

# 415\_Midterm\_Project

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## Before Starting the Project

### Data Set

The csv file named “strawb\_mar6.csv”, which contains strawberry cultivation data in different states and measurements.

### Main Focus

1. Pick three chemical treatments used for conventional strawberries in both states and contrast their use. Try to find chemicals with divergent use patterns between the states. Produce, tables, plots, and descriptions of the chemicals, how they are used, and how their use differs between California and Florida.
2. Compare the production and sales of organic and conventional strawberries and strawberries sold for processing. Show differences in price and volume between California and Florida. How do price, cost, and volume relationships change over the years?

### Data Cleaning

Install packages

```
# install required packages
#install.packages("knitr")
#install.packages("kableExtra")
#install.packages("tidyverse")
#install.packages("stringr")
#install.packages("ggplot2")
```

```
library(knitr)
library(kableExtra)
library(tidyverse)
```

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr      1.1.4      v readr      2.1.5
v forcats    1.0.0      v stringr    1.5.1
v ggplot2    3.5.1      v tibble     3.2.1
v lubridate  1.9.4      v tidyr      1.3.1
v purrr      1.0.2

-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter()      masks stats::filter()
x dplyr::group_rows()  masks kableExtra::group_rows()
x dplyr::lag()          masks stats::lag()
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become
```

```
library(stringr)
library(ggplot2)
```

Read File (From USDA-NASS starwberries.qmd)

```
# Label for the code chunk
#| label: read data - glimpse

# Read the CSV file "strawb_mar6.csv" into a data frame called 'strawberry'
# - 'col_names = TRUE' tells read_csv to treat the first row as column names
# - 'show_col_types = FALSE' suppresses printing of column type information
strawberry <- read_csv("strawb_mar6.csv",
                      col_names = TRUE,
                      show_col_types = FALSE)

# Source the file "my_functions.R" to load custom functions defined in that script
source("my_functions.R")

# remove columns from a data frame that contain only a single unique value
strawb <- strawberry |> drop_one_value_col()
```

Exploring Data

```

# assume data is a tibble
# n_show is the number of rows to show

show_unique <- function(data, n_rows=10 ){
  # make a tibble items to hold the data to show
  # browser()
  a <- n_rows * dim(data)[2] # number of cells in items
  items <- rep(" ", a) # items will coerce everything to char
  dim(items) <- c(n_rows ,dim(data)[2]) # shape items
  items <- as_tibble(items)
  colnames(items) <- colnames(data)
  # browser()
  for(i in 1:dim(data)[2]){

    col_items <- unique(data[,i])
    # row_ex is the number of rows needed
    # to make the column length conformable with items
    row_ex <- n_rows - dim(col_items)[1]
    if(row_ex >= 0){
      ex_rows <- tibble(rep(" ",row_ex))
      colnames(ex_rows) <- colnames(col_items)
      col_add <- rbind2(col_items, ex_rows)

    } else if(row_ex < 0){
      col_add <- col_items[1:10,]

    }

    items[,i] <- col_add

  }

  return(items)
}

#test <- show_unique(strawb, 10)

```

```

#|label: split strawb into census and survey pieces

strw_census <- strawb |> filter(Program == "CENSUS")

strw_survey <- strawb |> filter(Program == "SURVEY")

```

```
nrow(strawb) == (nrow(strw_census) + nrow(strw_survey))
```

```
[1] TRUE
```

```
# Remove columns that contain only a single unique value
s_census <- strw_census |> drop_one_value_col(prt_val = TRUE)
```

```
[1] "Looking for single value columns in data frame: strw_census"
```

```
[1] "Columns dropped:"
```

Program	Period	Week Ending
"CENSUS"	"YEAR"	NA

```
s_survey <- strw_survey |> drop_one_value_col(prt_val = TRUE)
```

```
[1] "Looking for single value columns in data frame: strw_survey"
```

```
[1] "Columns dropped:"
```

Program	Commodity	CV (%)
"SURVEY"	"STRAWBERRIES"	NA

```
# Preview up to 10 unique values per column in each data set
```

```
unique_cen <- s_census |> show_unique(nrows = 10)
```

```
unique_sur <- s_survey |> show_unique(nrows = 10)
```

```
# Drop redundant or unnecessary Data
```

```
strw_census <- s_census |> select(-`State ANSI`)
```

```
# Remove 'State ANSI', 'Week Ending', and 'Period' for Data Cleaning
```

```
strw_survey <- s_survey |> select(-`State ANSI`, -`Week Ending`, -Period)
```

```
# Removing intermediate and temporary data set
```

```
rm(s_census, s_survey, strawberry, items)
```

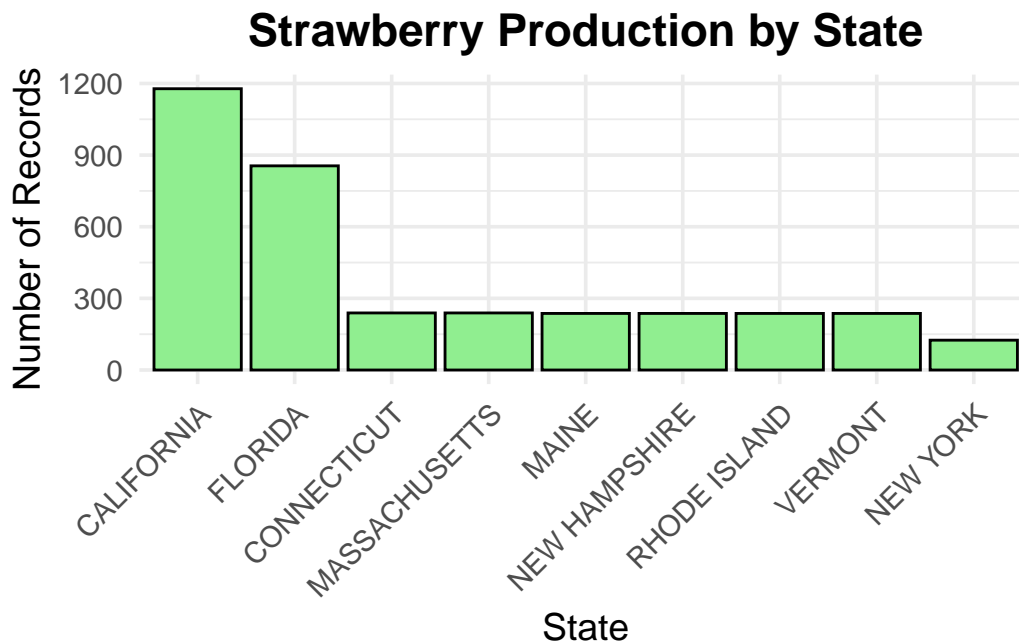
## Strawberry Growth Location

```
# Starberries' grown place :)
```

```
state_all <- strawb |> distinct(State)
```

```
state_all1 <- strawb |> group_by(State) |> count()
```

```
# Improved bar plot
ggplot(state_all1, aes(x = reorder(State, -n), y = n)) +
  geom_bar(stat = "identity", fill = "lightgreen", color = "black") +
  labs(
    title = "Strawberry Production by State",
    x = "State",
    y = "Number of Records"
  ) +
  theme_minimal(base_size = 14) +
  theme(
    axis.text.x = element_text(angle = 45, hjust = 1),
    plot.title = element_text(face = "bold", hjust = 0.5)
  )
)
```



```
# Show unique values in the 'Domain' column from both datasets
unique(strw_census$Domain)
```

```
[1] "NET GAIN"          "TOTAL"             "NET LOSS"
[4] "AREA OPERATED"    "ECONOMIC CLASS"    "FARM SALES"
[7] "NAICS CLASSIFICATION" "ORGANIC STATUS"
```

```
unique(strw_survey$Domain)
```

```
[1] "TOTAL"                "CHEMICAL, FUNGICIDE"  "CHEMICAL, INSECTICIDE"
[4] "CHEMICAL, OTHER"      "CHEMICAL, HERBICIDE"  "FERTILIZER"
```

## Part 1: Chemical Analysis

### Gather Chemical Information Data

```
#Select all the rows that contains chemical use information
chemical <- strw_survey[str_detect(strw_survey$`Domain Category`, regex("chemical", ignore_case))]

#Taking only CA and FL state data of chemical usage
chemical_CA <- chemical[chemical$State == "CALIFORNIA", ]
chemical_FL <- chemical[chemical$State == "FLORIDA", ]
print(chemical_CA)
```

```
# A tibble: 1,011 x 9
   Year State   Fruit   Category Item  Metric Domain `Domain Category` Value
  <dbl> <chr>   <chr>   <chr>   <chr> <chr>  <chr>   <chr>          <chr>
1  2023 CALIFORNIA STRAWB~ " MEASU~ <NA> <NA>  CHEMI~ CHEMICAL, FUNGIC~ (D)
2  2023 CALIFORNIA STRAWB~ " MEASU~ <NA> <NA>  CHEMI~ CHEMICAL, INSECT~ (D)
3  2023 CALIFORNIA STRAWB~ " MEASU~ <NA> <NA>  CHEMI~ CHEMICAL, INSECT~ (D)
4  2023 CALIFORNIA STRAWB~ " MEASU~ <NA> <NA>  CHEMI~ CHEMICAL, OTHER:~ (NA)
5  2023 CALIFORNIA STRAWB~ " MEASU~ " AV~ <NA>  CHEMI~ CHEMICAL, FUNGIC~ (D)
6  2023 CALIFORNIA STRAWB~ " MEASU~ " AV~ <NA>  CHEMI~ CHEMICAL, INSECT~ (D)
7  2023 CALIFORNIA STRAWB~ " MEASU~ " AV~ <NA>  CHEMI~ CHEMICAL, INSECT~ (D)
8  2023 CALIFORNIA STRAWB~ " MEASU~ " AV~ <NA>  CHEMI~ CHEMICAL, OTHER:~ (NA)
9  2023 CALIFORNIA STRAWB~ " MEASU~ " AV~ <NA>  CHEMI~ CHEMICAL, FUNGIC~ (D)
10 2023 CALIFORNIA STRAWB~ " MEASU~ " AV~ <NA>  CHEMI~ CHEMICAL, INSECT~ (D)
# i 1,001 more rows
```

```
print(chemical_FL)
```

```
# A tibble: 691 x 9
   Year State   Fruit   Category Item  Metric Domain `Domain Category` Value
  <dbl> <chr>   <chr>   <chr>   <chr> <chr>  <chr>   <chr>          <chr>
1  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA>  CHEMI~ CHEMICAL, FUNGIC~ (D)
```

```

2  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, FUNGIC~ (D)
3  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, FUNGIC~ (D)
4  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, HERBIC~ (D)
5  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, HERBIC~ (D)
6  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, HERBIC~ (D)
7  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, INSECT~ (D)
8  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, INSECT~ (D)
9  2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, INSECT~ (D)
10 2023 FLORIDA STRAWBERR~ " MEASU~ <NA> <NA> CHEMI~ CHEMICAL, OTHER:~ (D)
# i 681 more rows

```

```
rm(chemical)
```

## Chemical 1: Captan

```

# First we check the use of Captan
# Use library of stringr functions to select the captan information
# Filter rows from chemical_CA
captan_CA <- chemical_CA[str_detect(chemical_CA$`Domain Category`, regex("captan", ignore_case))]

# Filter rows from chemical_FL
captan_FL <- chemical_FL[str_detect(chemical_FL$`Domain Category`, regex("captan", ignore_case))]

captan_combined <- rbind(captan_CA, captan_FL)

rm(captan_CA)
rm(captan_FL)

print(captan_combined)

```

```
# A tibble: 20 x 9
```

	Year	State	Fruit	Category	Item	Metric	Domain	`Domain Category`	Value
	<dbl>	<chr>	<chr>	<chr>	<chr>	<chr>	<chr>	<chr>	<chr>
1	2023	CALIFORNIA	STRAWB~	" BEARI~	" ME~	<NA>	CHEMI~	CHEMICAL, FUNGIC~	603,~
2	2023	CALIFORNIA	STRAWB~	" BEARI~	" ME~	" AVG"	CHEMI~	CHEMICAL, FUNGIC~	1.693
3	2023	CALIFORNIA	STRAWB~	" BEARI~	" ME~	" AVG"	CHEMI~	CHEMICAL, FUNGIC~	15.9~
4	2023	CALIFORNIA	STRAWB~	" BEARI~	" ME~	" AVG"	CHEMI~	CHEMICAL, FUNGIC~	9.4
5	2023	CALIFORNIA	STRAWB~	" BEARI~	" ME~	" AVG"	CHEMI~	CHEMICAL, FUNGIC~	88
6	2021	CALIFORNIA	STRAWB~	" BEARI~	" ME~	<NA>	CHEMI~	CHEMICAL, FUNGIC~	253,~
7	2021	CALIFORNIA	STRAWB~	" BEARI~	" ME~	" AVG"	CHEMI~	CHEMICAL, FUNGIC~	1.662

```

8 2021 CALIFORNIA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 14.2~
9 2021 CALIFORNIA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 8.6
10 2021 CALIFORNIA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 45
11 2023 FLORIDA STRAWB~ " BEARI~ " ME~ <NA> CHEMI~ CHEMICAL, FUNGIC~ 144,~
12 2023 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 2.012
13 2023 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 10.5~
14 2023 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 5.2
15 2023 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 96
16 2021 FLORIDA STRAWB~ " BEARI~ " ME~ <NA> CHEMI~ CHEMICAL, FUNGIC~ 135,~
17 2021 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 2.025
18 2021 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 21.3~
19 2021 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 10.5
20 2021 FLORIDA STRAWB~ " BEARI~ " ME~ " AVG" CHEMI~ CHEMICAL, FUNGIC~ 61

```

```
# Now we get the table of captan use data in both California and Florida
```

```
# now we start to reorganize the data in the order of category at first, Item for the second
```

```

captan <- captan_combined |> group_by(Category, Item, State, Year) |> count(Value)

print(captan)

```

```
# A tibble: 20 x 6
```

```
# Groups:   Category, Item, State, Year [20]
```

	Category <chr>	Item <chr>	State <chr>	Year <dbl>	Value <chr>	n <int>
1	" BEARING - APPLICATIONS"	" MEASURED IN LB"	CALI~	2021	253,~	1
2	" BEARING - APPLICATIONS"	" MEASURED IN LB"	CALI~	2023	603,~	1
3	" BEARING - APPLICATIONS"	" MEASURED IN LB"	FLOR~	2021	135,~	1
4	" BEARING - APPLICATIONS"	" MEASURED IN LB"	FLOR~	2023	144,~	1
5	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2021	1.662	1
6	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2023	1.693	1
7	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2021	2.025	1
8	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2023	2.012	1
9	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2021	14.2~	1
10	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2023	15.9~	1
11	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2021	21.3~	1
12	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2023	10.5~	1
13	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	CALI~	2021	8.6	1
14	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	CALI~	2023	9.4	1
15	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	FLOR~	2021	10.5	1



16	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	FLOR~	2023	5.2	1
17	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~ CALI~	2021	45	1	
18	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~ CALI~	2023	88	1	
19	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~ FLOR~	2021	61	1	
20	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~ FLOR~	2023	96	1	

# This table directly shows the transition of captan use from year 2021 to 2023 in each state

#Then we can make a bar chart to compare the captan use in California and Florida

#Generate Data from 2021 and 2023

```
captan_2021 <- captan[str_detect(captan$Year, regex("2021", ignore_case = TRUE)), ]
```

```
captan_2023 <- captan[str_detect(captan$Year, regex("2023", ignore_case = TRUE)), ]
```

#Ensure the value under Value column is numeric, so in graph it can provide the correct relationship

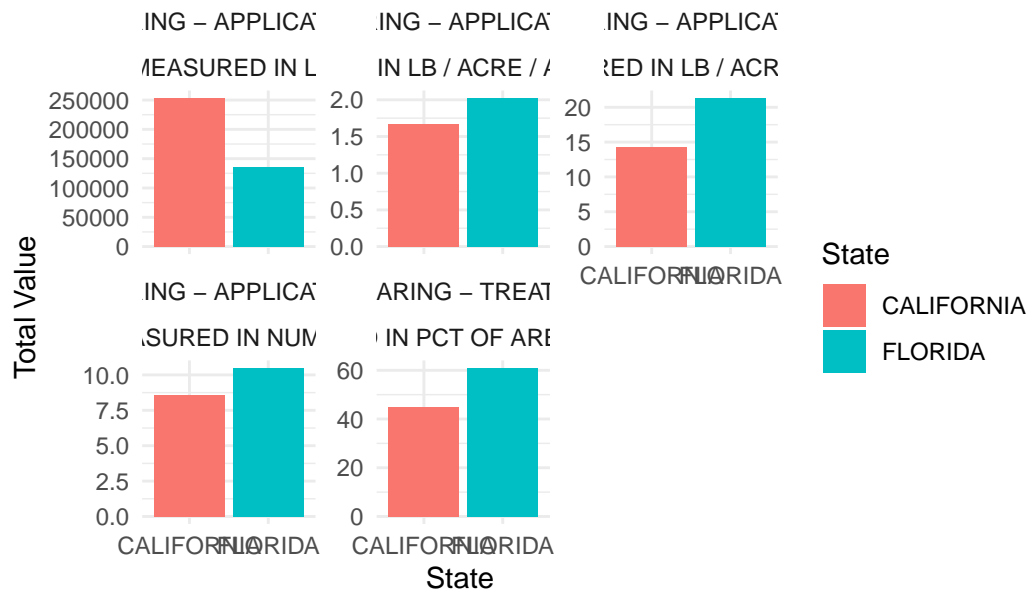
```
captan_2021$Value <- as.numeric(gsub(",", "", captan_2021$Value))
```

```
captan_2023$Value <- as.numeric(gsub(",", "", captan_2023$Value))
```

#Then we can make the graph (code perfection by chatgpt, for a better visual effect)

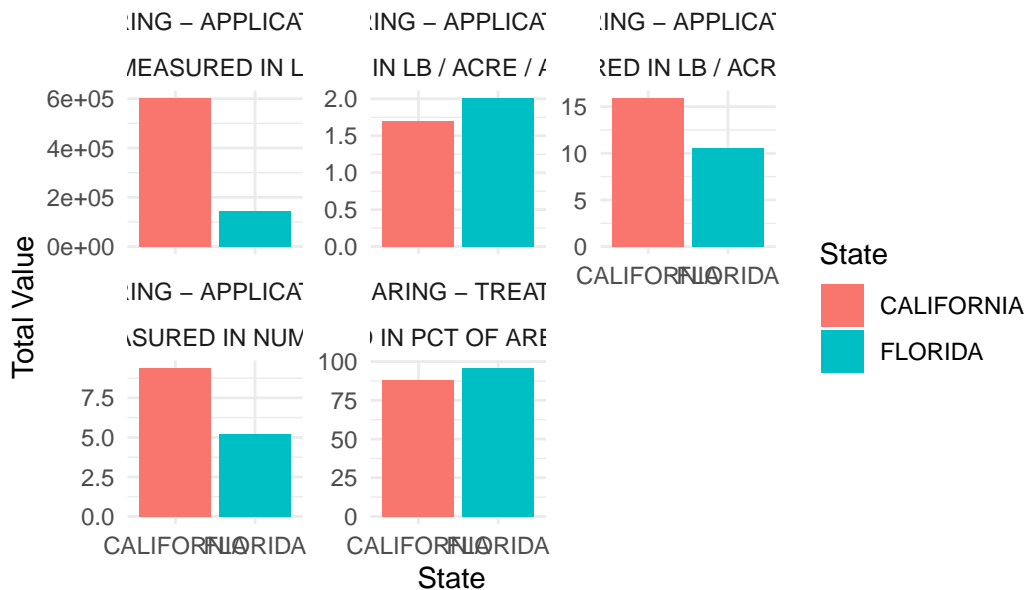
```
ggplot(captan_2021, aes(x = State, y = Value, fill = State)) +
  geom_bar(stat = "identity", position = "dodge") +
  facet_wrap(~ Category + Item, scales = "free_y") +
  labs(title = "Value of Captan Measurement Comparison by State (2021)",
       x = "State",
       y = "Total Value") +
  theme_minimal()
```

## Value of Captan Measurement Comparison by State (2021)



```
ggplot(captan_2023, aes(x = State, y = Value, fill = State)) +
  geom_bar(stat = "identity", position = "dodge") +
  facet_wrap(~ Category + Item, scales = "free_y") +
  labs(title = "Value of Captan Measurement Comparison by State (2023)",
       x = "State",
       y = "Total Value") +
  theme_minimal()
```

## Value of Captan Measurement Comparison by State (2023)



```
#Remove intermediate Data set
rm(captan_2021,captan_2023)
```

From the graph of 2021, California uses the fungicide that contains captan (a certified carcinogenic chemical) more than Florida in general. However, when we look up into other measurement, the average detection of captan use in California is less than Florida. This is because California Strawberry Production Firm is larger than Florida in size.

In 2023, There are some changes in the data. The measure in lb/acre/year and measure in number of California firm is higher than Florida firm, meaning that California uses captan fungicide more often per year, and the average number that detected captan is higher than Florida.

## Chemical 2: Thiram

```
#Follows the same step as captan
#Filter rows form chemical_CA
thiram_CA <- chemical_CA[str_detect(chemical_CA$`Domain Category`, regex("thiram", ignore_case = TRUE))

# Filter rows from chemical_FL
thiram_FL <- chemical_FL[str_detect(chemical_FL$`Domain Category`, regex("thiram", ignore_case = TRUE))
```

```
#Get the thiram information data and gather them into one data set
thiram_combined <- rbind(thiram_CA, thiram_FL)

#remove intermediate data set
rm(thiram_CA)
rm(thiram_FL)

#Table of Comparison of different year by same state
thiram <- thiram_combined |> group_by(Category, Item, State, Year) |> count(Value)

print(thiram)
```

```
# A tibble: 20 x 6
```

```
# Groups:   Category, Item, State, Year [20]
```

	Category <chr>	Item <chr>	State <chr>	Year <dbl>	Value <chr>	n <int>
1	" BEARING - APPLICATIONS"	" MEASURED IN LB"	CALI~	2021	96,3~	1
2	" BEARING - APPLICATIONS"	" MEASURED IN LB"	CALI~	2023	269,~	1
3	" BEARING - APPLICATIONS"	" MEASURED IN LB"	FLOR~	2021	142,~	1
4	" BEARING - APPLICATIONS"	" MEASURED IN LB"	FLOR~	2023	112,~	1
5	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2021	2.144	1
6	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2023	2.201	1
7	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2021	2.38	1
8	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2023	2.156	1
9	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2021	5.029	1
10	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2023	8.873	1
11	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2021	22.5~	1
12	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2023	12.4~	1
13	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	CALI~	2021	2.3	1
14	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	CALI~	2023	4	1
15	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	FLOR~	2021	9.5	1
16	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	FLOR~	2023	5.8	1
17	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	CALI~	2021	49	1
18	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	CALI~	2023	70	1
19	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	FLOR~	2021	61	1
20	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	FLOR~	2023	63	1

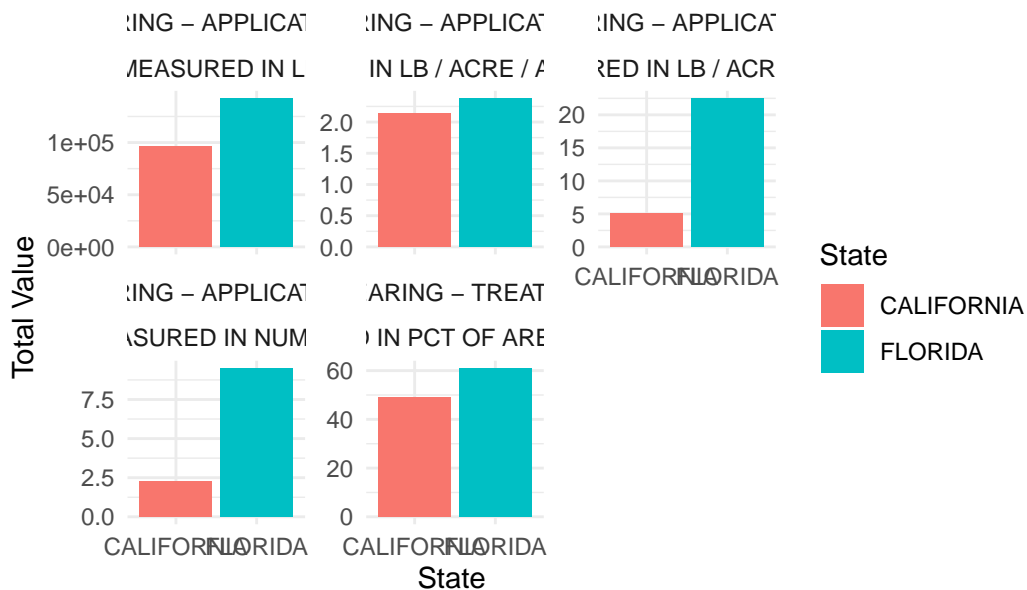
```
#Rearrange the data for bar chart
```

```
thiram_2021 <- thiram[str_detect(thiram$Year, regex("2021", ignore_case = TRUE)), ]
thiram_2023 <- thiram[str_detect(thiram$Year, regex("2023", ignore_case = TRUE)), ]
```

```
#Ensure the value under Value column is numeric, so in graph it can provide the right relation
thiram_2021$Value <- as.numeric(gsub(",", "", thiram_2021$Value))
thiram_2023$Value <- as.numeric(gsub(",", "", thiram_2023$Value))
#Same year, same domain category and item, but different state record comparison (code perfect)

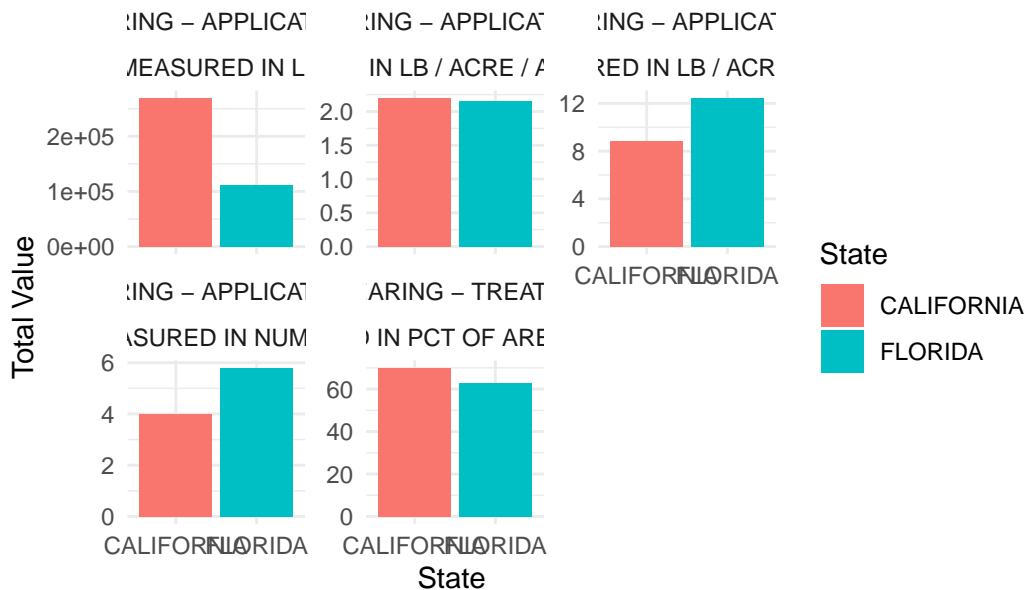
ggplot(thiram_2021, aes(x = State, y = Value, fill = State)) +
  geom_bar(stat = "identity", position = "dodge") +
  facet_wrap(~ Category + Item, scales = "free_y") +
  labs(title = "Value of Thiram Measurement Comparison by State (2021)",
       x = "State",
       y = "Total Value") +
  theme_minimal()
```

Value of Thiram Measurement Comparison by State (2021)



```
ggplot(thiram_2023, aes(x = State, y = Value, fill = State)) +
  geom_bar(stat = "identity", position = "dodge") +
  facet_wrap(~ Category + Item, scales = "free_y") +
  labs(title = "Value of Thiram Measurement Comparison by State (2023)",
       x = "State",
       y = "Total Value") +
  theme_minimal()
```

## Value of Thiram Measurement Comparison by State (2023)



```
# Remove intermediate data set
rm(thiram_2021, thiram_2023)
```

## Chemical 3: Propiconazole

```
#Follows the same step as captan
#Filter rows from chemical_CA
ppz_CA <- chemical_CA[str_detect(chemical_CA$`Domain Category`, regex("thiram", ignore_case = TRUE))]

# Filter rows from chemical_FL
ppz_FL <- chemical_FL[str_detect(chemical_FL$`Domain Category`, regex("thiram", ignore_case = TRUE))]

#Get the thiram information data and gather them into one data set
ppz_combined <- rbind(ppz_CA, ppz_FL)

#remove intermediate data set
rm(ppz_CA)
rm(ppz_FL)

#Table of Comparison of different year by same state
ppz1 <- ppz_combined |> group_by(Category, Item, State, Year) |> count(Value)
```

```
print(ppz1)
```

```
# A tibble: 20 x 6
```

```
# Groups:   Category, Item, State, Year [20]
```

	Category	Item	State	Year	Value	n
	<chr>	<chr>	<chr>	<dbl>	<chr>	<int>
1	" BEARING - APPLICATIONS"	" MEASURED IN LB"	CALI~	2021	96,3~	1
2	" BEARING - APPLICATIONS"	" MEASURED IN LB"	CALI~	2023	269,~	1
3	" BEARING - APPLICATIONS"	" MEASURED IN LB"	FLOR~	2021	142,~	1
4	" BEARING - APPLICATIONS"	" MEASURED IN LB"	FLOR~	2023	112,~	1
5	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2021	2.144	1
6	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2023	2.201	1
7	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2021	2.38	1
8	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2023	2.156	1
9	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2021	5.029	1
10	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	CALI~	2023	8.873	1
11	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2021	22.5~	1
12	" BEARING - APPLICATIONS"	" MEASURED IN LB / ACRE / ~	FLOR~	2023	12.4~	1
13	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	CALI~	2021	2.3	1
14	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	CALI~	2023	4	1
15	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	FLOR~	2021	9.5	1
16	" BEARING - APPLICATIONS"	" MEASURED IN NUMBER"	FLOR~	2023	5.8	1
17	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	CALI~	2021	49	1
18	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	CALI~	2023	70	1
19	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	FLOR~	2021	61	1
20	" BEARING - TREATED"	" MEASURED IN PCT OF AREA ~	FLOR~	2023	63	1

```
#Rearrange the data for bar chart
```

```
ppz_2021 <- ppz1[str_detect(ppz1$Year, regex("2021", ignore_case = TRUE)), ]
```

```
ppz_2023 <- ppz1[str_detect(ppz1$Year, regex("2023", ignore_case = TRUE)), ]
```

```
#Ensure the value under Value column is numeric, so in graph it can provide the right relation
```

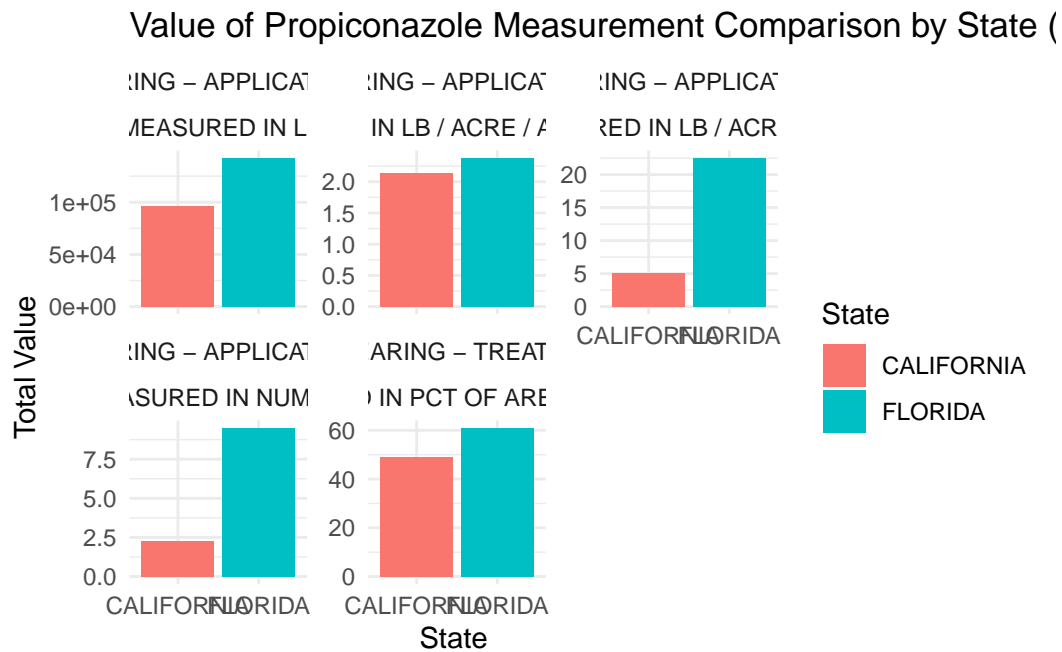
```
ppz_2021$Value <- as.numeric(gsub(",", "", ppz_2021$Value))
```

```
ppz_2023$Value <- as.numeric(gsub(",", "", ppz_2023$Value))
```

```
#Same year, same domain category and item, but different state record comparison (code perfect)
```

```
ggplot(ppz_2021, aes(x = State, y = Value, fill = State)) +
  geom_bar(stat = "identity", position = "dodge") +
  facet_wrap(~ Category + Item, scales = "free_y") +
  labs(title = "Value of Propiconazole Measurement Comparison by State (2021)",
       x = "State",
```

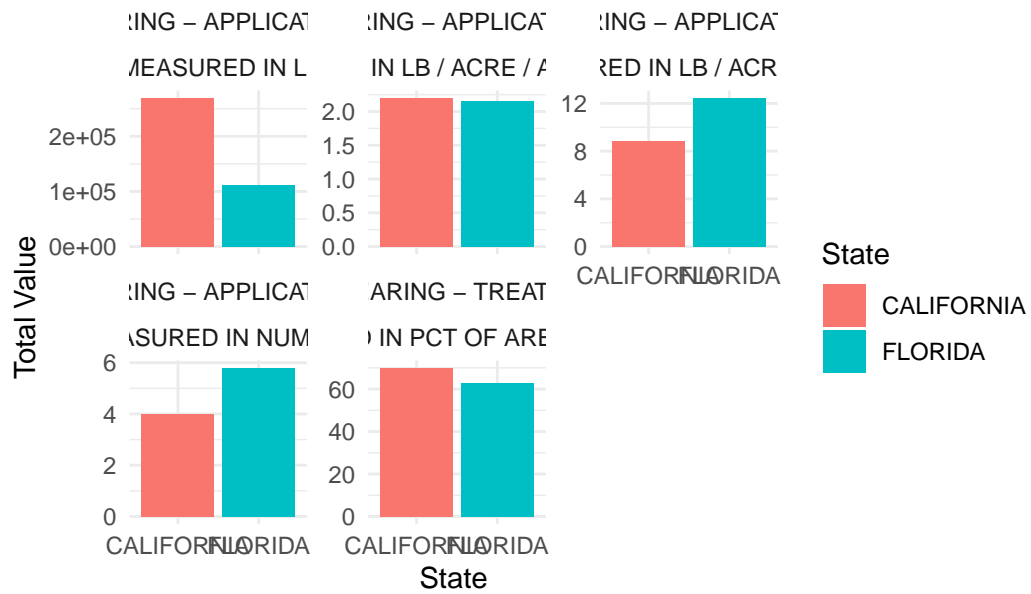
```
y = "Total Value") +
theme_minimal()
```



```
ggplot(ppz_2023, aes(x = State, y = Value, fill = State)) +
  geom_bar(stat = "identity", position = "dodge") +
  facet_wrap(~ Category + Item, scales = "free_y") +
  labs(title = "Value of Propiconazole Measurement Comparison by State (2023)",
        x = "State",
        y = "Total Value") +
  theme_minimal()
```



## Value of Propiconazole Measurement Comparison by State (



```
# Remove intermediate data set
rm(ppz_2021, ppz_2023)
```