0513367_陳中堅 Project Introduction

1	County	ID for each county	Year-Week (2012- 01 = 1 2019-22 = 35/) Time	Number of people with Dengoe Fever (confirmed exces)	Bretten Index	Average Temperature (°C)	Maximum Temperature (90)	Minimum Temperature (°C)	Relative Humelity (%)	Prospiration
2	Hualten	County ID	1me	Daugne Cases	0.5		1865/14286	14./1428571	Hunid	Ртеср
		-		<u> </u>	0.5	1661 1894			/5./° /5.43	
	Hushen	35	-	ů.				168		
	Hualten		,	Ů,	9	12.5		13.774,857	/6.43	
	Hualien	- 6	-	0	,	470	The second second second		79.71	
	Hualien	1	3	U.	0	1796	25-77-78-50-0	16,05714286	76.57	
	Hualien	1	5	0	0	1824		15.58571429	74.57	
	Hualien		1	"		19.8		17.51428571	75.29	
	Hualien	- 1	8	п	0.06	1964		17.57 42857	8.7	
	Hualien	1	9	0	0	19.59		17.37 42857	8243	
	Hualien	1	10		0	21.5		18.77142857	80	
	Hualien	1	11		0.07	20.17		17.14285714	60.29	
	Hualien	1	12	0	0	21.34			76.86	
15	Huslien	1	13	0		1961		165		
16	Huslien	1	14	0	0	2041	24.2	17.52857143	69.40	
1/	Hustien	1	15		0.13	23.25	2634285714	20,25714285	79.43	
18	Hushen	1	15	U	0.18	23.24	25.5	20,665/1429	79.43	
19	Haalien	1	1/	U	0.5	2349	27.4	20.5285/146	/8.14	
20	Haahen	1	18	U	0	246	27,64285714	22.185/1429	/9.43	
21	Hadren	1	19	U	0.17	24.3	28.6	22.04285714	75.43	
22	Hualten	1	20	U	0	26.05	30	221	74.14	
23	Hualien	1	21	0	0	2527	28.82957 143	22.642857 M	ivate Windows29	

-	A	Ti	C:	тэ	TC	T.	CI	TI	1	
1	ш.	Dengue	DI	Tave	liumd	Precp				
2	Husiica	30	0.467804	23.7231	77.32132	5.481886				
3	Pinghang	1178	4 136305	24 82372	74 07245	5 818786				
1	Taoyuan	312		23.04558		5.558822				
5	Taipei	558		23.54752						
6	Chinyi City	42	0 162222	23 9355	78 12711	5 30 1800				
7	Chiayi Cou	59	0.162222	23.9355	78.12711	5.391809				
8	New Taiper									
2	Mizoli	55			79 44726	4 263592				
10	Nantou	53	1.21292	19.41899	81.93463	6.980775				
	Perapha.	43		23.79556		3.381499				
	Hainchn Cit					4 981137				
1.3	Hisinchu Co			23.09202						_
14	Yilson	31		22.88928						
15	Taichung	486								
10	Talnan	23859		24,71191						_
17	Kachsiena	35972		25.77742		5.743463				
18	Changhua	141	1 231912	23 06346						
19	Yulin	68		21.35403		5.977933				_
20	Tratura	44		24.76432		5.054677				
21	Kaching	20	0.58023	22 8671	76 738	2.258				

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.as px?dc=1&dt=4&disease=061&position=1)und er the Notifiable Disease Surveillance System (NDSS) of the Taiwan Center for Disease Control.(from January 2012 to May 2019)

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.

4	Breteau Index	Амаяда Таправания (°G)	Maximum Temperature (°G)	Minimum Temperature (*G)	Relative Humidity (%)	Precpitation (mm)	Population (persons)	Population Density (per km2)
2	TRT	Tave	Tmax	Tmin	TInmid	Precp	Pop	Pop_Den
7-0	().:5	16.61	18.65714286	14./14285/1	/5./1	0.57	336704.9965	/2.//498/11
1	0	18.94	21.55714286	16.8	75.43	0.36	336676.9953	72.71998282
.5	0	19.3	21.91428571	17.1	76.43	2.4	336648.9965	72.72498711
6	U	U		13.77142857	79.71	د د	336621	/2./
7	0	17.96	20.41428571	16.05714286	76.57	1	336586.9948	72.7
5-3	O	18.24		15.58571429	14.51	1.2	886552,9981	12.1
9	0	19.8		17.51428571	75.29	0.4	336518.9948	
10	0.06	19.64	22.24285714	17.57142857	81.71	9.58	336485	72.7
11	U	19.58	22.65/14286	17.37142857	82.43	7.21	336443.2422	
12	0	21.5	24.44285714	18.77142857	80	2.58	336401.4896	72.7
133	0.07	20.17	23.61428571	17.14285714	80.29	8.51	886859.7422	12.1
14	U	21.34		18.67142857	/6.86	4.93	336378	12.1
1.5	O	19.61	23.71428571	16.5	63.14	0.07	336276.263	72.7
16>	O	20.41	24.2	17.52857148	69.48	1.5	886257.4887	72.67498709
17	0.13	23.26	26.34285714	20.25714286	79.43	1.17	336196.9782	72.64998279
18	0.18	23.24	26.6	20.68571429	79.43	1.2%	336136.4837	72.6249871
19	0.5	23.49		20.52857143	78.14	5.33	336076	
20	n	24.67	27.64285714	22.18571429	79.43	5.36	336061.9991	72.6
21	0.17	24.9	28.6	22.04285714	75.43	2.75	886047,9988	
22	0	26.03	30	22.7	74.14	1	336033.9991	72.6
22.73	O	2:5.21	28.82857143	22.64285714	1:5.29	8.64	336020	Activate Window

- 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/lefe ulien/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("RColorBrewer")

require("sf")

shp/gadm36_T WN 2.shp") print(taiwan.m ap, n = 22plot(taiwan.ma p[1]) #plot(taiwan.m ap[2:13], max.plot = 12plot(st_geometr y(taiwan.map)) #plot(taiwan.m ap["NAME_2"], axes = TRUE) st_geometry(tai wan.map)

```
path =
                       ggplot(data =
"../input/data-
deng-
                       taiwan.map) +
bi/plotTW.csv"
                        geom_sf(aes(fill =
df <-
                       NAME_2)) +
read.csv(path,
                        scale fill manual(name
h=T)
                       = "縣市區".
df <-
                                  values =
data.frame(df)
                       colorRampPalette(brewe
distill data <- df
                       r.pal(8, "Accent"))(22)) +
kable(distill dat
                       labs(title = "Taiwan
                       map")
summary(distill_
data)
                       print(distill data$ID)
taiwan.map <-
                       distill data$ID <-
st read("../input
                       as.character(distill_data$
/tw-map-
                       head(distill_data)
                       tmp <-
                       (min(distill_data$Dengue)
                       max(distill_data$Dengue
                       tmp
                       distill data$Dengue Nor
                       nalize <-
                       (distill data$Denaue -
                       min(distill_data$Dengue))
                       (max(distill_data$Dengu
```

```
taiwan.map[c("NAME
_2", "geometry")]
my.taiwan.map$NAM
E 2 <-
as.character(my.taiw
an.map$NAME 2)
head(my.taiwan.map)
my.taiwan.map.data
left_join(my.taiwan.m
ap, distill data,
c("NAME_2" = "ID"))
dim(mv.taiwan.map.d
ata)
aaplot(data =
my.taiwan.map.data)
 geom_sf(aes(fill =
Dengue Nornalize)) +
scale fill distiller(name
="Count", palette =
"RdYIGn")
scale_fill_distiller(palett
e = "Spectral", name =
"Dengue Cases")
```

my.taiwan.map <-

```
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 = "RdYIGn")
 #scale fill distiller(palette = "Spectral", name = "台灣瘧疾
Denguge 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(mv.taiwan.map.data\$Bl.breaks = c(-Inf. 1, 2, 3, ...)
Inf),right = FALSE)# divide into 4 section for clearly
revealization
#BI is Mosquito larva density parameter
aaplot(data = mv.taiwan.map.data) +
geom_sf(aes(fill = BI)) +
# scale_fill_distiller(name="Count", palette = "RdYIGn")
scale fill distiller(palette = "Spectral", name = "Mosquito
larva density distribution")
ggplot(data = my.taiwan.map.data) +
 aeom 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

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.

