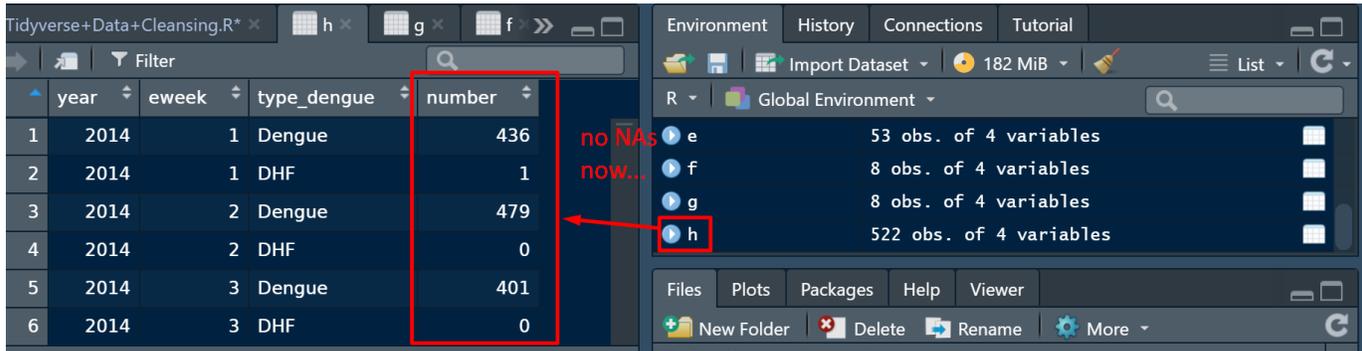
### 2. ANOTHER WAY OF SHOWING ALL NAs IN ALL COLUMNS

```
#6b. Another way of showing all NAs in all columns  
g = dengue %>%  
  filter(!complete.cases(.))
```



3. SHOWING ALL NO NAs (FILLED COLUMNS) NOW

```
#6c. Showing All NO NAs (filled columns) now...  
h = dengue %>%  
  filter(complete.cases(.))
```



## G. MUTATE DATA

```
#-----  
#7. Mutate data  
  
#using "eweek" to create a new column called "date"..  
i = dengue %>%  
  mutate(date = ymd(paste0(year, "-01-01"))+weeks(eweeek))
```

using eweeek to create a new column called "date" .....

so 1+7 = 8

eweeek2 = 14

so 1+ 14 = 15

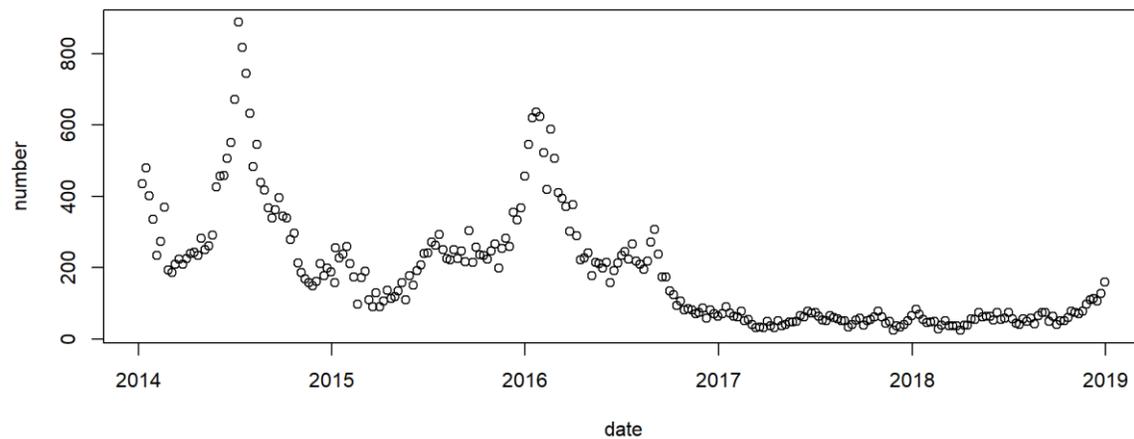
	year	eweeek	type_dengue	number	date
1	2014	1	Dengue	436	2014-01-08
2	2014	1	DHF	1	2014-01-08
3	2014	2	Dengue	479	2014-01-15
4	2014	2	DHF	0	2014-01-15
5	2014	3	Dengue	401	2014-01-22
6	2014	3	DHF	0	2014-01-22
7	2014	4	Dengue	336	2014-01-29

Showing 1 to 8 of 530 entries, 5 total columns

## H. FILTER, MUTATE THEN PLOT

### 1. DENGUE.CSV

```
#-----  
#8. Filter, Mutate then Plot  
  
#8a. Dengue.csv  
dengue %>%  
  filter(complete.cases(.)) %>%           #remove all NAs  
  
  filter(type_dengue=='Dengue') %>%      #filter out only "Dengue" type  
  
  mutate(date = ymd(paste0(year,"-01-01"))+weeks(eweeek)) %>%  
  #create a new column called "date"  
  
  select(date,number) %>%  
  #selecting out only "date" and "number" columns to plot  
  
  plot()
```



2. ANOTHER EXAMPLE FOR FILTER, MUTATE THEN PLOT  
(VACCINATION.XLS)

```
#-----  
#8b. Another Example for Filter, Mutate then Plot (vaccination.xls)  
#https://www.alvinang.sg/s/vaccination.xlsx  
  
vaccination <- read_excel("vaccination.xlsx")  
  
vaccination %>%  
  filter(complete.cases(.)) %>% #remove all NAs  
  
  filter(vaccination_type=='Poliomyelitis') %>%  
  #filter out only 'Poliomyelitis'  
  
  mutate(date = ymd(paste0(year, "-01-01"))) %>%  
  #create a new column called "date"  
  
  mutate(doses = no_of_doses_in_thousands) %>%  
  #rename the column  
  
  select(date, doses) %>%  
  #selecting out only "date" and "doses" columns to plot  
  
plot()
```

### 3. FILTER, MUTATE THEN EXPORT TO CSV

```
# Export to CSV
dengue_filtered <- dengue %>%
  filter(complete.cases(.)) %>%
  filter(type_dengue=='Dengue') %>%
  mutate(date = ymd(paste0(year, "-01-01"))+weeks(eweek)) %>%
  select(date,number)

write_csv(dengue_filtered, path = "dengue_filtered.csv")
```

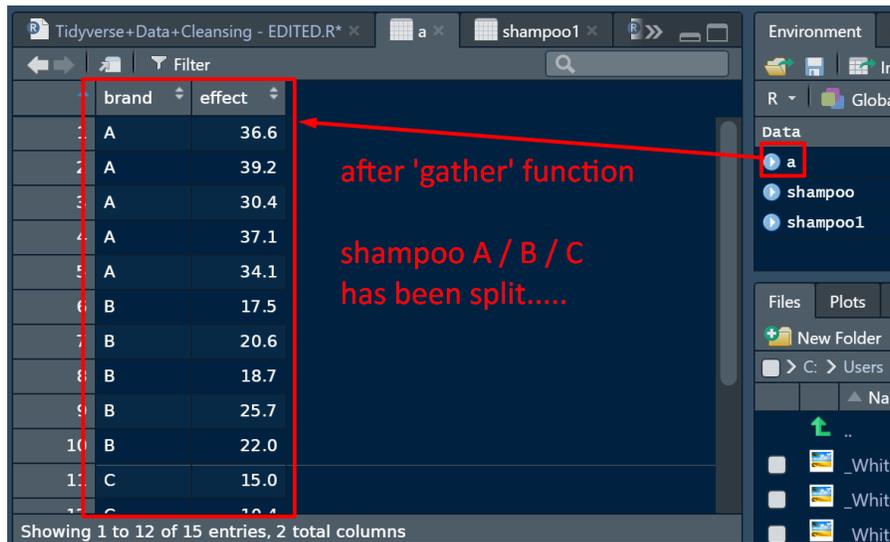
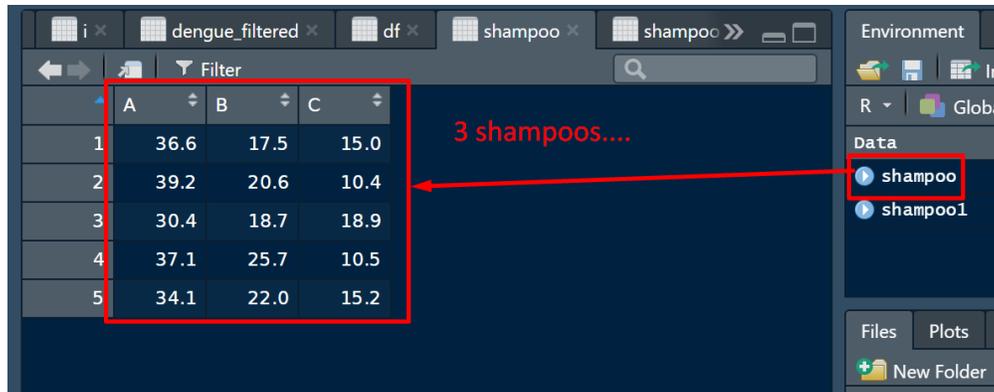
The screenshot shows the RStudio interface. On the left, the 'Environment' pane displays the 'dengue\_filtered' data frame with 261 observations and 2 variables. Below it, the 'Files' pane shows the 'dengue\_filtered.csv' file in the 'Downloads' directory. A red box highlights the 'dengue\_filtered' data frame in the Environment pane and the 'dengue\_filtered.csv' file in the Files pane. A red arrow points from the text 'this has now been created (write csv) into the working directory folder.....' to the 'dengue\_filtered.csv' file. The console shows a warning message: 'Warning message: The path argument of write\_csv() is deprecated as of readr 1.4.0.'

Environment	Size	Modified
dengue_filtered	261 obs. of 2 variables	
e	53 obs. of 4 variables	
f	8 obs. of 4 variables	
g	8 obs. of 4 variables	
h	522 obs. of 4 variables	

Name	Size	Modified
DAY_1_with_Dr_Alvin.ipynb	48.2 KB	Apr 18, 2022, 9:22 A
DB.Browser.for.SQLite-3.12.2-win32...	15 MB	Nov 26, 2021, 5:50 P
dengue_filtered.csv	3.7 KB	Apr 23, 2022, 6:27 P
dengue-clusters-kml (1).kml	884.2 KB	Mar 10, 2022, 3:00 P

#### 4. USING GATHER TO PIVOT DATA

```
shampoo =  
  data.frame('A'=c(36.6, 39.2, 30.4, 37.1, 34.1),  
            'B' = c(17.5, 20.6, 18.7, 25.7, 22.0),  
            'C'=c(15.0, 10.4, 18.9, 10.5, 15.2))  
  
shampoo1 <- as_tibble(shampoo)  
  
a = shampoo1 %>%  
  gather(brand, effect)
```



## I. JOINS

```
#-----  
#11. Data Joins  
  
df1 = data_frame(name=c('Ally', 'Steve', 'John'), age=c(45,46,47))  
df2 = data_frame(name=c('Ally', 'Belinda', 'John'), age=c(45,48,47))
```

	name	age
1	Ally	45
2	Steve	46
3	John	47

	name	age
1	Ally	45
2	Belinda	48
3	John	47

## 1. LEFT JOIN

```
#11a. Left Join  
left_join(df1, df2, by='name')
```

	name	age
1	Ally	45
2	Steve	46
3	John	47

df1

	name	age
1	Ally	45
2	Belinda	48
3	John	47

df2

	name	age.x	age.y
1	Ally	45	45
2	Steve	46	NA
3	John	47	47

LEFT JOIN

## 2. RIGHT JOIN

```
#11b. Right Join  
right_join(df1,df2,by='name')
```

	name	age
1	Ally	45
2	Steve	46
3	John	47

df1

	name	age
1	Ally	45
2	Belinda	48
3	John	47

df2

	name	age.x	age.y
1	Ally	45	45
2	John	47	47
3	Belinda	NA	48

RIGHT JOIN

### 3. INNER JOIN

```
#11c. Inner Join  
inner_join(df1,df2,by='name')
```



#### 4. FULL JOIN

```
#11d. Full Join  
full_join(df1,df2,by='name')
```

	name	age
1	Ally	45
2	Steve	46
3	John	47

df1

	name	age
1	Ally	45
2	Belinda	48
3	John	47

df2

	name	age.x	age.y
1	Ally	45	45
2	Steve	46	NA
3	John	47	47
4	Belinda	NA	48

FULL  
JOIN

## J. GROUPBY

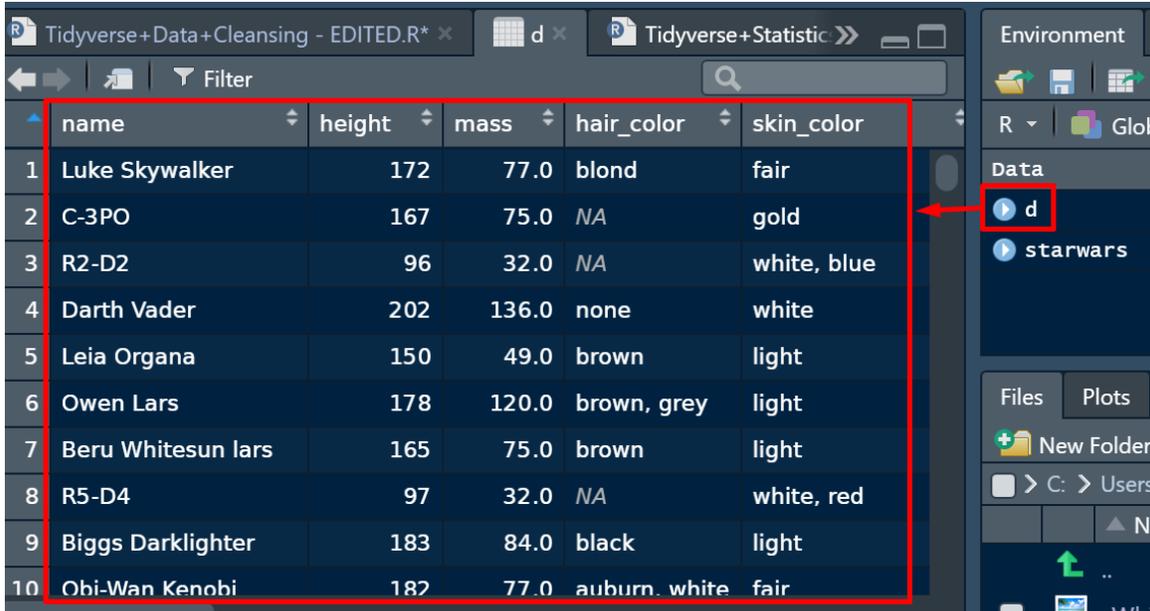
	extra	group	ID
6	3.4	1	6
7	3.7	1	7
8	0.8	1	8
9	0.0	1	9
10	2.0	1	10
11	1.9	2	1
12	0.8	2	2
13	1.1	2	3
14	0.1	2	4
15	-0.1	2	5
16	4.4	2	6

```
s1 = sleep %>%  
  group_by(group) %>%  
  summarize(avg_extra=mean(extra))
```

	group	avg_extra
1	1	0.75
2	2	2.33

## K. REMOVING COLUMN

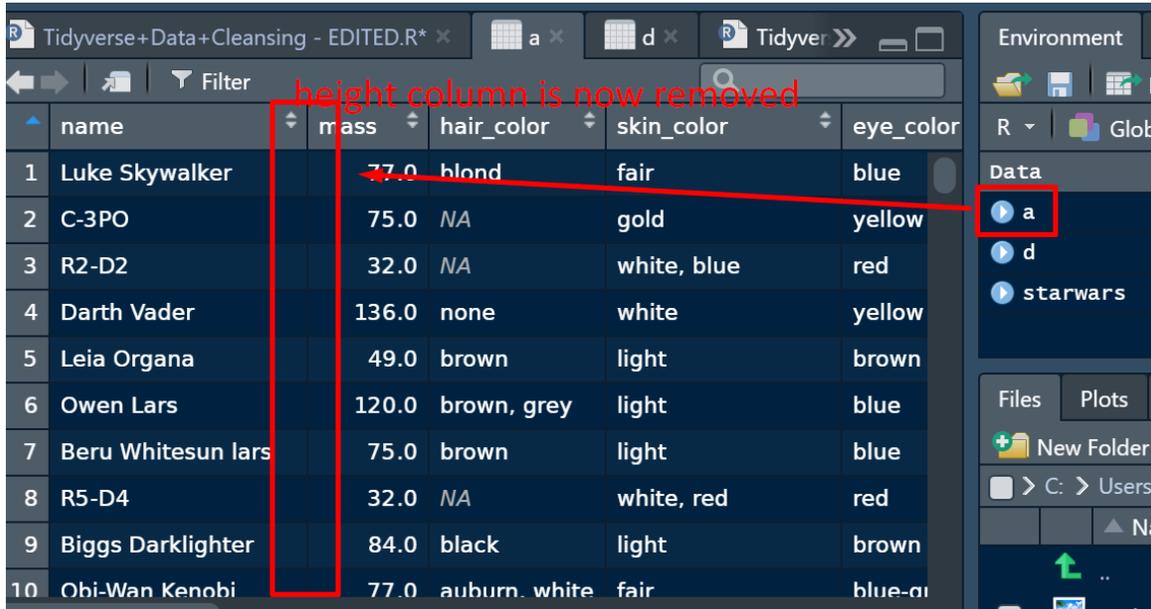
```
data("starwars", package = "dplyr")
d = starwars
```



	name	height	mass	hair_color	skin_color
1	Luke Skywalker	172	77.0	blond	fair
2	C-3PO	167	75.0	NA	gold
3	R2-D2	96	32.0	NA	white, blue
4	Darth Vader	202	136.0	none	white
5	Leia Organa	150	49.0	brown	light
6	Owen Lars	178	120.0	brown, grey	light
7	Beru Whitesun lars	165	75.0	brown	light
8	R5-D4	97	32.0	NA	white, red
9	Biggs Darklighter	183	84.0	black	light
10	Obi-Wan Kenobi	182	77.0	auburn, white	fair

## 1. REMOVING THE 'HEIGHT' COLUMN

```
#remove the 'height' column  
a = select(starwars, -height)
```



height column is now removed

	name	mass	hair_color	skin_color	eye_color
1	Luke Skywalker	77.0	blond	fair	blue
2	C-3PO	75.0	NA	gold	yellow
3	R2-D2	32.0	NA	white, blue	red
4	Darth Vader	136.0	none	white	yellow
5	Leia Organa	49.0	brown	light	brown
6	Owen Lars	120.0	brown, grey	light	blue
7	Beru Whitesun lars	75.0	brown	light	blue
8	R5-D4	32.0	NA	white, red	red
9	Biggs Darklighter	84.0	black	light	brown
10	Obi-Wan Kenobi	77.0	auburn, white	fair	blue-gr

2. RENAME THE 'NAME' COLUMN

```
#rename the 'name' column  
b = starwars %>%  
  rename(BLABLABLA = name)
```

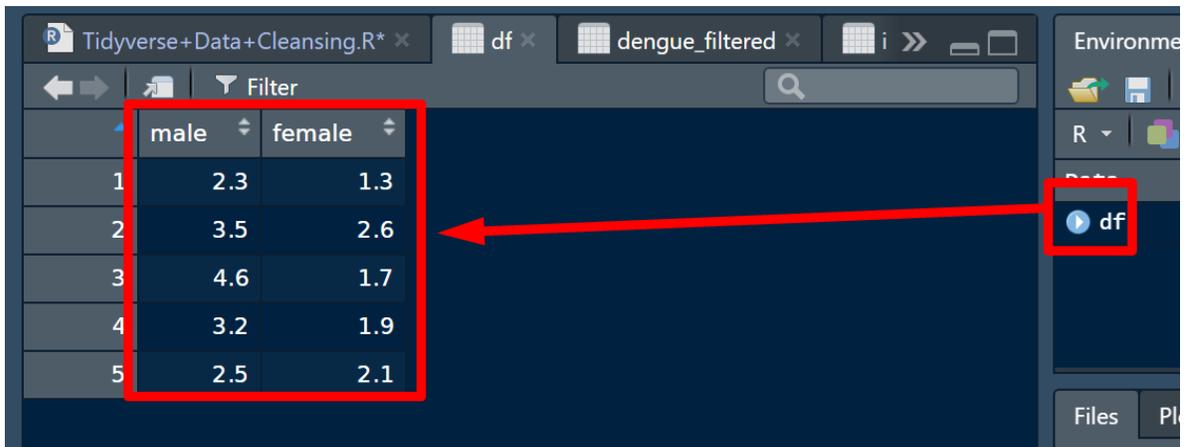
name column has been renamed

	BLABLABLA	height	mass	hair_color	skin_color
1	Luke Skywalker	172	77.0	blond	fair
2	C-3PO	167	75.0	NA	gold
3	R2-D2	96	32.0	NA	white, blue
4	Darth Vader	202	136.0	none	white
5	Leia Organa	150	49.0	brown	light skin - white
6	Owen Lars	178	120.0	brown, grey	light
7	Beru Whitesun lars	165	75.0	brown	light
8	R5-D4	97	32.0	NA	white, red
9	Biggs Darklighter	183	84.0	black	light
10	Obi-Wan Kenobi	182	77.0	auburn, white	fair

## L. DIFFERENCES BETWEEN TIBBLE VS DATAFRAME

### 1. TIBBLE

```
#-----  
#15. Differences between Tibble vs Dataframe  
#15a. Tibble  
  
df <- tibble(  
  'male' = c(2.3, 3.5, 4.6, 3.2, 2.5),  
  'female' = c(1.3, 2.6, 1.7, 1.9, 2.1)  
)  
df
```



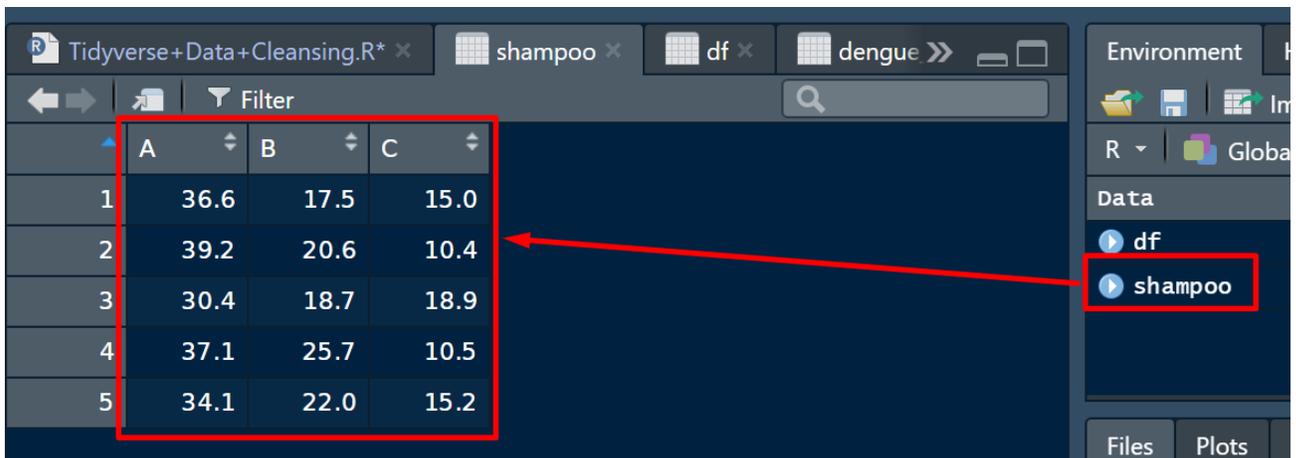
The screenshot shows the RStudio environment pane with a variable named 'df'. The variable is highlighted with a red box. A red arrow points from the 'df' variable to a preview of the data, which is also highlighted with a red box. The data is a tibble with two columns: 'male' and 'female'.

	male	female
1	2.3	1.3
2	3.5	2.6
3	4.6	1.7
4	3.2	1.9
5	2.5	2.1

## 2. DATAFRAME

### #15b. Dataframe

```
shampoo = data.frame(  
  'A' = c(36.6, 39.2, 30.4, 37.1, 34.1),  
  'B' = c(17.5, 20.6, 18.7, 25.7, 22.0),  
  'C' = c(15.0, 10.4, 18.9, 10.5, 15.2))
```



The screenshot shows the RStudio interface. The main window displays a dataframe with 5 rows and 3 columns (A, B, C). The data is as follows:

	A	B	C
1	36.6	17.5	15.0
2	39.2	20.6	10.4
3	30.4	18.7	18.9
4	37.1	25.7	10.5
5	34.1	22.0	15.2

The Environment pane on the right shows the 'shampoo' object under the 'Data' section, which is highlighted with a red box. A red arrow points from this box to the first row of the dataframe in the main window.

### 3. AS TIBBLE

```
#15c. As Tibble  
shampoo1 <- as_tibble(shampoo)
```

even after tibbling,  
you don't see any  
difference....

	A	B	C
1	36.6	17.5	15.0
2	39.2	20.6	10.4
3	30.4	18.7	18.9
4	37.1	25.7	10.5
5	34.1	22.0	15.2

#### 4. COMPARING CONVERSION

## Data Frame

```
df1 <- data.frame(  
  gender = c("Female", "Female", "Male"),  
  height = c(152, 171.5, 165),  
  weight = c(81, 93, 78),  
  age = c(42, 38, 26),  
  row.names = c('Ally', 'Belinda', 'Alvin')  
)
```

	gender	height	weight	age
Ally	Female	152.0	81	42
Belinda	Female	171.5	93	38
Alvin	Male	165.0	78	26

## Tibble

```
df2 <- tibble(  
  gender = c("Female", "Female", "Male"),  
  height = c(152, 171.5, 165),  
  weight = c(81, 93, 78),  
  age = c(42, 38, 26),  
  row.names = c('Ally', 'Belinda', 'Alvin')  
)
```

	gender	height	weight	age	row.names
1	Female	152.0	81	42	Ally
2	Female	171.5	93	38	Belinda
3	Male	165.0	78	26	Alvin

Not Much Difference....

- There's not much visible difference between a Data Frame vs Tibble....
- Except that Tibble adds an extra column....

## 5. COMPARING STRUCTURE (STR)

### Data Frame

```
> str(df1)
'data.frame':   3 obs. of  4 variables:
 $ gender: chr  "Female" "Female" "Male"
 $ height: num  152 172 165
 $ weight: num   81  93  78
 $ age   : num   42  38  26
```

### Tibble

```
> str(df2)
tibble [3 x 5] (S3: tbl_df/tbl/data.frame)
 $ gender : chr [1:3] "Female" "Female" "Male"
 $ height : num [1:3] 152 172 165
 $ weight : num [1:3] 81 93 78
 $ age    : num [1:3] 42 38 26
 $ row.names: chr [1:3] "Ally" "Belinda" "Alvin"
>
```

The Structure of a Data Frame vs Tibble also don't show much difference....

- Even if you look at the structure...they display the same things....

## Data Frame

```
> df1$ge  
[1] "Female" "Female" "Male"
```

Even though the proper column name is called "gender" .... If you use a DataFrame, You can misspell it as \$ge and it will still Show the column items....

This might cause future errors if you accidentally Call out the wrong column with similar column "ge" headings.....

## Tibble

```
> df2$ge  
NULL  
warning message:  
unknown or uninitialised column: `ge`  
> df2$gender  
[1] "Female" "Female" "Male"
```

However, for Tibble, you are not able to Display the column items if you misspell The column name.....it will show an error...

You have to type out the whole "\$gender" To get the items.....

This prevents future errors.....

- But you are not able to use short forms for the column names to call out the items...

## 7. COMPARING DISPLAY

```
1 library(tidyverse)
2 library(tibble)
3 library(tidyr)
4 library(dplyr)
5 library(readxl)
6 library(ggplot2)
7 library(lubridate)
8
9 penguins = read.csv('penguins.csv', header = TRUE)
10
11 as.data.frame(penguins)
12
13 as_tibble(penguins)
```

if we run this code line 11 we converted the penguins.csv into dataframe format

we see on the right that the data is messy and all over the place

```
> as.data.frame(penguins)
  rowid species island bill_length_mm bill_depth_mm flipper_length_m
1         1  Adelle Torgersen      39.1          18.7             181
2         2  Adelle Torgersen      39.5          17.4             186
3         3  Adelle Torgersen      40.3           18              195
4         4  Adelle Torgersen      NA            NA                NA
5         5  Adelle Torgersen      36.7          19.3             193
6         6  Adelle Torgersen      39.3          20.6             190
7         7  Adelle Torgersen      38.9          17.8             181
8         8  Adelle Torgersen      39.2          19.6             195
9         9  Adelle Torgersen      34.1          18.1             193
10        10  Adelle Torgersen       42           20.2             190
```

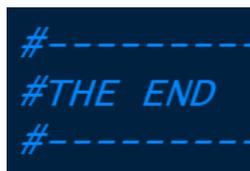
- The Penguins dataset can be found here: <https://www.alvinang.sg/s/penguins.csv>
- For dataframe as shown above, we see that it displays very messily in the console....

```
1 library(tidyverse)
2 library(tibble)
3 library(tidyr)
4 library(dplyr)
5 library(readxl)
6 library(ggplot2)
7 library(lubridate)
8
9 penguins = read.csv('penguins.csv', header = TRUE)
10
11 as.data.frame(penguins)
12
13 as_tibble(penguins)
```

now that we use tibble, we see that the data is displayed neatly on the right console....

```
110      4775  male 2009
111      3825  female 2009
[ Reached max. # of options ("max.print") -- omitted 233 rows ]
> as_tibble(penguins)
# A tibble: 344 x 7
  rowid species island bill_length_mm bill_depth_mm flipper_length_mm
  <int> <fct>   <fct>         <dbl>         <dbl>         <int>
1     1  Adelle Torgersen      39.1           18.7           181
2     2  Adelle Torgersen      39.5           17.4           186
3     3  Adelle Torgersen      40.3            18            195
4     4  Adelle Torgersen      NA              NA              NA
5     5  Adelle Torgersen      36.7           19.3           193
6     6  Adelle Torgersen      39.3           20.6           190
7     7  Adelle Torgersen      38.9           17.8           181
8     8  Adelle Torgersen      39.2           19.6           195
9     9  Adelle Torgersen      34.1           18.1           193
10    10  Adelle Torgersen       42            20.2           190
# ... with 334 more rows, and 3 more variables: body_mass_g <int>,
# sex <fct>, year <int>
```

- For tibble, we now see that its displayed neatly....in the console....



---

## ABOUT THE AUTHOR

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### ABOUT DR. ALVIN ANG

Dr. Alvin Ang earned his Ph.D., Masters and Bachelor degrees from NTU, Singapore. He was a previously a Professor, Scientist and Financial Consultant. Currently, he owns multiple self-started businesses and is a Personal/Business Advisor.

More about him at [www.AlvinAng.sg](http://www.AlvinAng.sg)