

# 0513367\_陳中堅

## Project Introduction

R\_2020

### ► Data Source

► Dengue case surveillance data: In Taiwan, dengue fever (台灣瘧疾) is classified as a notifiable infectious disease and suspected cases must be reported to a clinic for diagnosis within 24 hours.

► These data obtained from the web-based National Infectious Disease Statistics System (<https://nidss.cdc.gov.tw/en/SingleDisease.aspx?dc=1&dt=4&disease=061&position=1>) under the Notifiable Disease Surveillance System (NDSS) of the Taiwan Center for Disease Control. (from January 2012 to May 2019)

County	ID for each county	Year/Week (2012-2019-22 = 381)	Number of people with Dengue Fever (confirmed cases)	Breiman Index	Average Temperature (°C)	Maximum Temperature (°C)	Minimum Temperature (°C)	Relative Humidity (%)	Precipitation
County ID		Time	Dengue Cases	BI	Tave	Tmax	Tmin	Humid	Prep
1	Hualien	1	0	0.3	19.61	18.62/14.28	14.1/4.82/1	75.17	
4	Hualien	1	0	0	19.34	21.22/14.28	14.8	75.43	
5	Hualien	1	0	0	13.3	21.9/14.28/1	1/1	76.43	
6	Hualien	1	0	0	15.6	12.1/14.28/1	1/1	79.17	
7	Hualien	1	0	0	17.96	20.4/14.28/1	16.05/14.28	76.57	
8	Hualien	1	0	0	18.21	21.37/14.28	15.85/14.28	74.57	
9	Hualien	1	0	0	16.8	22.57/14.28	17.51/14.28	75.24	
10	Hualien	1	0	0.06	19.61	22.40/14.28	17.57/14.28	81.7	
17	Hualien	1	0	0	19.73	23.65/14.28	17.37/14.28	82.43	
17	Hualien	1	0	0	21.5	24.40/14.28	16.77/14.28	80	
17	Hualien	1	0	0.07	20.17	23.66/14.28	17.14/14.28	80.24	
14	Hualien	1	0	0	21.34	24.5	16.67/14.28	76.06	
15	Hualien	1	0	0	19.61	23.71/14.28	16.5	63.14	
15	Hualien	1	0	0	20.41	24.2	17.52/14.28	69.43	
17	Hualien	1	0	0.13	23.26	26.34/14.28	16.25/14.28	79.43	
18	Hualien	1	0	0.18	23.24	26.6	16.86/14.28	79.43	
19	Hualien	1	0	0.3	25.43	27.4	16.32/14.28	78.74	
20	Hualien	1	0	0	24.67	27.64/14.28	12.18/14.28	79.43	
21	Hualien	1	0	0.17	24.3	26.6	12.04/14.28	75.43	
22	Hualien	1	0	0	20.93	30	22.1	74.74	
23	Hualien	1	0	0	25.27	28.62/14.28	22.64/14.28	75.24	

	A	B	C	D	E	F	G	H	I
1	ID	Dengue	BI	Tave	Humid	Prep			
2	Mandara	30	0.467804	28.7251	77.32132	5.481886			
3	Pangloss	1178	0.136165	28.82172	74.07245	5.818788			
4	1305/ANAN	312	0.500015	23.01558	77.20095	5.58322			
5	Lubad	558	0.176382	23.54752	72.74812	6.732502			
6	Chisley City	42	0.182232	23.0346	78.12711	5.101800			
7	Chisley Cou	59	0.162222	23.93551	78.12711	5.391809			
8	New Taipei	523	0.302842	23.77915	73.04752	6.191266			
9	Moodi	66	0.276786	22.8688	79.44726	4.261526			
10	NANOU	23	0.21282	18.11899	81.93103	0.930775			
11	Poukua	43	0.144553	23.79556	79.17615	3.381409			
12	Haincha Cit	64	0.780129	21.00202	76.8951	4.981137			
13	Haincha Cou	90	0.700129	23.09202	76.8951	3.193137			
14	Yilan	31	0.403178	22.98038	80.58902	7.731517			
15	Taipei	486	0.490147	23.00424	74.09046	4.814522			
16	Linan	23859	0.380281	21.71191	71.8181	5.131137			
17	Kuoshinhou	25972	0.32447	25.77742	73.33075	5.742463			
18	Chingmau	141	0.231912	23.06346	81.0868	4.522097			
19	Yilan	68	0.171532	21.25103	82.19857	5.977993			
20	Takima	44	0.350465	24.70432	75.77424	5.056077			
21	Kechang	20	0.58023	22.8871	76.7318	0.928			

# Introduction

- According to the computerized database of the surveillance system by Taiwan Center for Disease Control (Taiwan-CDC) [29], in the period of 2006 to 2012, there were 10,094 confirmed cases of dengue virus infections, or an average of about 1442 cases per year. However, from 2012 to 2018, the Taiwan-CDC recorded 63,471 confirmed cases of dengue, or an average of about 9067 cases per year. Thus, the annual average number of dengue cases has increased by 529% in the past seven years compared with the period from 2006 to 2012.

	Bureau Index	Average Temperature (°C)	Maximum Temperature (°C)	Minimum Temperature (°C)	Relative Humidity (%)	Precipitation (mm)	Population (persons)	Population Density (per km <sup>2</sup> )
1								
2	BT	Tave	Tmax	Tmin	Humid	Precp	Pop	Pop_Den
3	0.5	16.61	18.65/14286	14.7/14285/1	75.71	0.57	336704.9965	72.7/498/11
4	0	18.94	21.55714286	16.0	75.43	0.36	336676.9953	72.71990202
5	0	19.5	21.91428571	17.1	76.45	2.4	336648.9965	72.72498711
6	0	19.6	22.77142857	15.7/14285/	79.71	5	3366621	72.7
7	0	17.96	20.11428571	16.05714286	76.57	1	336586.9948	72.7
8	0	18.24	21.37142857	15.985/1429	74.57	1.2	336752.9951	72.7
9	0	19.0	22.57142857	17.51428571	75.29	0.4	336518.9948	72.7
10	0.06	19.64	22.24285714	17.17142857	81.71	9.58	336485	72.7
11	0	19.58	22.65714286	17.27142857	84.45	7.21	336448.2422	72.7
12	0	21.5	24.11428571	18.77142857	80	2.58	336401.1896	72.7
13	0.07	20.17	23.61428571	17.14285/14	80.29	5.57	336359.7422	72.7
14	0	21.34	24.5	18.67142857	76.86	19.3	336318	72.7
15	0	19.64	23.71428571	16.5	63.14	0.07	336276.263	72.7
16	0	20.41	24.2	17.5285/143	69.43	1.5	336257.4422	72.6/498/109
17	0.13	23.26	26.31428571	20.25714286	79.43	1.17	336196.9782	72.61990279
18	0.18	23.24	26.6	20.685/1429	79.43	7.25	336156.4837	72.6249871
19	0.5	23.19	27.1	20.5285/143	78.11	5.33	336076	72.6
20	0	24.67	27.64285714	22.18571429	78.41	5.16	336061.9991	72.6
21	0.17	24.9	28.0	22.04285/14	77.43	2.75	336047.9988	72.6
22	0	26.03	30	22.7	74.14	4	336033.9991	72.6
23	0	26.27	28.8285/143	22.64285/14	75.29	8.64	336020	Activate Win72.6

- Analysis of this data is primarily intended to determine the geographic distribution of dengue cases across Taiwan's provinces.
- Simultaneously with the visualization of the density distribution of mosquito larvae, providing convincing proof of the leading cause of dengue fever, plus the influence of population density has led to the difference in rate as well as the number of cases between Taiwan's provinces.

# Code

```
#code at kaggle
#https://www.kaggle.com/lefeulien/notebook0f66b16139

library(tidyverse) #
metapackage of all tidyverse
packages

list.files(path = "../input")

library(caret)

install.packages("geojsonio")

library("geojsonio")

require("dplyr")

require("stringr")

require("data.table")

require("ggplot2")

require("maptools")

require("knitr")

require("kableExtra")

require("mapproj")

require("sf")

require("RColorBrewer")

path =
"../input/data-
deng-
bi/plotTW.csv"
df <-
read.csv(path,
h=T)
df <-
data.frame(df)
distill_data <- df
kable(distill_data)
summary(distill_data)

taiwan.map <-
st_read("../input
/tw-map-
shp/gadm36_T
WN_2.shp")
print(taiwan.map, n = 22)

plot(taiwan.map[1])
#plot(taiwan.map[2:13],
max.plot = 12)
plot(st_geometry(taiwan.map))
#plot(taiwan.map["NAME_2"],
axes = TRUE)
st_geometry(taiwan.map)

ggplot(data =
taiwan.map) +
geom_sf(aes(fill =
NAME_2)) +
scale_fill_manual(name =
"縣市區",
values =
colorRampPalette(brewer.pal(8, "Accent"))(22)) +
labs(title = "Taiwan
map")

print(distill_data$ID)

distill_data$ID <-
as.character(distill_data$
ID)
head(distill_data)

tmp <-
(min(distill_data$Dengue)
-
max(distill_data$Dengue
))
tmp

distill_data$Dengue_Nor
malize <-
(distill_data$Dengue -
min(distill_data$Dengue))
/
(max(distill_data$Dengue)
-

my.taiwan.map <-
taiwan.map[c("NAME
_2", "geometry")]
my.taiwan.map$NAME
E_2 <-
as.character(my.taiwan.map$NAME_2)
head(my.taiwan.map)

my.taiwan.map.data
<-
left_join(my.taiwan.map, distill_data,
c("NAME_2" = "ID"))

dim(my.taiwan.map.data)

ggplot(data =
my.taiwan.map.data) +
geom_sf(aes(fill =
Dengue_Normalize)) +
#
scale_fill_distiller(name
="Count", palette =
"RdYlGn")
scale_fill_distiller(palette =
"Spectral", name =
"Dengue Cases")

my.taiwan.map.data$Dengue3 <-
cut(my.taiwan.map.data$Dengue, breaks = c(-Inf, 61,
501, 1001, Inf), right = FALSE) # divide into 4 section for
clearly revealization

ggplot(data = my.taiwan.map.data) +
geom_sf(aes(fill = Dengue3)) +
#scale_fill_distiller(name="Count", palette = "RdYlGn")
#scale_fill_distiller(palette = "Spectral", name = "台灣瘧疾
Dengue Cases") +

scale_fill_manual(breaks=c("[-Inf,61)", "[61,501)",
"[501,1e+03)", "[1e+03, Inf)"),
name = "Dengue Cases",
values = c("lightyellow", "yellow", "orange",
"red"),
labels = c("0-60", "61-500", "501-1000", ">1001"))

my.taiwan.map.data$BI3 <-
cut(my.taiwan.map.data$BI, breaks = c(-Inf, 1, 2, 3,
Inf), right = FALSE) # divide into 4 section for clearly
revealization

#BI is Mosquito larva density parameter
ggplot(data = my.taiwan.map.data) +
geom_sf(aes(fill = BI)) +
# scale_fill_distiller(name="Count", palette = "RdYlGn")
scale_fill_distiller(palette = "Spectral", name = "Mosquito
larva density distribution")

ggplot(data = my.taiwan.map.data) +
geom_sf(aes(fill = BI3)) +
scale_fill_manual(breaks=c("[-Inf,1)", "[1,2)", "[2,3)", "[3,
Inf)"),
name = "Mosquito larva density distribution",
values = c("lightyellow", "yellow", "orange",
"red"),
labels = c("0-1", "1-2", "2-3", "3-4"))
```



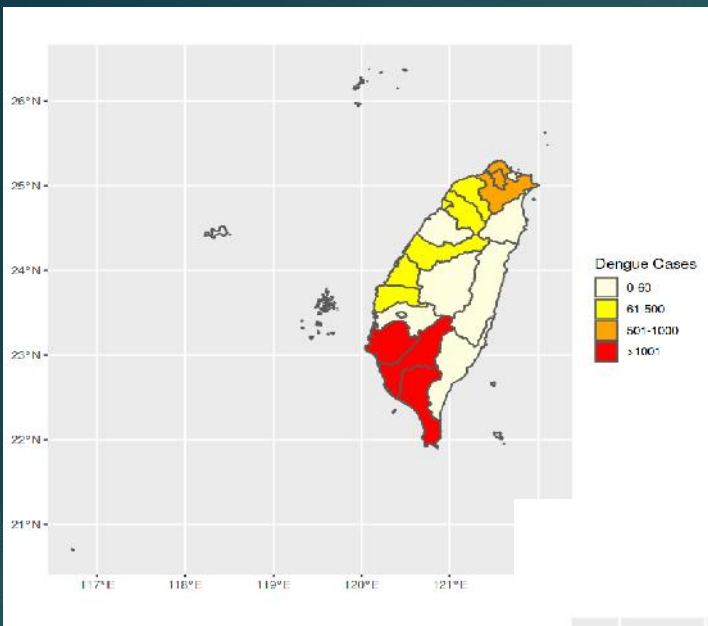
# Project Questions

Where is the area with the most mosquitoes in Taiwan?

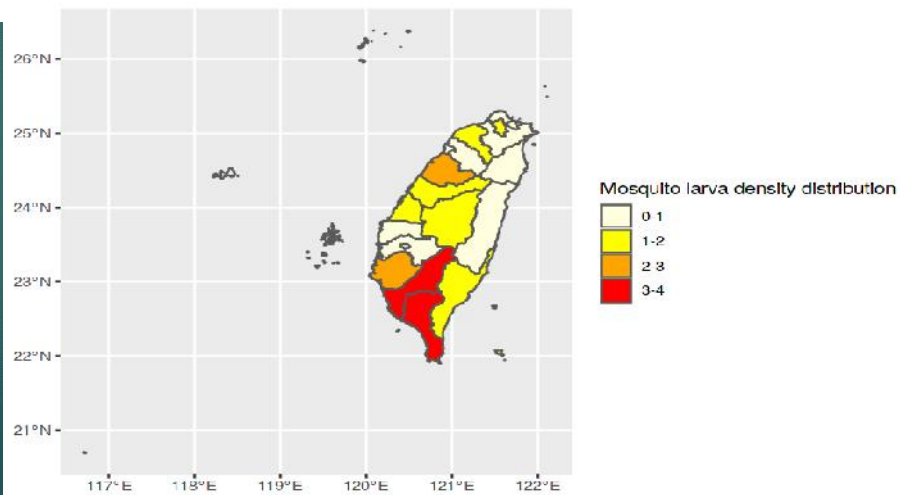
Which is the region with the most dengue cases?

Analysis the relationship between number of Dengue cases with mosquito density and provincial population density

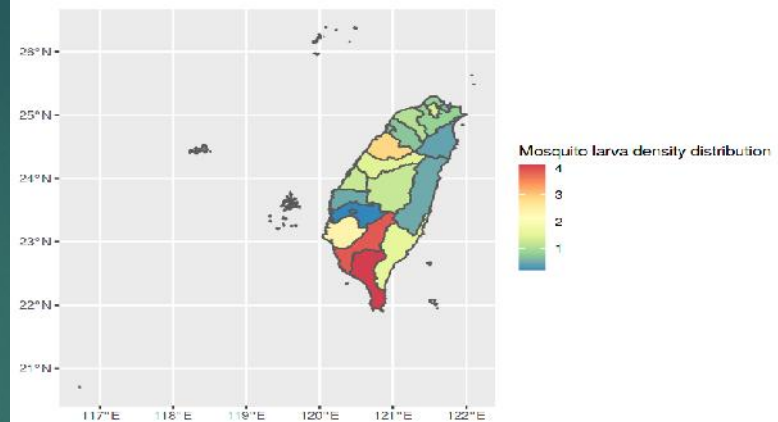





台灣瘧疾分佈  
Jan 2012 to May  
2019



台灣孑孓  
密度分佈



- 
- ▶ Result analysis
  - ▶ From the visualization graphs we can see the significant relationship between about Mosquito larva density and the number cases of Dengue, It also reveal an important relationship between cases of Dengue and population density with temperature of each Taiwan's province.
  - ▶ We can see that BI, mosquito larval distribution density is an important parameter affecting the number of dengue cases in the regions. However, there are some exceptions. For example, the Miaoli area has a high density of mosquito larvae but the incidence is low, most likely because this is a sparsely populated area, there are many mosquitoes in the wilderness, so there is no Affect the dengue fever.
  - ▶ Similarly, the Xinbei area with low mosquito density but high prevalence, it can only be explained because it's high population density.