

Stawberries

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Strawberries: Data

This is a project about acquiring strawberry data from the USDA-NASS system and then cleaning, organizing, and exploring the data in preparation for data analysis. To get started, I acquired the data from the USDA NASS system and downloaded them in a csv.

Data cleaning and organization references

[“An introduction to data cleaning with R”](#) by Edwin de Jonge and Mark van der Loo

[“Problems, Methods, and Challenges in Comprehensive Data Cleansing”](#) by Heiko Müller and Johann-Christoph Freytag

Questions about Strawberries

- Where they are grown? By whom?
- Are they really loaded with carcinogenic poisons?
- Are they really good for your health? Bad for your health?
- Are organic strawberries carriers of deadly diseases?
- When I go to the market should I buy conventional or organic strawberries?
- Do Strawberry farmers make money?
- How do the strawberries I buy get to my market?

Strawberry data source and parameters

The data set for this assignment has been selected from:

[\[USDA_NASS_strawb_2024SEP25\]](#).

The data have been stored on NASS here: [USDA_NASS_strawb_2024SEP25](#) .

For the assignment, I stored the csv I downloaded on the MA615 Blackboard as strawberries25_v3.csv.

The data was originally collected at the county, state, and national levels, but the degree of missingness at the state level was too high, so I dropped the county-level data.

There are 5,359 rows and 21 column in the initial data set. The only complete year is 2022, although there is data for years 2018 through 2024.

To work with the data, define a function to remove columns with only single value in all its rows.

To work with this data, split the Census data from the Survey data.

Census data cleaning and organizing

we're examining census data because it's different from survey data

we're isolating organic data

Note that straw_cen has only one year: 2022

Current stats Census date has been isolated and split between Organic and Conventional strawberries

imputed values

We'll start with missing value interpolation for census data:

We will first interpolate the Bearing "Grown" values. Then we can interpolate Bearing and Non-bearing from those values as we discussed in lecture.

Now if GROWN is NA, there are three possibilities:

- 1 Bearing and Non-bearing are both NA
- 2 Neither Bearing nor Non-bearing are NA
- 3 One of Bearing nor Non-bearing are NA

The solution for each of these cases is different:

1 Interpolate from the total for area grown(a check of the dataframe shows you that the total is never NA). We'll try and calculate the average ratio of the missing parts across the non-missing data and split the difference from the total across those

2 Sum Bearing and non-bearing. Easy.

3 Calculate the average proportion of grown that bearing or non bearing(as the case may be) is for the state in question. Then we use that to figure out the value.

Applying our functions now to strawberry census:

Now that we have imputed the values, we still have an adjustment to make. Because we are using proportional averages, the values might no longer add up to our totals. So we need to write a function that will scale our imputed values so that the totals add up.

The happy news is because we are doing this by index, we still know which are the values we have imputed and therefore which values we need to scale.

Imputing acres bearing and non-bearing

We can go one better with imputation where we impute case 1 for the “GROWN” categories together with other case 1s for each. A slight modification to the function above plus some other changes achieves this.

CENSUS TABLES

Survey data cleaning and organizing

Shift data into alignment function

Examine Domain

now look at totals

there are two markets for Strawberries – Fresh Marketing and Processing

make a table for each

from the Survey Totals

we have reports for

Markets: Fresh and Processing Operations: Growing and Production

Florida - California - 2018 -2023

California and Florida chemicals

California and Florida fertilizers

```
library(ggplot2)
library(dplyr)
library(tidyverse)
library(PubChemR)
chemical<- read.csv("new.csv", header = TRUE)
```

#Question 1: Does different chemicals has different effects to the strawberry growth?

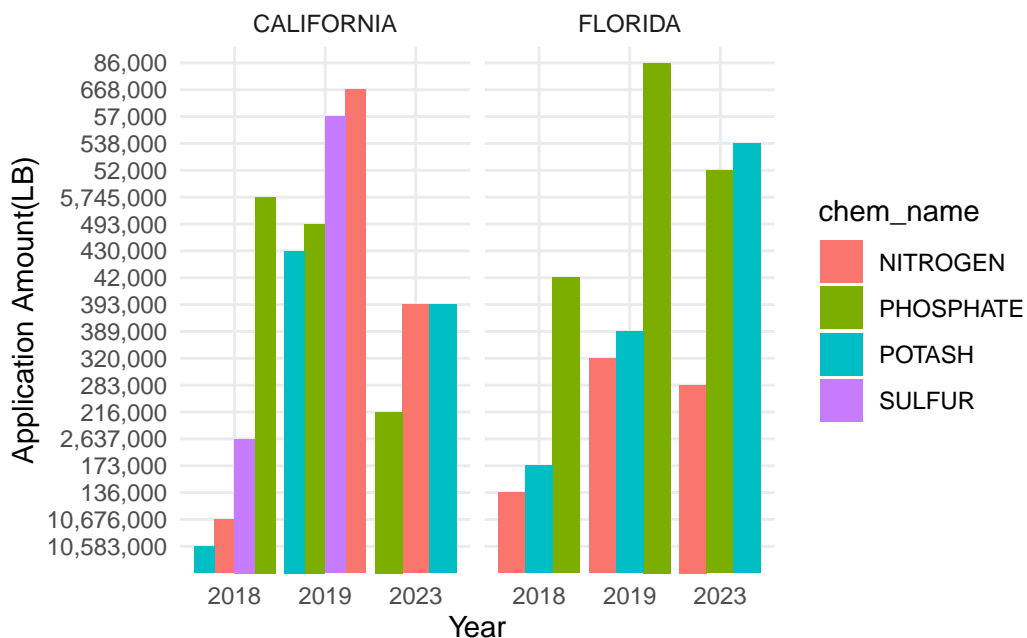
In the first step, I plan to use cleaned data named “new” from USDA-NASS data cleaning-ver7.qmd file. Since only Florida and California used chemicals, it is easy to start with total chemical used.

```
#Total Chemical
data_chem<- chemical %>%
  filter(Year %in% c("2018", "2019", "2023") & State %in% c("CALIFORNIA", "FLORIDA")& chem_n
filter(Value != "(D)") %>% filter(Value != "(NA)")
data_chem
```

	Year	State	mk2	measure	other	chem_name	Value
1	2023	CALIFORNIA	APPLICATIONS	LB	<NA>	NITROGEN	393,000
2	2023	CALIFORNIA	APPLICATIONS	LB	<NA>	PHOSPHATE	216,000
3	2023	CALIFORNIA	APPLICATIONS	LB	<NA>	POTASH	393,000
4	2023	FLORIDA	APPLICATIONS	LB	<NA>	NITROGEN	283,000
5	2023	FLORIDA	APPLICATIONS	LB	<NA>	PHOSPHATE	52,000
6	2023	FLORIDA	APPLICATIONS	LB	<NA>	POTASH	538,000
7	2019	CALIFORNIA	APPLICATIONS	LB	<NA>	NITROGEN	668,000
8	2019	CALIFORNIA	APPLICATIONS	LB	<NA>	PHOSPHATE	493,000
9	2019	CALIFORNIA	APPLICATIONS	LB	<NA>	POTASH	430,000
10	2019	CALIFORNIA	APPLICATIONS	LB	<NA>	SULFUR	57,000
11	2019	FLORIDA	APPLICATIONS	LB	<NA>	NITROGEN	320,000
12	2019	FLORIDA	APPLICATIONS	LB	<NA>	PHOSPHATE	86,000
13	2019	FLORIDA	APPLICATIONS	LB	<NA>	POTASH	389,000
14	2018	CALIFORNIA	APPLICATIONS	LB	<NA>	NITROGEN	10,676,000
15	2018	CALIFORNIA	APPLICATIONS	LB	<NA>	PHOSPHATE	5,745,000

16	2018	CALIFORNIA APPLICATIONS	LB	<NA>	POTASH	10,583,000
17	2018	CALIFORNIA APPLICATIONS	LB	<NA>	SULFUR	2,637,000
18	2018	FLORIDA APPLICATIONS	LB	<NA>	NITROGEN	136,000
19	2018	FLORIDA APPLICATIONS	LB	<NA>	PHOSPHATE	42,000
20	2018	FLORIDA APPLICATIONS	LB	<NA>	POTASH	173,000

```
# Check the range of values in Value
ggplot(data_chem, aes(x = factor(Year), y = Value, fill = chem_name))+
  geom_bar(stat = "identity", position = "dodge")+
  facet_wrap(~ State)+
  labs(
    x = "Year",
    y = "Application Amount(LB)")+
  theme_minimal()
```



From the histogram, we can find that In both states, Sulfur is consistently the least used chemical across all years, while Potash and Phosphate dominate in terms of usage in various years.

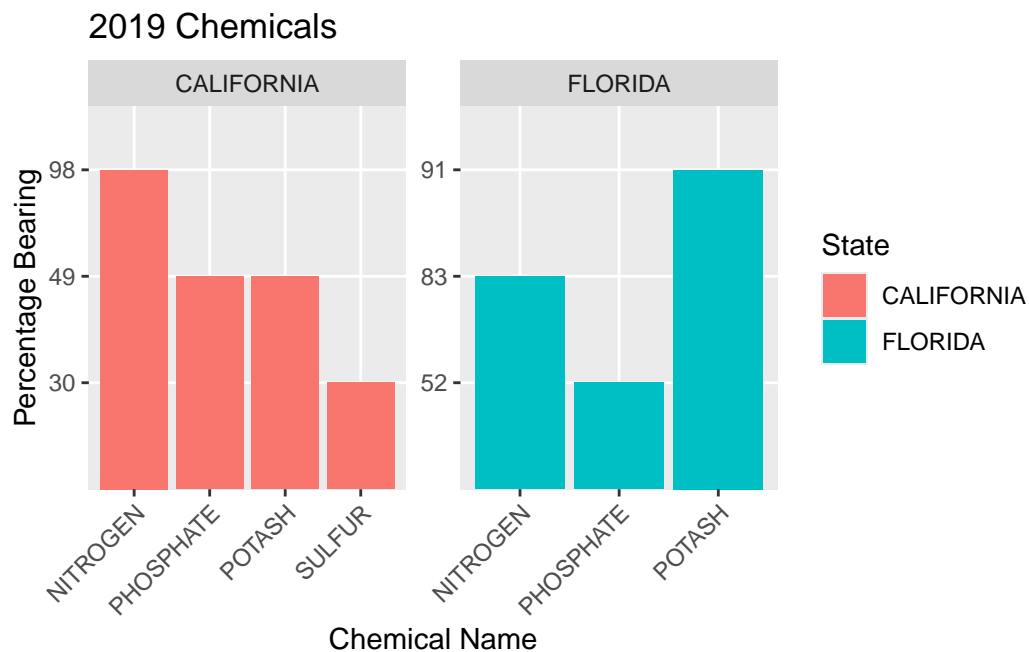
#Plot “PCT OF AREA BEARING”

```
data_chem1<- chemical %>%
  filter(Year == 2019 & State %in% c("CALIFORNIA", "FLORIDA")& chem_name %in% c("NITROGEN", "PHOSPHATE", "POTASH", "SULFUR"))
```

```

filter(Value != "(D)") %>% filter(Value != "(NA)")
#Plot the growth rate under different chemicals.
ggplot(data_chem1, aes(x = chem_name, y = Value, fill = State))+
  geom_bar(stat = "identity", position = "dodge")+
  facet_wrap(~ State, scales = "free")+
  labs(title = "2019 Chemicals",
       x = "Chemical Name",
       y = "Percentage Bearing",
       fill = "State")+
  theme(axis.text.x = element_text(angle= 45, hjust =1))

```



```

theme_minimal()

```

List of 136

```

$ line                                     :List of 6
..$ colour                               : chr "black"
..$ linewidth                             : num 0.5
..$ linetype                             : num 1
..$ lineend                               : chr "butt"
..$ arrow                                 : logi FALSE
..$ inherit.blank                         : logi TRUE
..- attr(*, "class")= chr [1:2] "element_line" "element"

```

```

$ rect                                     :List of 5
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..$ colour    : chr "black"
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..$ linetype  : num 1
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_rect" "element"
$ text                                     :List of 11
..$ family    : chr ""
..$ face      : chr "plain"
..$ colour    : chr "black"
..$ size      : num 11
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..$ vjust     : num 0.5
..$ angle     : num 0
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.. ..- attr(*, "unit")= int 8
..$ debug     : logi FALSE
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ title                                     : NULL
$ aspect.ratio : NULL
$ axis.title   : NULL
$ axis.title.x :List of 11
..$ family    : NULL
..$ face      : NULL
..$ colour    : NULL
..$ size      : NULL
..$ hjust     : NULL
..$ vjust     : num 1
..$ angle     : NULL
..$ lineheight : NULL
..$ margin    : 'margin' num [1:4] 2.75points 0points 0points 0points
.. ..- attr(*, "unit")= int 8
..$ debug     : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.title.x.top :List of 11
..$ family    : NULL
..$ face      : NULL
..$ colour    : NULL
..$ size      : NULL

```

```

..$ hjust          : NULL
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..$ angle          : NULL
..$ lineheight     : NULL
..$ margin         : 'margin' num [1:4] 0points 0points 2.75points 0points
.. ..- attr(*, "unit")= int 8
..$ debug          : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.title.x.bottom : NULL
$ axis.title.y        :List of 11
..$ family          : NULL
..$ face            : NULL
..$ colour          : NULL
..$ size            : NULL
..$ hjust           : NULL
..$ vjust           : num 1
..$ angle           : num 90
..$ lineheight      : NULL
..$ margin          : 'margin' num [1:4] 0points 2.75points 0points 0points
.. ..- attr(*, "unit")= int 8
..$ debug           : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.title.y.left   : NULL
$ axis.title.y.right  :List of 11
..$ family          : NULL
..$ face            : NULL
..$ colour          : NULL
..$ size            : NULL
..$ hjust           : NULL
..$ vjust           : num 1
..$ angle           : num -90
..$ lineheight      : NULL
..$ margin          : 'margin' num [1:4] 0points 0points 0points 2.75points
.. ..- attr(*, "unit")= int 8
..$ debug           : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.text           :List of 11
..$ family          : NULL
..$ face            : NULL
..$ colour          : chr "grey30"

```



```

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..$ vjust         : NULL
..$ angle         : NULL
..$ lineheight    : NULL
..$ margin        : NULL
..$ debug         : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.text.x          :List of 11
..$ family           : NULL
..$ face             : NULL
..$ colour           : NULL
..$ size             : NULL
..$ hjust            : NULL
..$ vjust            : num 1
..$ angle            : NULL
..$ lineheight       : NULL
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.. ..- attr(*, "unit")= int 8
..$ debug            : NULL
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..$ face             : NULL
..$ colour           : NULL
..$ size             : NULL
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..$ angle            : NULL
..$ lineheight       : NULL
..$ margin           : 'margin' num [1:4] 0points 0points 2.2points 0points
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..$ debug            : NULL
..$ inherit.blank    : logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.text.x.bottom   : NULL
$ axis.text.y          :List of 11
..$ family           : NULL
..$ face             : NULL
..$ colour           : NULL
..$ size             : NULL

```

```

..$ hjust          : num 1
..$ vjust          : NULL
..$ angle          : NULL
..$ lineheight     : NULL
..$ margin         : 'margin' num [1:4] 0points 2.2points 0points 0points
.. ..- attr(*, "unit")= int 8
..$ debug          : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.text.y.left : NULL
$ axis.text.y.right :List of 11
..$ family         : NULL
..$ face           : NULL
..$ colour         : NULL
..$ size           : NULL
..$ hjust          : num 0
..$ vjust          : NULL
..$ angle          : NULL
..$ lineheight     : NULL
..$ margin         : 'margin' num [1:4] 0points 0points 0points 2.2points
.. ..- attr(*, "unit")= int 8
..$ debug          : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.text.theta  : NULL
$ axis.text.r      :List of 11
..$ family         : NULL
..$ face           : NULL
..$ colour         : NULL
..$ size           : NULL
..$ hjust          : num 0.5
..$ vjust          : NULL
..$ angle          : NULL
..$ lineheight     : NULL
..$ margin         : 'margin' num [1:4] 0points 2.2points 0points 2.2points
.. ..- attr(*, "unit")= int 8
..$ debug          : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
$ axis.ticks       : list()
..- attr(*, "class")= chr [1:2] "element_blank" "element"
$ axis.ticks.x     : NULL
$ axis.ticks.x.top : NULL

```

```

$ axis.ticks.x.bottom      : NULL
$ axis.ticks.y             : NULL
$ axis.ticks.y.left       : NULL
$ axis.ticks.y.right      : NULL
$ axis.ticks.theta        : NULL
$ axis.ticks.r             : NULL
$ axis.minor.ticks.x.top   : NULL
$ axis.minor.ticks.x.bottom : NULL
$ axis.minor.ticks.y.left  : NULL
$ axis.minor.ticks.y.right : NULL
$ axis.minor.ticks.theta   : NULL
$ axis.minor.ticks.r       : NULL
$ axis.ticks.length        : 'simpleUnit' num 2.75points
  ..- attr(*, "unit")= int 8
$ axis.ticks.length.x      : NULL
$ axis.ticks.length.x.top  : NULL
$ axis.ticks.length.x.bottom : NULL
$ axis.ticks.length.y      : NULL
$ axis.ticks.length.y.left : NULL
$ axis.ticks.length.y.right : NULL
$ axis.ticks.length.theta  : NULL
$ axis.ticks.length.r      : NULL
$ axis.minor.ticks.length  : 'rel' num 0.75
$ axis.minor.ticks.length.x : NULL
$ axis.minor.ticks.length.x.top : NULL
$ axis.minor.ticks.length.x.bottom : NULL
$ axis.minor.ticks.length.y : NULL
$ axis.minor.ticks.length.y.left : NULL
$ axis.minor.ticks.length.y.right : NULL
$ axis.minor.ticks.length.theta : NULL
$ axis.minor.ticks.length.r : NULL
$ axis.line                : list()
  ..- attr(*, "class")= chr [1:2] "element_blank" "element"
$ axis.line.x              : NULL
$ axis.line.x.top          : NULL
$ axis.line.x.bottom       : NULL
$ axis.line.y              : NULL
$ axis.line.y.left         : NULL
$ axis.line.y.right        : NULL
$ axis.line.theta          : NULL
$ axis.line.r              : NULL
$ legend.background        : list()
  ..- attr(*, "class")= chr [1:2] "element_blank" "element"

```

```

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$ legend.spacing               : 'simpleUnit' num 11points
  ..- attr(*, "unit")= int 8
$ legend.spacing.x             : NULL
$ legend.spacing.y             : NULL
$ legend.key                   : list()
  ..- attr(*, "class")= chr [1:2] "element_blank" "element"
$ legend.key.size              : 'simpleUnit' num 1.2lines
  ..- attr(*, "unit")= int 3
$ legend.key.height            : NULL
$ legend.key.width             : NULL
$ legend.key.spacing           : 'simpleUnit' num 5.5points
  ..- attr(*, "unit")= int 8
$ legend.key.spacing.x         : NULL
$ legend.key.spacing.y         : NULL
$ legend.frame                 : NULL
$ legend.ticks                 : NULL
$ legend.ticks.length          : 'rel' num 0.2
$ legend.axis.line             : NULL
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  ..$ family                   : NULL
  ..$ face                     : NULL
  ..$ colour                   : NULL
  ..$ size                     : 'rel' num 0.8
  ..$ hjust                    : NULL
  ..$ vjust                    : NULL
  ..$ angle                    : NULL
  ..$ lineheight               : NULL
  ..$ margin                   : NULL
  ..$ debug                    : NULL
  ..$ inherit.blank: logi TRUE
  ..- attr(*, "class")= chr [1:2] "element_text" "element"
$ legend.text.position          : NULL
$ legend.title                 :List of 11
  ..$ family                   : NULL
  ..$ face                     : NULL
  ..$ colour                   : NULL
  ..$ size                     : NULL
  ..$ hjust                    : num 0
  ..$ vjust                    : NULL
  ..$ angle                    : NULL
  ..$ lineheight               : NULL

```

```

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..$ debug           : NULL
..$ inherit.blank: logi TRUE
..- attr(*, "class")= chr [1:2] "element_text" "element"
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$ legend.position.inside : NULL
$ legend.direction      : NULL
$ legend.byrow          : NULL
$ legend.justification  : chr "center"
$ legend.justification.top : NULL
$ legend.justification.bottom : NULL
$ legend.justification.left : NULL
$ legend.justification.right : NULL
$ legend.justification.inside : NULL
$ legend.location       : NULL
$ legend.box            : NULL
$ legend.box.just       : NULL
$ legend.box.margin     : 'margin' num [1:4] 0cm 0cm 0cm 0cm
..- attr(*, "unit")= int 1
$ legend.box.background : list()
..- attr(*, "class")= chr [1:2] "element_blank" "element"
$ legend.box.spacing    : 'simpleUnit' num 11points
..- attr(*, "unit")= int 8
[list output truncated]
- attr(*, "class")= chr [1:2] "theme" "gg"
- attr(*, "complete")= logi TRUE
- attr(*, "validate")= logi TRUE

```

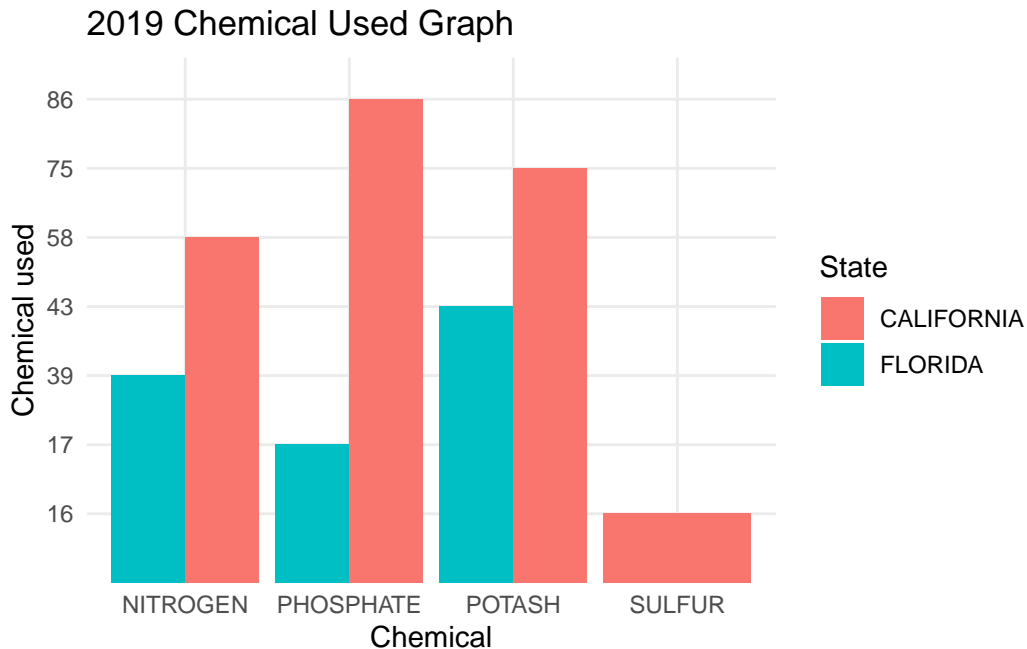
#Plot “LB / ACRE / YEAR”

```

data_chem2<- chemical %>%
  filter(Year == 2019 & State %in% c("CALIFORNIA", "FLORIDA"))& chem_name %in% c("NITROGEN", "PHOSPHORUS")
filter(Value != "(D)") %>% filter(Value != "(NA)")

#Plot the use of chemical graph
ggplot(data_chem2, aes(x = chem_name, y = Value, fill = State))+
  geom_bar(stat = "identity", position = "dodge")+
  labs(title = "2019 Chemical Used Graph",
       x = "Chemical",
       y = "Chemical used")+
  theme_minimal()

```



Compare the two plots, we can easily find that California uses more chemicals than Florida in 2019. Among them, California uses Nitrogen most while Florida uses Potash most.

Different chemicals have different effects to strawberry bearing. While some may restrict the growth, others can help increase it. For example, enough Nitrogen has positive effects to the area bearing, but too much Potash makes the area bearing smaller.

#Question 2: How does the Fungicide affect the strawberry growth?

Chemicals are used in strawberry Sterilization. I wrote a new csv file named “new1” for my EDA. It contains all information about chemical types and names.

Take AZOXYSTROBIN as an example, lets check if it is hazard to humans.

```
chem<- read.csv("new1.csv")

chem_data<- chem %>%
  filter(type== "FUNGICIDE")%>% filter(Value != "(D)") %>% filter(Value != "(NA)")
head(chem_data)
```

	Year	State	mk1	mk2	measure	other	type
1	2023	CALIFORNIA	BEARING	APPLICATIONS	LB	<NA>	FUNGICIDE
2	2023	CALIFORNIA	BEARING	APPLICATIONS	LB	<NA>	FUNGICIDE
3	2023	CALIFORNIA	BEARING	APPLICATIONS	LB	<NA>	FUNGICIDE
4	2023	CALIFORNIA	BEARING	APPLICATIONS	LB	<NA>	FUNGICIDE

```

5 2023 CALIFORNIA BEARING APPLICATIONS      LB <NA> FUNGICIDE
6 2023 CALIFORNIA BEARING APPLICATIONS      LB <NA> FUNGICIDE
      chem_name chem_index  Value
1      AZOXYSTROBIN      128810  3,300
2 BORAX DECAHYDRATE      11102   2,800
3      BOSCALID      128008   6,600
4      CAPTAN      81301 603,100
5      CYPRODINIL      288202 30,300
6      FENHEXAMID      90209   8,600

```

```
#Compare the use of Azoxystrobin
```

```
chem_data1<- chem_data %>% filter(State %in% c("FLORIDA", "CALIFORNIA") & mk2=="TREATED" & c
chem_data1
```

```

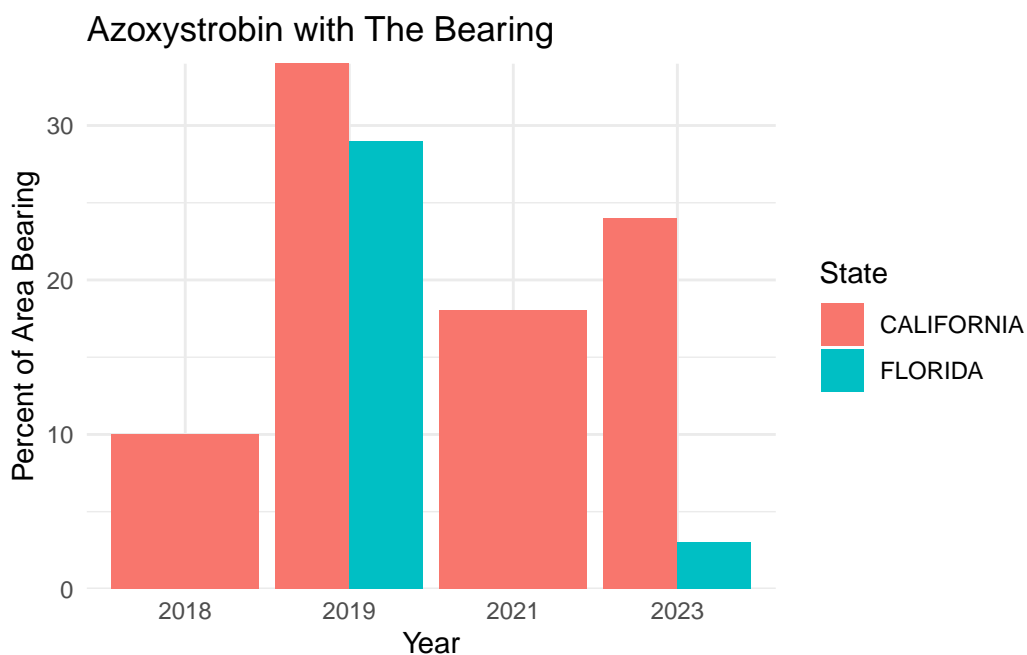
      Year      State      mk1      mk2      measure other      type
1 2023 CALIFORNIA BEARING TREATED PCT OF AREA BEARING  AVG FUNGICIDE
2 2023  FLORIDA BEARING TREATED PCT OF AREA BEARING  AVG FUNGICIDE
3 2021 CALIFORNIA BEARING TREATED PCT OF AREA BEARING  AVG FUNGICIDE
4 2019 CALIFORNIA BEARING TREATED PCT OF AREA BEARING  AVG FUNGICIDE
5 2019  FLORIDA BEARING TREATED PCT OF AREA BEARING  AVG FUNGICIDE
6 2018 CALIFORNIA BEARING TREATED PCT OF AREA BEARING  AVG FUNGICIDE
      chem_name chem_index Value
1 AZOXYSTROBIN      128810    24
2 AZOXYSTROBIN      128810     3
3 AZOXYSTROBIN      128810    18
4 AZOXYSTROBIN      128810    34
5 AZOXYSTROBIN      128810    29
6 AZOXYSTROBIN      128810    10

```

```

chem_data1$Year <- as.factor(chem_data1$Year)
chem_data1$Value <- as.numeric(chem_data1$Value)
ggplot(chem_data1, aes(x= Year, y= Value, fill= State))+
  geom_bar(stat = "identity", position = "dodge")+
  scale_y_continuous(limits = c(0, max(chem_data1$Value, na.rm = TRUE)), expand = c(0, 0))+
  labs(title = "Azoxystrobin with The Bearing",
       x = "Year",
       y = "Percent of Area Bearing")+
  theme_minimal()

```



From the graph, we can find that Florida didn't use Azoxystrobin in 2018, and 2021.

```
chem_data<- chem %>%
  filter(type== "FUNGICIDE")%>% filter(Value != "(D)") %>% filter(Value != "(NA)")
head(chem_data)
```

	Year	State	mk1	mk2	measure	other	type
1	2023	CALIFORNIA	BEARING APPLICATIONS		LB	<NA>	FUNGICIDE
2	2023	CALIFORNIA	BEARING APPLICATIONS		LB	<NA>	FUNGICIDE
3	2023	CALIFORNIA	BEARING APPLICATIONS		LB	<NA>	FUNGICIDE
4	2023	CALIFORNIA	BEARING APPLICATIONS		LB	<NA>	FUNGICIDE
5	2023	CALIFORNIA	BEARING APPLICATIONS		LB	<NA>	FUNGICIDE
6	2023	CALIFORNIA	BEARING APPLICATIONS		LB	<NA>	FUNGICIDE

	chem_name	chem_index	Value
1	AZOXYSTROBIN	128810	3,300
2	BORAX DECAHYDRATE	11102	2,800
3	BOSCALID	128008	6,600
4	CAPTAN	81301	603,100
5	CYPRODINIL	288202	30,300
6	FENHEXAMID	90209	8,600

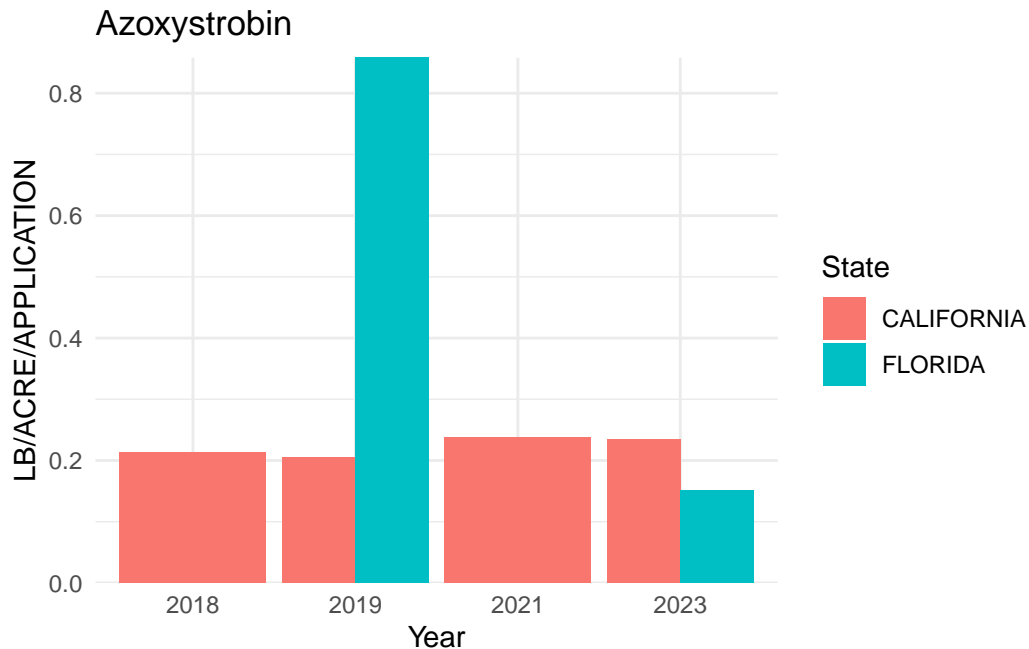

```
chem_data1<- chem_data %>% filter(State %in% c("FLORIDA", "CALIFORNIA") & measure=="LB / ACRE")
chem_data1
```

	Year	State	mk1	mk2	measure	other	type
1	2023	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG FUNGICIDE		
2	2023	FLORIDA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG FUNGICIDE		
3	2021	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG FUNGICIDE		
4	2019	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG FUNGICIDE		
5	2019	FLORIDA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG FUNGICIDE		
6	2018	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG FUNGICIDE		

	chem_name	chem_index	Value
1	AZOXYSTROBIN	128810	0.234
2	AZOXYSTROBIN	128810	0.151
3	AZOXYSTROBIN	128810	0.237
4	AZOXYSTROBIN	128810	0.205
5	AZOXYSTROBIN	128810	0.858
6	AZOXYSTROBIN	128810	0.213

```
chem_data1$Year <- as.factor(chem_data1$Year)
chem_data1$Value <- as.numeric(chem_data1$Value)

ggplot(chem_data1, aes(x= Year, y= Value, fill= State))+
  geom_bar(stat = "identity", position = "dodge")+
  scale_y_continuous(limits = c(0, max(chem_data1$Value, na.rm = TRUE)), expand = c(0, 0))+
  labs(title = "Azoxystrobin",
       x = "Year",
       y = "LB/ACRE/APPLICATION")+
  theme_minimal()
```



From the research, the best rate for Azoxystrobin is about 0.65-0.975 lb/Acre/Application. Using the histogram, we can find the highest rates are below the limit, so we can conclude that the use of it is normative.

However, it is hard to predict the relationship between the use of Azoxystrobin and the bearing rate of strawberries. In our histograms, we can find that the use of Azoxystrobin in California are similar in 2018 and 2019, but the bearing area under it have a large difference between 2018 and 2019. However, in Florida, although there is 0.8 lb/Acre/Application, the percent of area bearing is still higher. Therefore, there might be other factors that affect the growth of strawberries.

#Question 3: How does a hazard pesticide affect the growth of strawberries?

Next goal is trying to explore the use of deadly carcinogens. Malathion is an [organophosphate insecticide](#) which acts as an [acetylcholinesterase inhibitor](#). We start by choosing the true columns.

```
chem_data2<- chem %>% filter(type=="INSECTICIDE") %>% filter(State %in% c("FLORIDA", "CALIFORNIA"))
chem_data2
```

	Year	State	mk1	mk2	measure	other
1	2023	CALIFORNIA	BEARING APPLICATIONS		LB	<NA>
2	2023	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION		AVG
3	2023	CALIFORNIA	BEARING APPLICATIONS		LB / ACRE / YEAR	AVG

4	2023	CALIFORNIA	BEARING APPLICATIONS	NUMBER	AVG
5	2023	CALIFORNIA	BEARING TREATED	PCT OF AREA BEARING	AVG
6	2023	FLORIDA	BEARING APPLICATIONS	LB	<NA>
7	2023	FLORIDA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG
8	2023	FLORIDA	BEARING APPLICATIONS	LB / ACRE / YEAR	AVG
9	2023	FLORIDA	BEARING APPLICATIONS	NUMBER	AVG
10	2023	FLORIDA	BEARING TREATED	PCT OF AREA BEARING	AVG
11	2021	CALIFORNIA	BEARING APPLICATIONS	LB	<NA>
12	2021	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG
13	2021	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / YEAR	AVG
14	2021	CALIFORNIA	BEARING APPLICATIONS	NUMBER	AVG
15	2021	CALIFORNIA	BEARING TREATED	PCT OF AREA BEARING	AVG
16	2021	FLORIDA	BEARING APPLICATIONS	LB	<NA>
17	2021	FLORIDA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG
18	2021	FLORIDA	BEARING APPLICATIONS	LB / ACRE / YEAR	AVG
19	2021	FLORIDA	BEARING APPLICATIONS	NUMBER	AVG
20	2021	FLORIDA	BEARING TREATED	PCT OF AREA BEARING	AVG
21	2019	CALIFORNIA	BEARING APPLICATIONS	LB	<NA>
22	2019	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG
23	2019	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / YEAR	AVG
24	2019	CALIFORNIA	BEARING APPLICATIONS	NUMBER	AVG
25	2019	CALIFORNIA	BEARING TREATED	PCT OF AREA BEARING	AVG
26	2019	FLORIDA	BEARING APPLICATIONS	LB	<NA>
27	2019	FLORIDA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG
28	2019	FLORIDA	BEARING APPLICATIONS	LB / ACRE / YEAR	AVG
29	2019	FLORIDA	BEARING APPLICATIONS	NUMBER	AVG
30	2019	FLORIDA	BEARING TREATED	PCT OF AREA BEARING	AVG
31	2018	CALIFORNIA	BEARING APPLICATIONS	LB	<NA>
32	2018	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG
33	2018	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / YEAR	AVG
34	2018	CALIFORNIA	BEARING APPLICATIONS	NUMBER	AVG
35	2018	CALIFORNIA	BEARING TREATED	PCT OF AREA BEARING	AVG
36	2018	FLORIDA	BEARING APPLICATIONS	LB	<NA>
37	2018	FLORIDA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG
38	2018	FLORIDA	BEARING APPLICATIONS	LB / ACRE / YEAR	AVG
39	2018	FLORIDA	BEARING APPLICATIONS	NUMBER	AVG
40	2018	FLORIDA	BEARING TREATED	PCT OF AREA BEARING	AVG

	type	chem_name	chem_index	Value
1	INSECTICIDE	MALATHION	57701	19,400
2	INSECTICIDE	MALATHION	57701	1.807
3	INSECTICIDE	MALATHION	57701	2.398
4	INSECTICIDE	MALATHION	57701	1.3
5	INSECTICIDE	MALATHION	57701	19

6	INSECTICIDE	MALATHION	57701	(D)
7	INSECTICIDE	MALATHION	57701	(D)
8	INSECTICIDE	MALATHION	57701	(D)
9	INSECTICIDE	MALATHION	57701	(D)
10	INSECTICIDE	MALATHION	57701	(D)
11	INSECTICIDE	MALATHION	57701	29,100
12	INSECTICIDE	MALATHION	57701	2.027
13	INSECTICIDE	MALATHION	57701	2.443
14	INSECTICIDE	MALATHION	57701	1.2
15	INSECTICIDE	MALATHION	57701	30
16	INSECTICIDE	MALATHION	57701	(D)
17	INSECTICIDE	MALATHION	57701	(D)
18	INSECTICIDE	MALATHION	57701	(D)
19	INSECTICIDE	MALATHION	57701	(D)
20	INSECTICIDE	MALATHION	57701	(D)
21	INSECTICIDE	MALATHION	57701	56,700
22	INSECTICIDE	MALATHION	57701	1.991
23	INSECTICIDE	MALATHION	57701	5.571
24	INSECTICIDE	MALATHION	57701	2.8
25	INSECTICIDE	MALATHION	57701	28
26	INSECTICIDE	MALATHION	57701	(D)
27	INSECTICIDE	MALATHION	57701	(D)
28	INSECTICIDE	MALATHION	57701	(D)
29	INSECTICIDE	MALATHION	57701	(D)
30	INSECTICIDE	MALATHION	57701	(D)
31	INSECTICIDE	MALATHION	57701	8,000
32	INSECTICIDE	MALATHION	57701	1.799
33	INSECTICIDE	MALATHION	57701	3.062
34	INSECTICIDE	MALATHION	57701	1.7
35	INSECTICIDE	MALATHION	57701	7
36	INSECTICIDE	MALATHION	57701	(D)
37	INSECTICIDE	MALATHION	57701	(D)
38	INSECTICIDE	MALATHION	57701	(D)
39	INSECTICIDE	MALATHION	57701	(D)
40	INSECTICIDE	MALATHION	57701	(D)

```
chem_data2<- chem_data2 %>% filter(measure== "LB / ACRE / APPLICATION") %>% filter(Value != 0)
chem_data2
```

	Year	State	mk1	mk2	measure	other
1	2023	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG	
2	2021	CALIFORNIA	BEARING APPLICATIONS	LB / ACRE / APPLICATION	AVG	

```

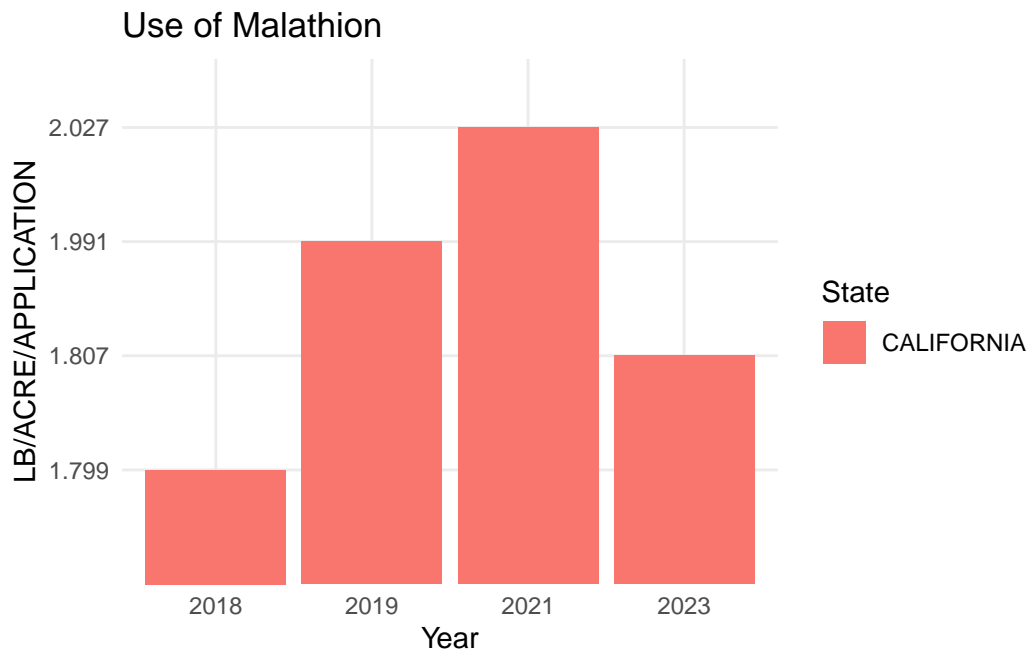
3 2019 CALIFORNIA BEARING APPLICATIONS LB / ACRE / APPLICATION    AVG
4 2018 CALIFORNIA BEARING APPLICATIONS LB / ACRE / APPLICATION    AVG
      type chem_name chem_index Value
1 INSECTICIDE MALATHION      57701 1.807
2 INSECTICIDE MALATHION      57701 2.027
3 INSECTICIDE MALATHION      57701 1.991
4 INSECTICIDE MALATHION      57701 1.799

```

```

#
chem_data2$Year <- as.factor(chem_data2$Year)
ggplot(chem_data2, aes(x= Year, y= Value, fill= State))+
  geom_bar(stat = "identity", position = "dodge")+
  labs(title = "Use of Malathion",
       x = "Year",
       y = "LB/ACRE/APPLICATION")+
  theme_minimal()

```



```

#Input hazard information

GHS_searcher<-function(result_json_object){
  result<-result_json_object
  for (i in 1:length(result[["result"]][["Hierarchies"]][["Hierarchy"]])){

```

```

    if(result[["result"]][["Hierarchies"]][["Hierarchy"]][[i]][["SourceName"]]=="GHS Classif
      return(i)
    }
  }
}

hazards_retriever<-function(index,result_json_object){
  result<-result_json_object
  hierarchy<-result[["result"]][["Hierarchies"]][["Hierarchy"]][[index]]
  i<-1
  output_list<-rep(NA,length(hierarchy[["Node"]]))
  while(str_detect(hierarchy[["Node"]][[i]][["Information"]][["Name"]], "H") & i<length(hiera
    output_list[i]<-hierarchy[["Node"]][[i]][["Information"]][["Name"]]
    i<-i+1
  }
  return(output_list[!is.na(output_list)])
}
chemical_vec<-c("azoxystrobin","malathion")
#
result_f<-get_pug_rest(identifier = "azoxystrobin", namespace = "name", domain = "compound",
hazards_retriever(GHS_searcher(result_f),result_f)

```

```

[1] "H331: Toxic if inhaled [Danger Acute toxicity, inhalation]"
[2] "H300: Health Hazards"
[3] "Hazard Statement Codes"
[4] "H370: Causes damage to organs [Danger Specific target organ toxicity, single exposure]"
[5] "H400: Very toxic to aquatic life [Warning Hazardous to the aquatic environment, acute h
[6] "H400: Environmental Hazards"
[7] "H410: Very toxic to aquatic life with long lasting effects [Warning Hazardous to the aqu

```

```

result_d<-get_pug_rest(identifier = "malathion", namespace = "name", domain = "compound",ope
hazards_retriever(GHS_searcher(result_d),result_d)

```

```

[1] "H302: Harmful if swallowed [Warning Acute toxicity, oral]"
[2] "H300: Health Hazards"
[3] "Hazard Statement Codes"
[4] "H317: May cause an allergic skin reaction [Warning Sensitization, Skin]"
[5] "H320: Causes eye irritation [Warning Serious eye damage/eye irritation]"

```

```
[6] "H331: Toxic if inhaled [Danger Acute toxicity, inhalation]"
[7] "H341: Suspected of causing genetic defects [Warning Germ cell mutagenicity]"
[8] "H350: May cause cancer [Danger Carcinogenicity]"
[9] "H370: Causes damage to organs [Danger Specific target organ toxicity, single exposure]"
[10] "H372: Causes damage to organs through prolonged or repeated exposure [Danger Specific target organ toxicity, repeated exposure]"
[11] "H373: May causes damage to organs through prolonged or repeated exposure [Warning Specific target organ toxicity, repeated exposure]"
[12] "H400: Very toxic to aquatic life [Warning Hazardous to the aquatic environment, acute toxicity]"
[13] "H400: Environmental Hazards"
[14] "H410: Very toxic to aquatic life with long lasting effects [Warning Hazardous to the aquatic environment, long term toxicity]"
```

According to the records, Malathion can damage organs and genetic defects as well as being environment hazards, the best rate for using is 1.25 - 2.5 lb/ Acre/ Application. In California, the highest rate is 2.027 lb/ Acre/ Application while the lowest rate is 1.799 lb/Acre/Application, both of them are under the limit. We can then explore the bearing area to find the relationship between them.

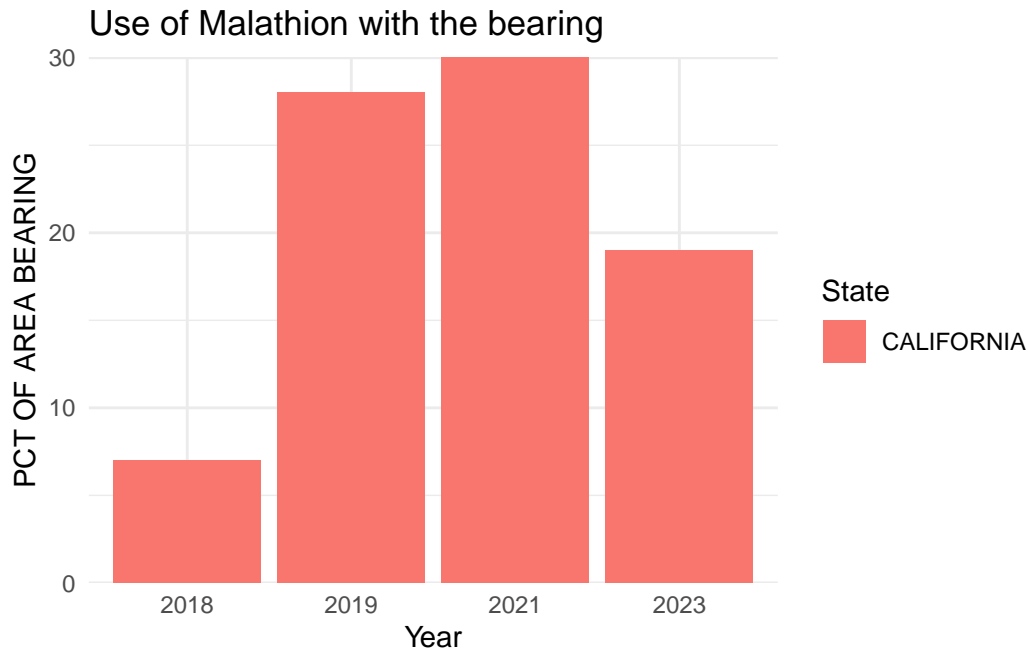
```
chem_data3<- chem %>% filter(type=="INSECTICIDE") %>% filter(State %in% c("FLORIDA", "CALIFORNIA"))
chem_data3<- chem_data3 %>% filter(measure== "PCT OF AREA BEARING") %>% filter(Value != "(D)")
chem_data3
```

	Year	State	mk1	mk2	measure	other	type
1	2023	CALIFORNIA	BEARING	TREATED	PCT OF AREA BEARING	AVG	INSECTICIDE
2	2021	CALIFORNIA	BEARING	TREATED	PCT OF AREA BEARING	AVG	INSECTICIDE
3	2019	CALIFORNIA	BEARING	TREATED	PCT OF AREA BEARING	AVG	INSECTICIDE
4	2018	CALIFORNIA	BEARING	TREATED	PCT OF AREA BEARING	AVG	INSECTICIDE

	chem_name	chem_index	Value
1	MALATHION	57701	19
2	MALATHION	57701	30
3	MALATHION	57701	28
4	MALATHION	57701	7

```
chem_data3$Year <- as.factor(chem_data3$Year)
chem_data3$Value <- as.numeric(chem_data3$Value)

ggplot(chem_data3, aes(x= Year, y= Value, fill= State))+
  geom_bar(stat = "identity", position = "dodge")+
  scale_y_continuous(limits = c(0, max(chem_data3$Value, na.rm = TRUE)), expand = c(0, 0))+
  labs(title = "Use of Malathion with the bearing",
       x = "Year",
       y = "PCT OF AREA BEARING")+
  theme_minimal()
```



We can find that the use of Malathion truly has some good effect to the bearing rate, even though it is dangerous to humans. For instance, in 2018, the use of Malathion was little, leading to the low bearing rate, this is because Malathion is a great insecticide, in that year, little use of Malathion could not restrict the pests.

Chemicals used in strawberry cultivaion

Six deadly carcinogens from WHO list

[captafol](#)

[ethylene dibromide](#) also

[glyphosate](#) See also 1

2

3

4

[malathion](#) 1 2

[diazinon](#) 1 2 3

[dichlorophenyltrichloroethane \(DDT\)](#) 1 2 [3]([https://www.epa.gov/ingredients-used-pesticide-products/ddt-brief-history-and-status#:~:text=DDT%20\(dichloro%2Ddiphenyl%2Dtrichloroethane,both%20mi](https://www.epa.gov/ingredients-used-pesticide-products/ddt-brief-history-and-status#:~:text=DDT%20(dichloro%2Ddiphenyl%2Dtrichloroethane,both%20mi)

For contrast

[Azadirachtin 1 2 3](#)

Sources of agricultural chemical information

for EPA number lookup [epa numbers](#)

[Active Pesticide Product Registration Informational Listing](#)

[CAS for Methyl Bromide](#)

[pesticide chemical search](#)

[toxic chemical dashboard](#)

[pubChem](#)

The EPA PC (Pesticide Chemical) Code is a unique chemical code number assigned by the EPA to a particular pesticide active ingredient, inert ingredient or mixture of active ingredients.

Investigating toxic pesticides

[start here with chem PC code](#)

[step 2](#) to get label (with warnings) for products using the chemical

[Pesticide Product and Label System](#)

[Search by Chemical](#)

[CompTox Chemicals Dashboard](#)

[Active Pesticide Product Registration Informational Listing](#)

[OSHA chemical database](#)

[Pesticide Ingredients](#)

[NPIC Product Research Online \(NPRO\)](#)

[Databases for Chemical Information](#)

[Pesticide Active Ingredients](#)

[TSCA Chemical Substance Inventory](#)

[glyphosate](#)