## **Avocado Prices Prediction**

#### Approach:

#### a. Extracting data from a large Dataset

**Data Cleansing:** Data cleansing is the process of preparing data for analysis by removing or modifying data that is incorrect, incomplete, irrelevant, duplicated, or improperly formatted. This data is usually not necessary or helpful when it comes to analysing data because it may hinder the process or provide inaccurate results. This process identifies incomplete, incorrect, inaccurate or irrelevant parts of the data and then replacing modifying, or deleting the dirty or coarse data. Data cleansing may be performed interactively with data wrangling tools, or as batch processing through scripting.

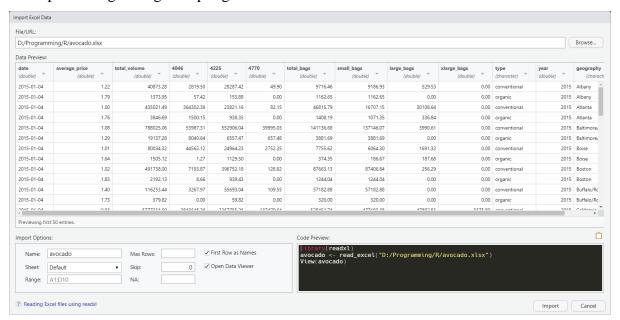


Fig 1.1: Importing dataset in RStudio.

```
| State | Stat
```

**Fig 1.2:** Glimpse of the data.

Fig 1.3: Summary of the attributes.

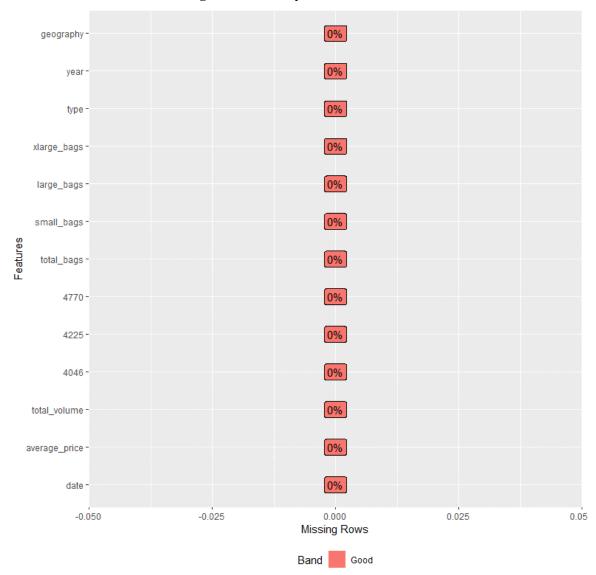
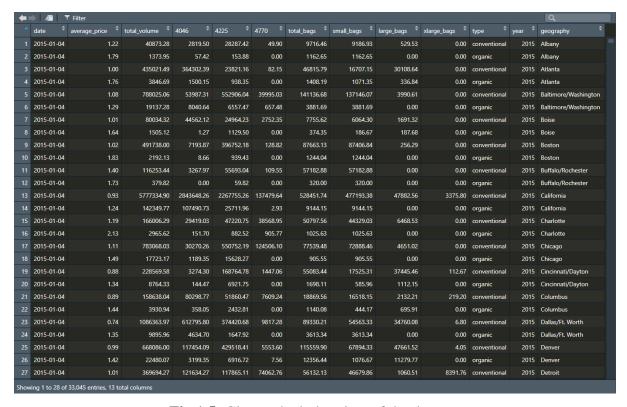


Fig 1.4: Missing values information using plot\_missing() method.



**Fig 1.5:** Cleansed tabular view of the dataset.

## Counting missing values in the dataset:

```
> sum(is.na(data))
[1] 0
```

#### b. Exploratory Analysis

In this phase of the project, we will concentrate mainly on the following aspect of the time series forecasting analysis which consists of:

**Seasonal Patterns:** In this section we will focus on constant patterns that occur frequently from year to year and from month to month in both types of avocados conventional and organic.

```
library(dplyr)
library(ggplot2)
library(DataExplorer)

options(repr.plot.width = 8, repr.plot.height = 4)
ggplot(
    data,
    aes(x = average_price, fill = type)
) +
geom_density() +
facet_wrap(~type) +
theme_minimal() +
theme(
    plot.title = element text(hjust = 0.5),
```

```
legend.position = "bottom"
) +
labs(title="Avocado Price by Type") +
scale_fill_brewer(palette = "Set2")
```

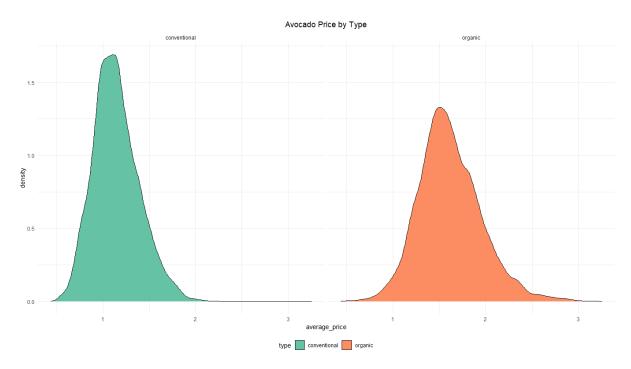


Fig 2.1: Density plots of the different type of avocadoes.

## **Output:**

#### **Types of Avocados:**

In this section we will analyze the different types of avocados that we have in this dataset. Basically, we have two types of avocados: **Conventional and Organic.** 

## **Summary:**

 Organic avocados: Based on the price changes throughout time we can see that they are more expensive. • **Conventional avocados:** Based on price changes throughout time we can see that they are less expensive.

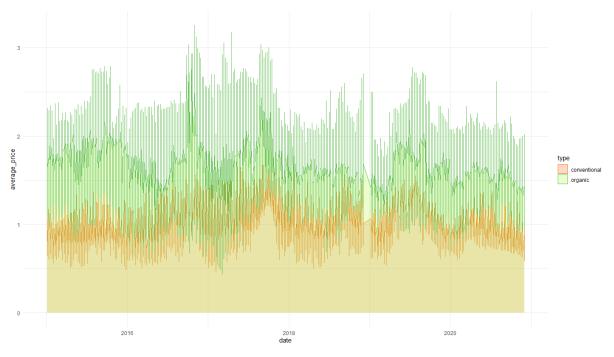


Fig 2.2: Range of average prices of avocados throughout the years.

# **Univariate Analysis:**

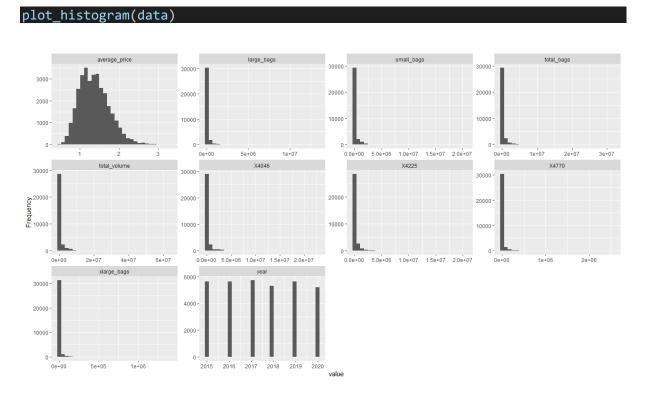


Fig 2.3: Histogram for every continuous feature.

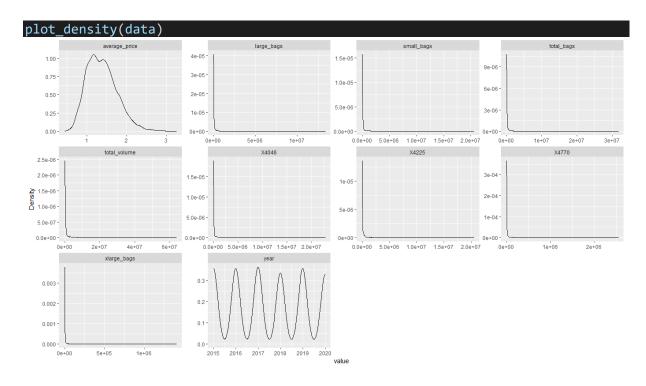


Fig 2.4: Density comparison plot.

The data is provided from 2015 till 2020. Next is to convert the year column to factor to treat it as categorical variable and also create a new column called month from date.

```
avocado$year = as.factor(avocado$year)
avocado$date = as.Date(avocado$date)
avocado$month = factor(months(avocado$date), levels = month.name)
```

Trend of Avocado Prices: The trend of avocado prices in last six years by avocado type.

```
options(repr.plot.width = 7, repr.plot.height = 5)
ggplot(avocado, aes(avocado$type, avocado$average_price)) +
geom_boxplot(aes(colour = avocado$year)) +
labs(
    colour = "Year",
    x = "Type",
    y ="Average Price",
    title = "Boxplot - Average price per year by type."
)
```

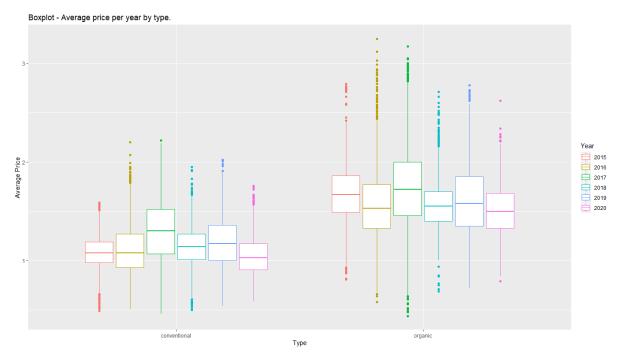


Fig 2.5: Average price of avocados per year by type of avocados.

## Monthly trend of avocado prices by Avocado type in last years

**Step 1:** Group the dataset by Year, Month and Avocado Type. Calculate the monthly average price for Avocado respectively for each year.

```
grouped = avocado %>%
   group_by(year, month, type) %>%
   select(year, month, type, average_price) %>%
   summarise(average_price = mean(average_price))
```

**Step 2:** Plot a line chart showing the trend for both Conventional and Organic Avocado.

```
options(repr.plot.width = 12, repr.plot.height = 5)
ggplot(data = grouped, aes(x = month, y = average_price, colour = year, group
= year)) +
labs(
    colour = "Year",
    x = "Month",
    y ="Average Price",
    title = "Line Plot - Average monthly prices by type for every year."
) +
geom_line() +
theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
facet_grid(. ~grouped$type)
```

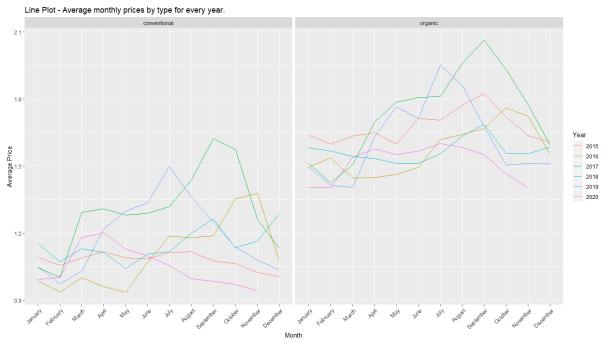


Fig 2.6: Average price of avocados per month by type of avocados for every year.

#### Group by geography:

```
grouped_geography_conv = avocado %>%
    select(year, geography, type, average_price) %>%
    filter(type == 'conventional')
min_con = round(min(grouped_geography_conv$average_price), 1) - 0.1
max_con = round(max(grouped_geography_conv$average_price), 1) + 0.1
grouped_geography_org = avocado %>%
    select(year, geography, type, average_price) %>%
    filter(type == 'organic')
min_org = round(min(grouped_geography_org$average_price), 1) - 0.1
max_org = round(max(grouped_geography_org$average_price), 1) + 0.1
options(repr.plot.width = 10, repr.plot.height = 12)
ggplot(grouped_geography_conv, aes(x = geography, y = average_price)) +
geom_tufteboxplot() +
facet_grid(.~grouped_geography_conv$year, scales = "free") +
labs(
    colour = "Year",
    x = "Geography",
    y ="Average Price",
    title = "Average prices of Conventional Avocados for each geography by yea
scale_y_continuous(breaks = c(seq(min_con, max_con, 0.2)), limits = c(min_con,
 max_con)) +
```

```
coord_flip() +
theme(axis.text.x = element text(angle = 90, vjust = 0))
```

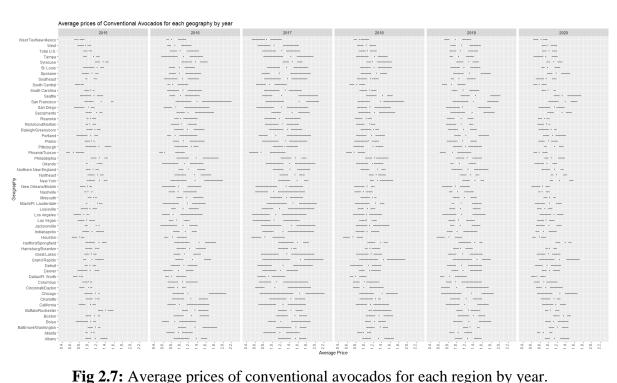


Fig 2.7: Average prices of conventional avocados for each region by year.

```
options(repr.plot.width = 12, repr.plot.height = 12)
ggplot(grouped_geography_org, aes(x = geography, y = average_price)) +
geom_tufteboxplot() +
facet_grid(.~grouped_geography_org$year, scales = "free") +
labs(
   colour = "Year",
   x = "Geography",
   y ="Average Price",
   title = "Average prices of Organic Avocados for each geography by year"
scale_y_continuous(breaks = c(seq(min_org, max_org, 0.2)), limits = c(min_org,
max_org)) +
coord_flip() +
theme(axis.text.x = element_text(angle = 90, vjust = 0))
```

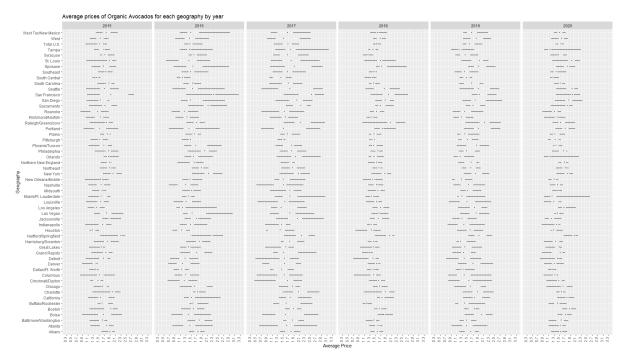


Fig 2.8: Average prices of organic avocados for each region by year.

## **Multivariate Analysis:**





Fig 2.9: Correlation heatmap for all the continuous categories.

#### **Principal Component Analysis**

```
data.pca <- prcomp(
    select(data, 2, 3, 4, 5, 6, 7, 8, 9, 10),
    center = TRUE,
    scale = TRUE
)
print(data.pca)
summary(data.pca)</pre>
```

## c. Applying DMBI/ML algorithms

#### 1. Linear Regression

Linear regression is one of the most commonly used predictive modelling techniques to predict the value of a continuous variable Y based on one or more input predictor variables X. The aim is to establish a mathematical formula between the response variable (Y) and the predictor variables (X's). Linear regression model can be used to learn which features are important by examining coefficients. If a coefficient is close to zero, the corresponding feature is considered to be less important than if the coefficient was a large positive or negative value. That's how the linear regression model generates the output. Coefficients are multiplied with corresponding input variables, and in the end, the bias (intercept) term is added.

**Step 1:** It is observed in the dataset that the price of avocados goes down as volume sold goes up. Test if this relationship is statistically significant.

```
library(sjPlot)
library(ggfortify)
library(readxl)

# Import the dataset.
avocado <- read_excel("avocado.xlsx")
data = avocado

conmod <- data %>%
```

```
filter(type == "conventional") %>%
  mutate(Volume = log(`total_volume`)) %>%
  lm(average_price ~ Volume, data = .)

orgmod <- data %>%
  filter(type == "organic") %>%
  mutate(Volume = log(`total_volume`)) %>%
  lm(average_price ~ Volume, data = .)
```

Save the model in order to display the results in a table. Use the tab\_model function from the siPlot package for this purpose.

```
# Printing the output in a table rather than in the console.

tab_model(conmod, orgmod)
```

#### **Step 2: Regression Diagnostics**

Examine if the fitted model is any good with autoplot and ggfortify. This will give a ggplot2 version of the regression diagnostics plot from base R.

```
#Regression diagnostics.
autoplot(conmod)
autoplot(orgmod)
```

#### 2. Decision Trees

Decision tree is a type of supervised learning algorithm that can be used in both regression and classification problems. It works for both categorical and continuous input and output variables. It builds classification models in the form of a tree-like structure, just like its name. It is also used to create data models that will predict class labels or values for the decision-making process. The models are built from the training dataset fed to the system (supervised learning). Using a decision tree, we can visualize the decisions that make it easy to understand and thus it is a popular data mining technique.

```
library(rsample)  # Data splitting.
library(dplyr)  # Data wrangling.
library(rpart)  # Performing regression trees.
library(rpart.plot)  # Plotting regression trees.
library(readxl)

# Import the dataset.
avocado <- read_excel("avocado.xlsx")

# Drop the geography column.
data = avocado[, 1:12]</pre>
```

```
# Splitting the dataset in a 0.7 ratio by default order by years.
avocado train = data[1:23131,]
avocado_test = data[23132:33045,]
# Regressor.
m1 <- rpart(formula = average_price ~ ., data = avocado_train, method = "anova</pre>
print(m1)
# Summary of the decision tree regressor.
summary(m1)
# Plotting the tree.
rpart.plot(m1)
plotcp(m1)
# Predicting prices for the test split.
predictions <- predict(m1, avocado_test, type = 'vector')</pre>
# Summarizing accuracy.
# Calculating the Root Mean Squared Error.
RMSE(predictions, avocado_test$average_price)
# Calculating the Mean Squared Error.
mse <- mean((avocado_test$average_price - predictions)^2)</pre>
print(mse)
# Calculating the Mean Absolute Error.
MAE = function(actual, predicted) {
  mean(abs(actual - predicted))
print(MAE(avocado_test$average_price, predictions))
```

#### d. Visualization and Interpretation of results.

### 1. Linear Regression

	average_price			average_price		
Predictors	Estimates	CI	p	Estimates	CI	р
(Intercept)	1.66	1.62 - 1.70	< 0.001	1.89	1.86 - 1.93	< 0.001
Volume	-0.04	-0.040.04	<0.001	-0.03	-0.030.02	<0.001
Observations	16524			16521		
$R^2 / R^2$ adjusted	0.043 / 0.043			0.014 / 0.014		

Fig 4.1: Tabular data showing the relation between average avocado prices and volume.

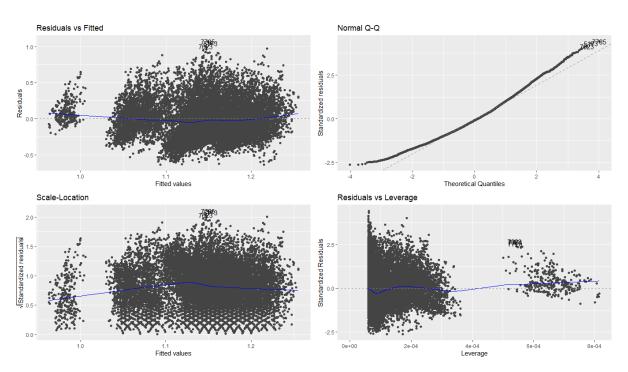


Fig 4.2: Regression diagnostic plots for the conventional avocados.

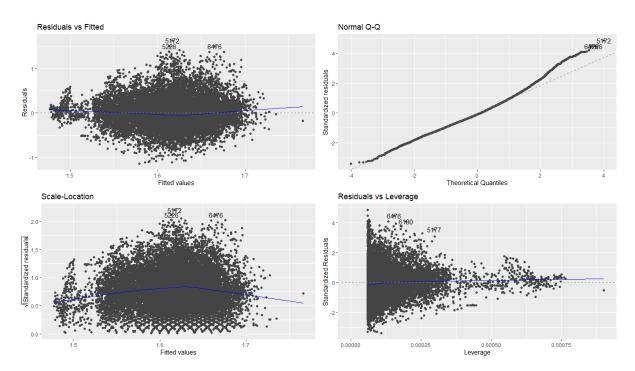


Fig 4.3: Regression diagnostic plots for the organic avocados

## 2. Decision Tree Regressor

Fig 4.4: Decision tree regressor model.

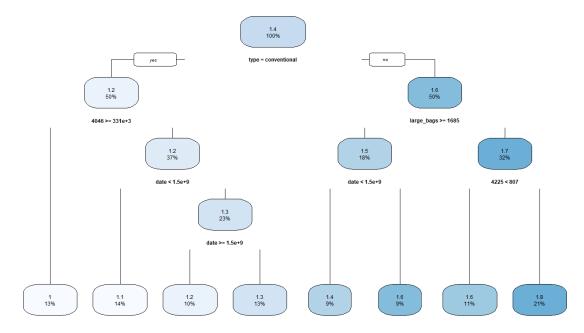
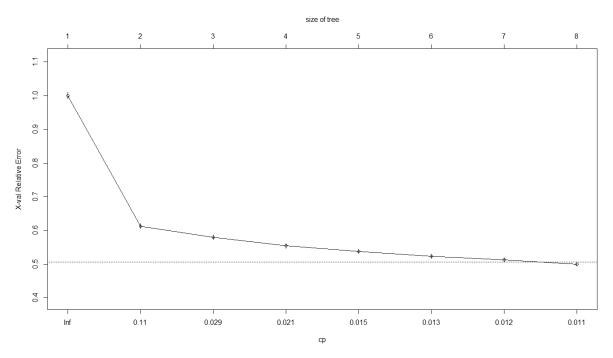


Fig 4.5: Visualizing the decision tree regressor model.



**Fig 4.6:** Visual representation of the cross-validation results in the model.

```
> # Predicting prices for the test split.
> predictions <- predict(m1, avocado_test, type = 'vector')
> # Summarizing accuracy.
> # Calculating the Root Mean Squared Error.
> RMSE[predictions, avocado_test$average_price)
[1] 0.2758131
> # Calculating the Mean Squared Error.
> mse <- mean((avocado_test$average_price - predictions)^2)
> print(mse)
[1] 0.07607286
> # Calculating the Mean Absolute Error.
> MAE = function(actual, predicted) {
+ mean(abs(actual - predicted))
+ }
> print(MAE(avocado_test$average_price, predictions))
[1] 0.2203961
```

Fig 4.7: Summarizing accuracy with RMSE, MSE and MAE.

#### e. Results and Discussions

#### 1. Linear Regression

Observations claiming the relationship between avocado prices and volumes where prices go down with increase in volumes sold were confirmed by the results. The p-value is shows that the negative coefficient is statistically significant.

It is observed from the Linear Regression plots, for both type of avocados (conventional and organic), that the model roughly follows a normal distribution as the deviations are not too steep. Furthermore, the standard deviation of the residuals does not exceed 3 and are mostly below 2. We can therefore conclude that the price of avocado drops as volume increases.

#### 2. Decision Tree Regressor

A decision tree is one of the most common classification algorithms that are implemented. The avocado dataset was first split in the ratio 0.7 for training and testing sets. The decision tree implemented in this project was plot on the training dataset. From the testing set, predictions were calculated. To summarize the accuracy of the model, different errors like Root Mean Squared Error, Mean Squared Error, Mean Absolute Error were calculated. MSE resulted in a 0.076 which is much closer to 0, indicating that the predictions were significantly accurate and closer to the actual values in the testing set. Lower values of MSE indicate good predictions, the closer the MSE is to 0, the accurate the prediction is.

<u>Conclusion</u>: The two algorithms that were used to create models were Linear Regression and Decision Tree Regressor.

From the Linear Regression plots, for both type of avocados (conventional and organic), we conclude that the model roughly follows a normal distribution as the deviations are not too steep (less than 2.5). We therefore conclude that the price of avocado drops as volume increases.

Decision Tree Regression model was implemented to predict the average prices of the avocados. Different errors were calculated to test the accuracy. They are:

- Root Mean Squared Error = 0.2758
- Mean Squared Error = 0.0760
- Mean Absolute Error = 0.2209

Thus, we can use R in order to analyse and design various algorithms on any dataset. Other functions can also be performed using R such as exploratory data analysis and functions in preprocessing of data like cleaning data and replacing missing values. Larger datasets can also be modified to work on smaller subsets of data to avoid problems like overfitting. Using R for carrying out such operations is very beneficial.