The Prediction of case’s recovery rate Corona Virus 2019 (COVID-19) worldwide

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faculty of technology and computer science

UNIVERSITY OF MALAYA

Kuala lumpur

2021

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**UNIVERSITY OF MALAYA**

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The prediction of cases based on average moving days of covid-19 worlwide  
Abstract

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Around 96,000 confirmed coronavirus disease cases have been reported in 2019 (COVID-2019) and 3300 reported deaths to date of 05/03/2020. Inhalation or contact with infected droplets transmits the disease and the incubation duration varies from two (2) to fourteen (14) days. This research then aimed to predict the Recovery rate and fatality rate from the confirm, deaths, recovery and active results by using Linear Regression, Random Forest and Decision Tree model to evaluate the prediction of the result. The Clustering method also be applied by clustering the cases based on the high, middle and low cases within days.

Acknowledgements

I would like to express my very great appreciation to Dr Woo Chaw Seng for his valuable and constructive teaching during the pandemic when the class need to be handle online. The struggle to make the class for the entire semester to be great and the knowledge received by student perfectly. His passion to conduct the class and the project for the course was really appreciated.

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# introduction

The discovery and spread of the novel coronavirus virus in 2019 challenge the planet with a new public health crisis (2019-nCoV) or also known as Extreme Acute Respiratory Syndrome (SARS-CoV-2). The virus originated in bats and was transmitted to humans in Wuhan, Hubei province, China in December 2019 through still unknown intermediate animals. Around 96,000 confirmed coronavirus disease cases have been reported in 2019 (COVID-2019) and 3300 reported deaths to date of 05/03/2020. Inhalation or contact with infected droplets transmits the disease and the incubation duration varies from two (2) to fourteen (14) days. Usually, the symptoms are fever, cough, sore throat, breathlessness, weakness, malaise and more. In most individuals, the disease is mild; pneumonia, acute respiratory distress syndrome (ARDS) and multi-organ dysfunction can progress in some, usually the elderly and those with comorbidities. The case fatality rate is estimated to range between 2% and 3%.

The COVID-19 case description is based on symptoms irrespective of travel history or interaction with reported cases. In patients with a recent, continuous cough, fever or loss or a changed sense of normal smell or taste, diagnosis is suspected (anosmia). A diagnostic test has been developed, and suspected cases are quarantined by countries. A critical source of information and expertise has been generated by this sudden burst of cases and their health data. Using various data storage systems, there is an immediate need to store such a vast volume of data in these situations. These data are used for the purpose of research and development on the virus, the pandemic, and initiatives to contain the virus and its aftermath.

## Problem Statement

New cases of COVID-19 (Coronavirus) are rising rapidly at staggering rates worldwide; more than 1.2 billion people have acquired an infection and about 65,000 have died of the disease to date. Every day, new coronavirus outbreaks are announced, related statistics fluctuate quickly, and huge amounts of data are produced at great speed, becoming a challenge for academics and professionals to convert this information into useful information. People around the world are still unaware with the growth rate of the new cases every day and take it slightly due to lack of understanding the given information such as the causes and future forecasting regarding on result collected.

## Background of Study

As the latest coronavirus (COVID-19) keeps the earth in limbo, it has become an urgent, significant challenge for mankind to end it as soon as possible. Every day, new coronavirus outbreaks are announced, related statistics fluctuate quickly, and huge amounts of data are produced. Big data may potentially help us understand the existence of the new coronavirus, offering a source of preventive and treatment-inspiring knowledge. Big data analysis may also promote the prediction of systemic changes that may arise from the current pandemic in our culture, economy, and lifestyle.

The dataset found in open website, Kaggle.com are provided with different datasets provide Global confirm cases, Global death cases and Global Recover cases. All datasets contains the features such as the list of country with longitude and latitude and the list of dates for every cases until 9th January 2021. The all 3 collection of datasets have 272 of rows and 358 columns.

### Objectives

The objectives are listed below to achieve the goal of this project:

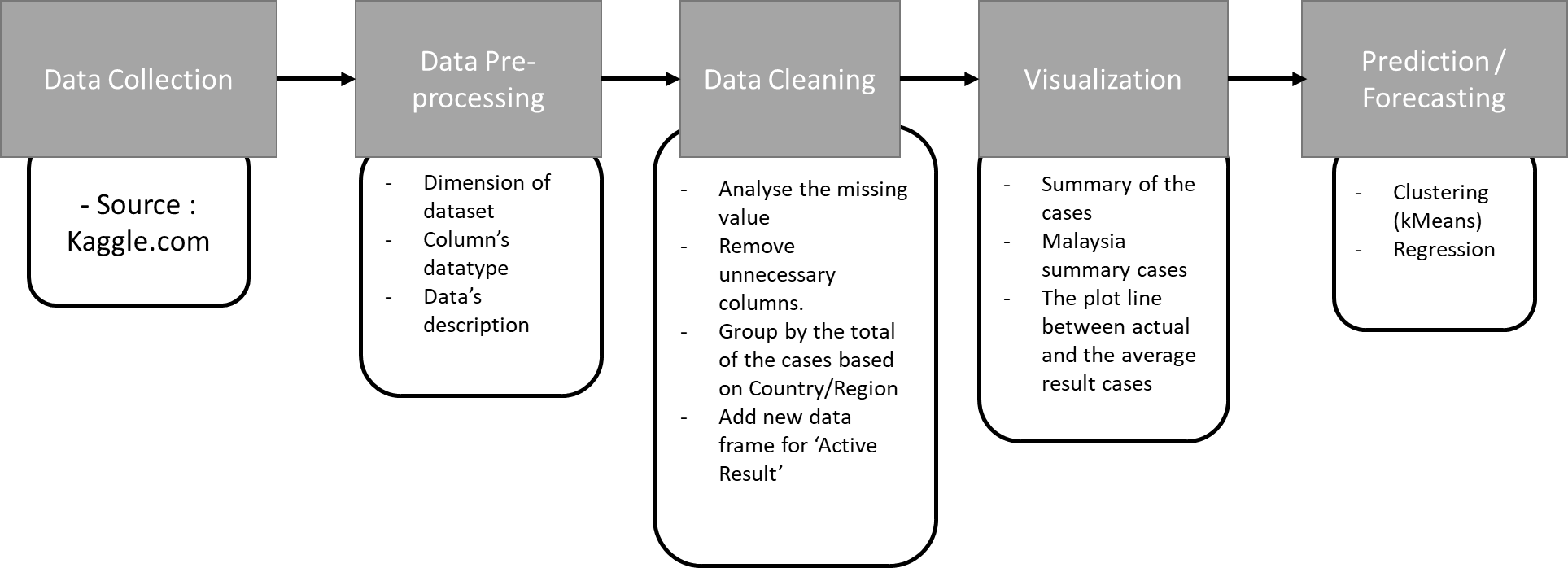
1. To improve understanding of COVID19 information through processing and visualization of massive and structured data.

2. To trace the trends and growth rate related to deaths and recovery cases expected during pandemic COVID19 around the world.

3. To predict the growth rate within the time related to data from confirm, deaths, recovery and active cases around the world.

# methdology

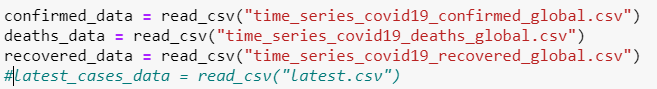
In recent years, with promising results, predictive medical analysis using machine learning techniques has experienced enormous growth. In numerous types of applications in various fields, machine learning algorithms are effectively applied. The study was carried out in several phases.



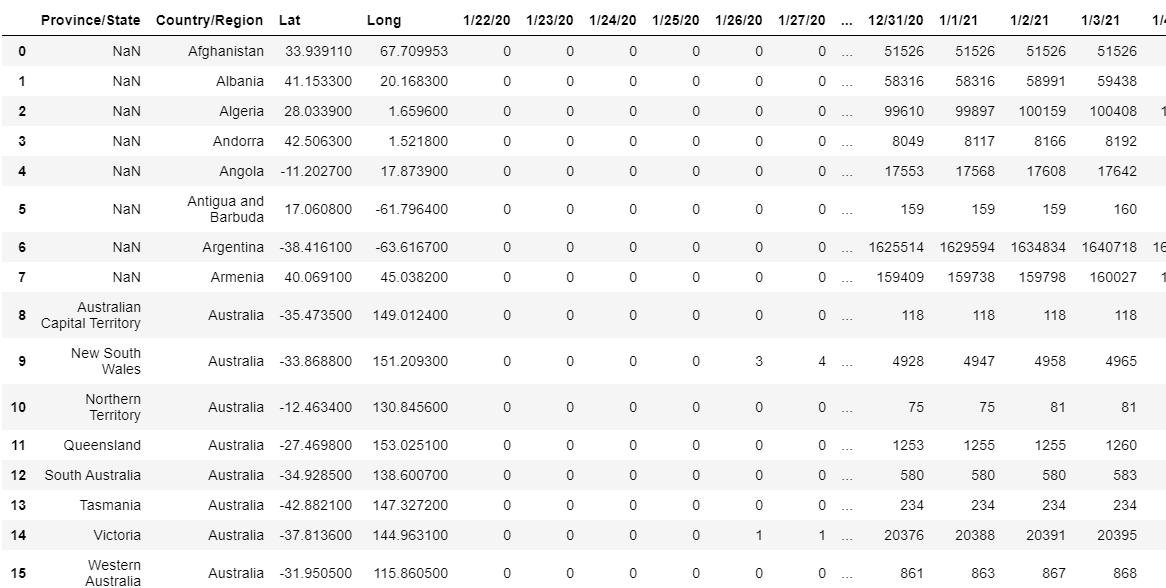
***Figure 2.0:*** *Flowchart of the basic data analysis methodology*

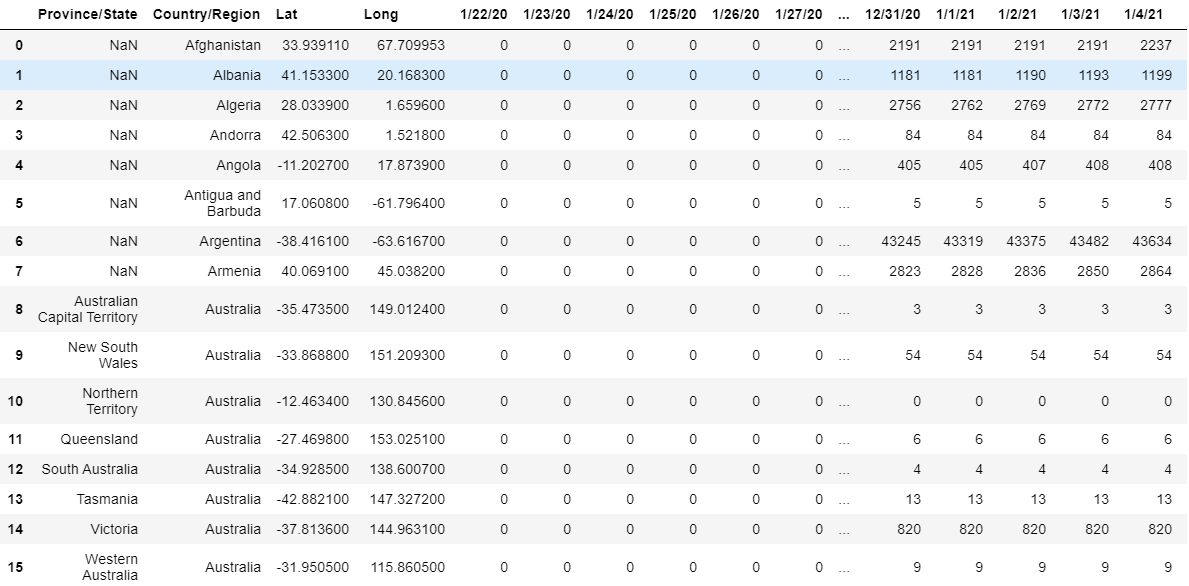
## Data Collection

For this project, the COVID-19 dataset from the Kaggle is taken for predictive analysis. There are 3 different datasets used in this project named as Global Confirmed Cases, Global Deaths Cases and Global Recovery Cases. All the datasets have the same dimension which is 272 of rows and 358 of columns before cleaning process. The features provided in the dataset is the province, country with its longitude and latitude and list of dates up until 9th January 2021.



***Figure 2.1.1:*** *query in python reading the datasets (csv file)*



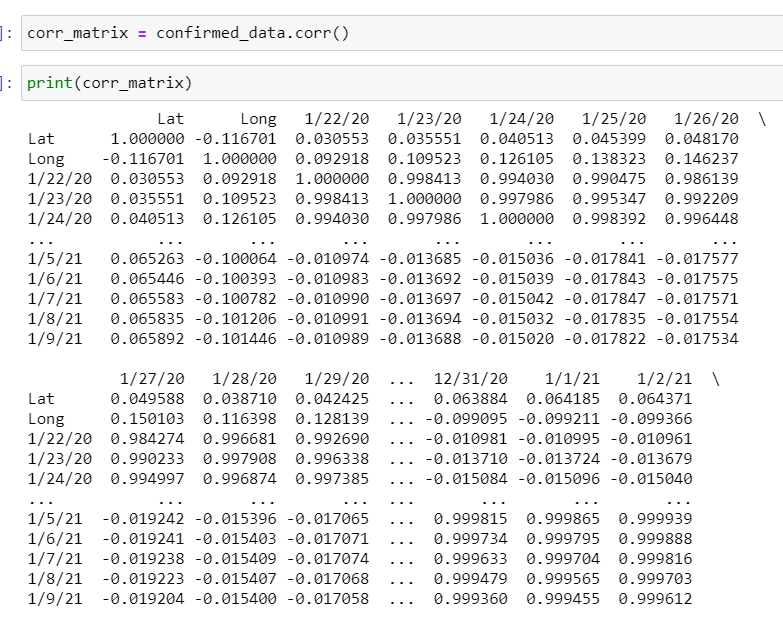
***Figure 2.1.2:*** *A glance of Global Confirmed Cases Dataset*

***Figure 2.1.3:*** *A glance of Global Deaths Cases Dataset*



***Figure 2.1.4:*** *A glance of Global Recovered Cases Dataset*

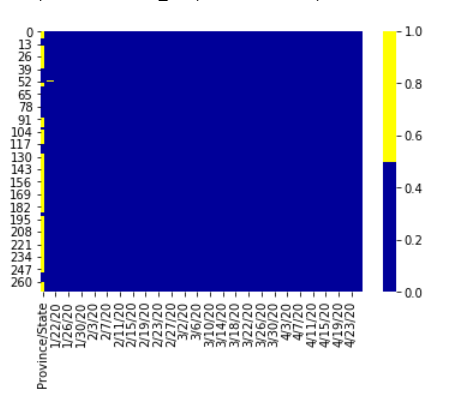
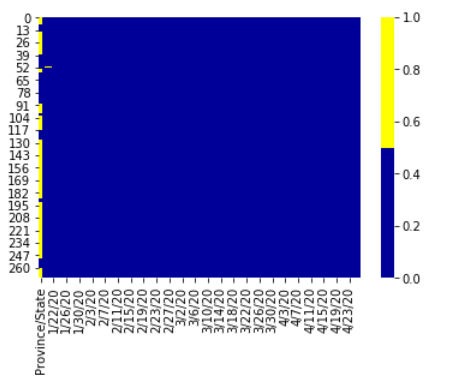
## Data Pre-processing

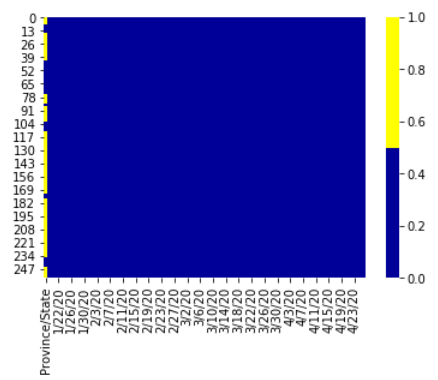
Using data preprocessing and data cleaning methodologies, the considered dataset was cleaned, then the resulting dataset was considered for multiple experiments using various classification algorithms. The datasets contain the information of the cases within the dates. With a glance through datasets, the datasets have the dimension of 272 of rows and 358 of columns before cleaning. This pre-processing part is the way to determine the data type of every columns, the description summary of the datasets, came out with the unique keys of every strings (object) columns and determine the correlation among the features of the datasets.

***Figure 2.2.1:*** *A glance correlation matrix of Global Confirmed Cases Dataset*

## Data Cleaning

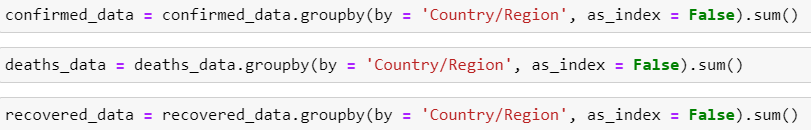
Data cleaning part is the crucial parts where it is the part to handle the missing value of every columns. Every column is in numerical and easy to compute the missing value with the mean or medium value. The summary of the total missing values was imputed by the percentage. The highest percentage represent the column that contains higher result of missing value.





***Figure 2.3.1:*** *summary of total missing values for every dataset*

As shown in figure 2.1.6 above, the columns of ‘province/State’ of each dataset have the most missing values. For this project, the column is removed. After that, every dataset was group by the country and sum the cases value for every column based on same country. It is much easier to visualize and further analysis. The dimension of the dataset’s changes to 191 of rows and 357 of rows.

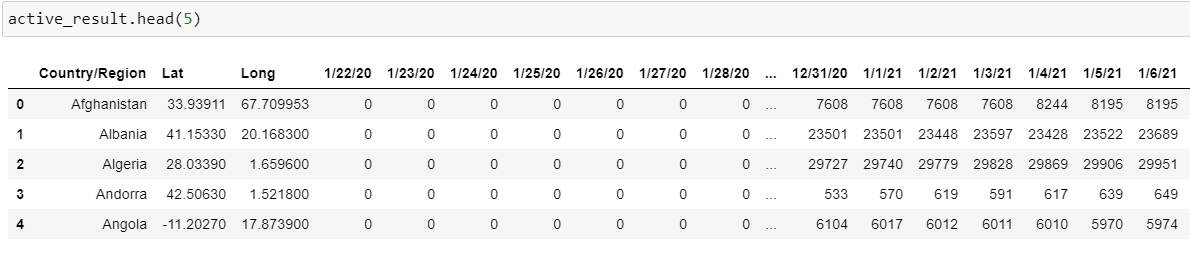


***Figure 2.3.2:*** *‘Group by’ python query based on similar country*

After handling the missing value of the datasets, the new data frame was created named ‘Active Results’ by compute using formula below:

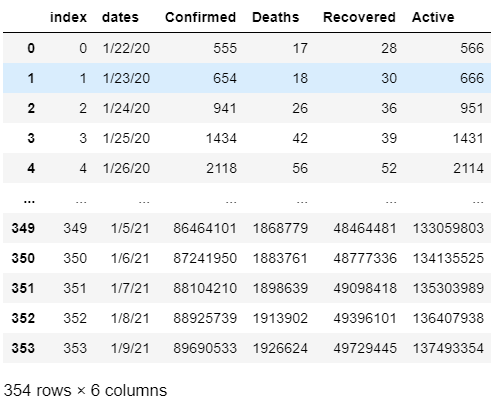
Active Result = Confirm Cases - (Deaths Cases + Recover Cases)

The result used the provided datasets that contained the details of the cases within dates. The country, longitude and latitude column remain same.



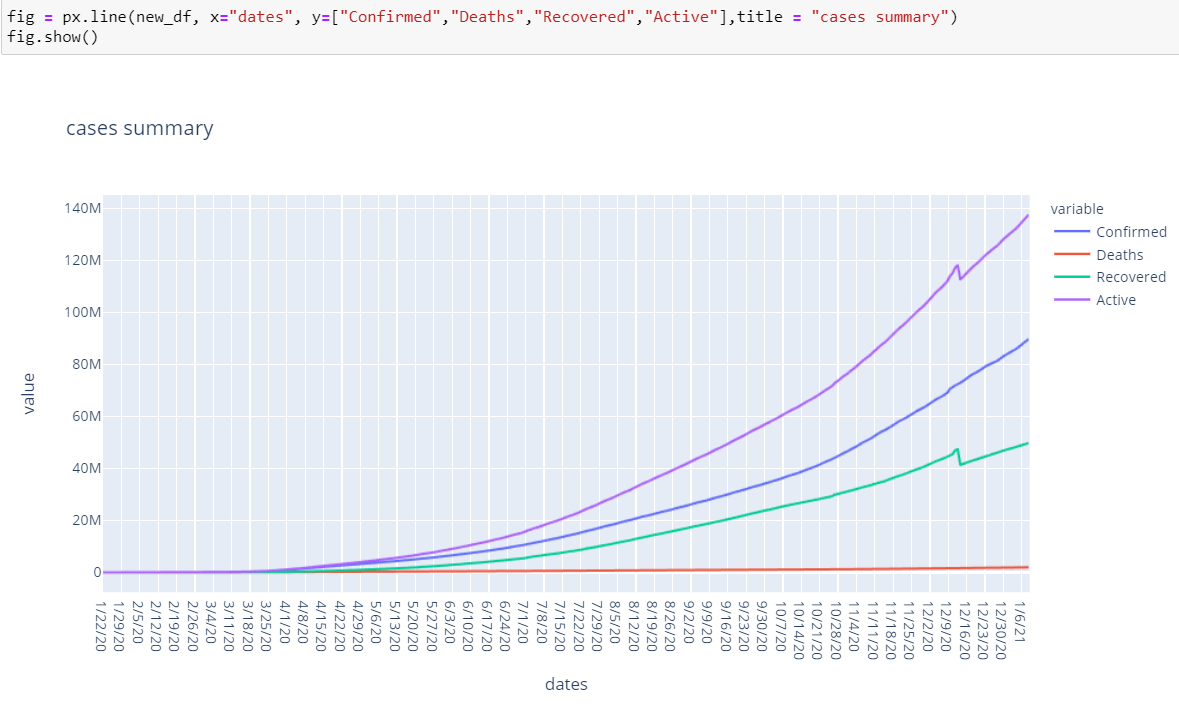
***Figure 2.3.3:*** *A glance of Global Active Cases Dataset*

Next, with all datasets, the summary of global result was computing, and the new data frame created to place all the values of the result. The data frame was created to easily make a prediction model later.



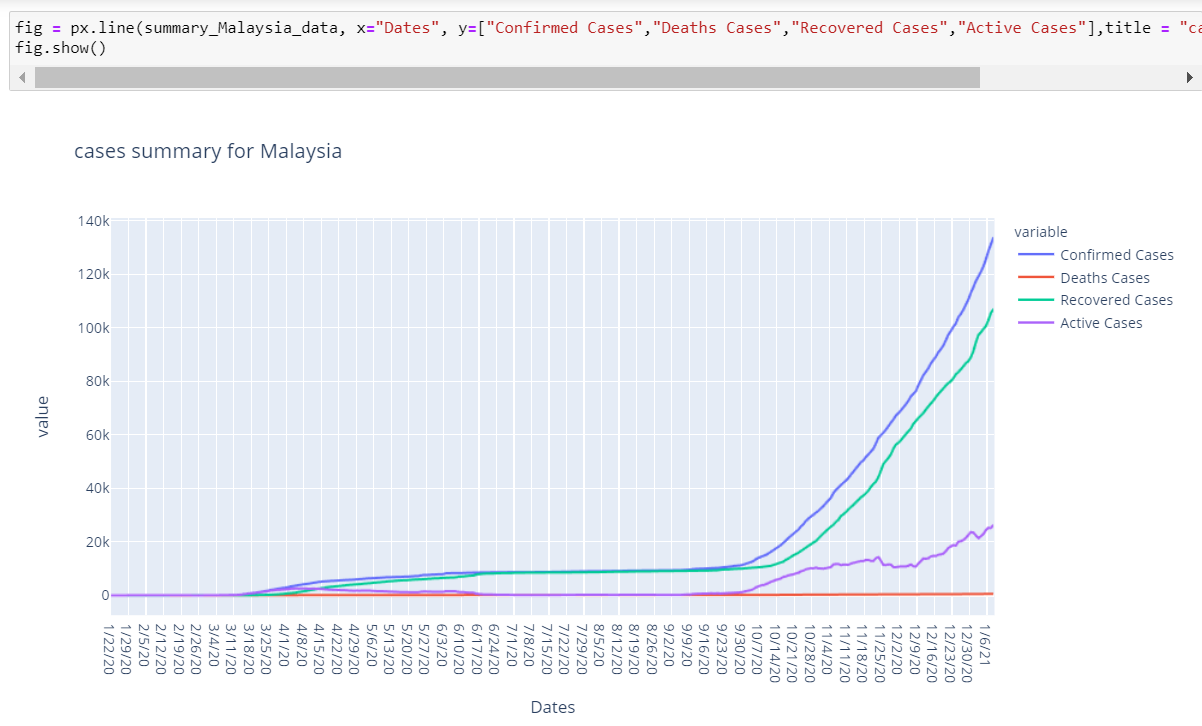
***Figure 2.3.4:*** *A glance of Global Cases Dataset*

## Data Visualisation

Visualizations is needed in data processing to view the outcome or information in an interesting and insightful, that will help clients and management to understand what the data is trying to show and what action can be taken with a second glance of the dashboard. sing visual elements in dashboards view, people can easily monitor and view the pattern or even the prediction trends.

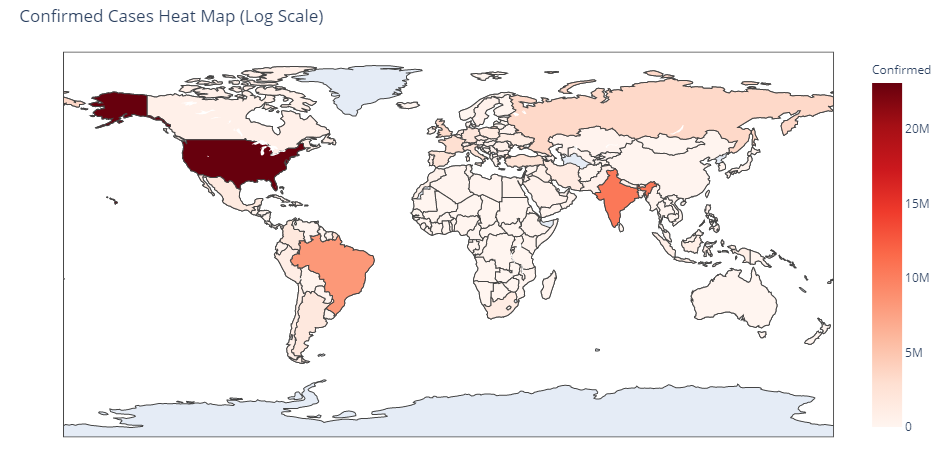
***Figure 2.4.1:*** *Plot of summary case all over the world*

The figure 2.4.1 above shows the plot line of the cases summary Data Frame that compute the sum of the cases for all country for every stated date. The plot line is nearly linear where the cases increasing every day for confirm, deaths and active cases but the deaths result line almost flatten where the deaths reported is small value.

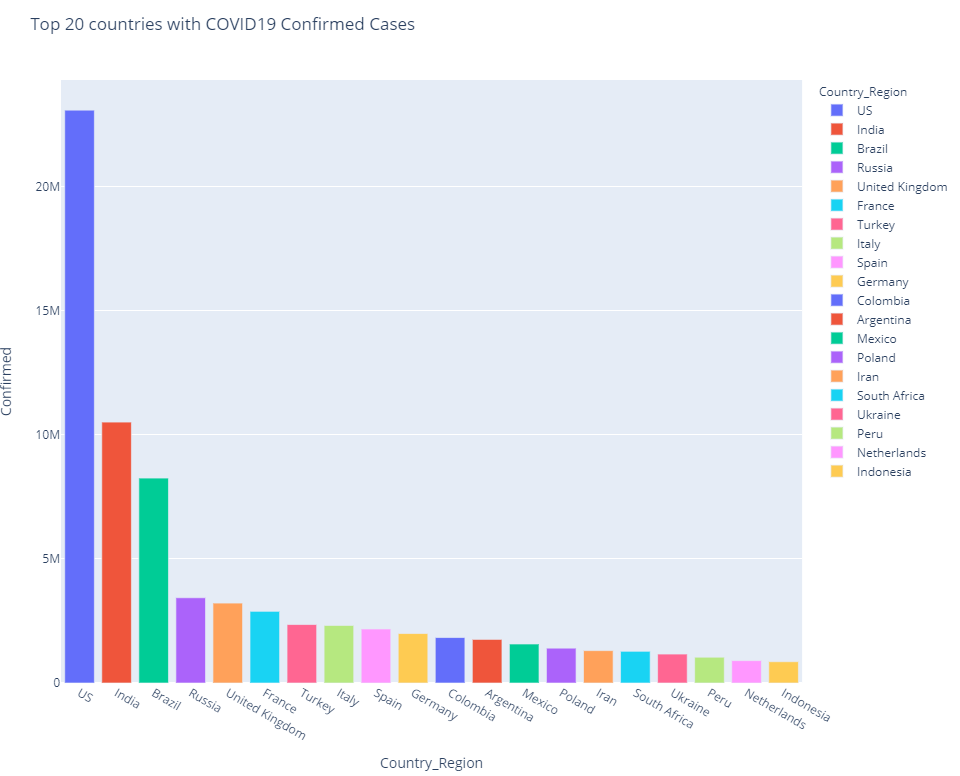


***Figure 2.4.2:*** *Plot of summary case of Malaysia*

Figure 2.4.2 above shows the plot line of Malaysia cases summary. The data of Malaysia’s cases from datasets are extract and group into new data frame named as ‘summary\_Malaysia\_data’ where it is contains the data of the total confirmed cases, deaths cases and recovered cases. From the graph, the ‘confirmed cases’ and ‘recovered cases’ are almost linear as the data increasing within continuous dates. The deaths cases are almost flat because the number of cases reported as death are small value and consistent.

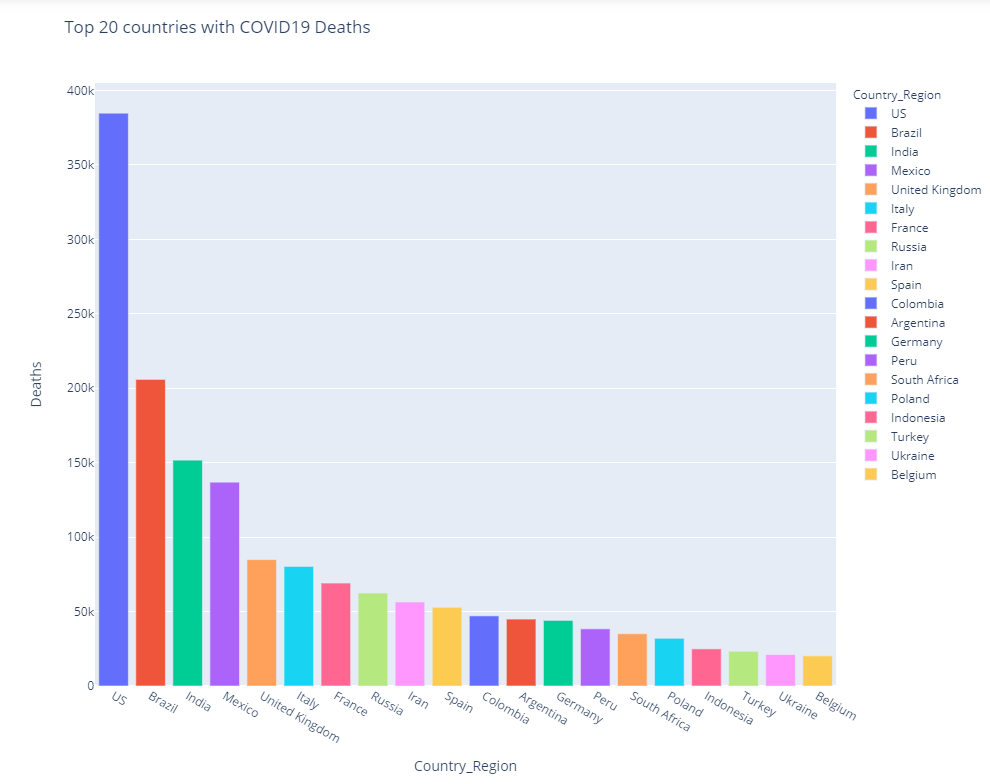


***Figure 2.4.3:*** *Confirm cases Heat Map*

***Figure 2.4.3:*** *Heat Map based on Confirmed Cases*

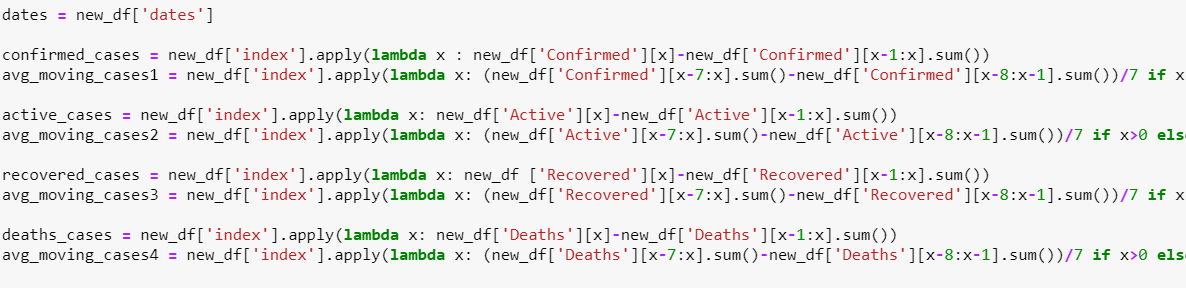
***Figure 2.4.4:*** *Bar Chart based on Top 20 Countries Confirmed cases COVID-19*

Figure 2.4.3 is the heat map based on confirmed cases and figure 2.4.4 is the bar chart of top 20 Countries confirmed cases of COVID-19. Both figures were extract from the date 13th January 2021 of dataset. From heat map figure, the darker color of red was the country had the highest confirm cases and the result shows United States (US) had the most highest cases among countries with total cases more than 23 million.

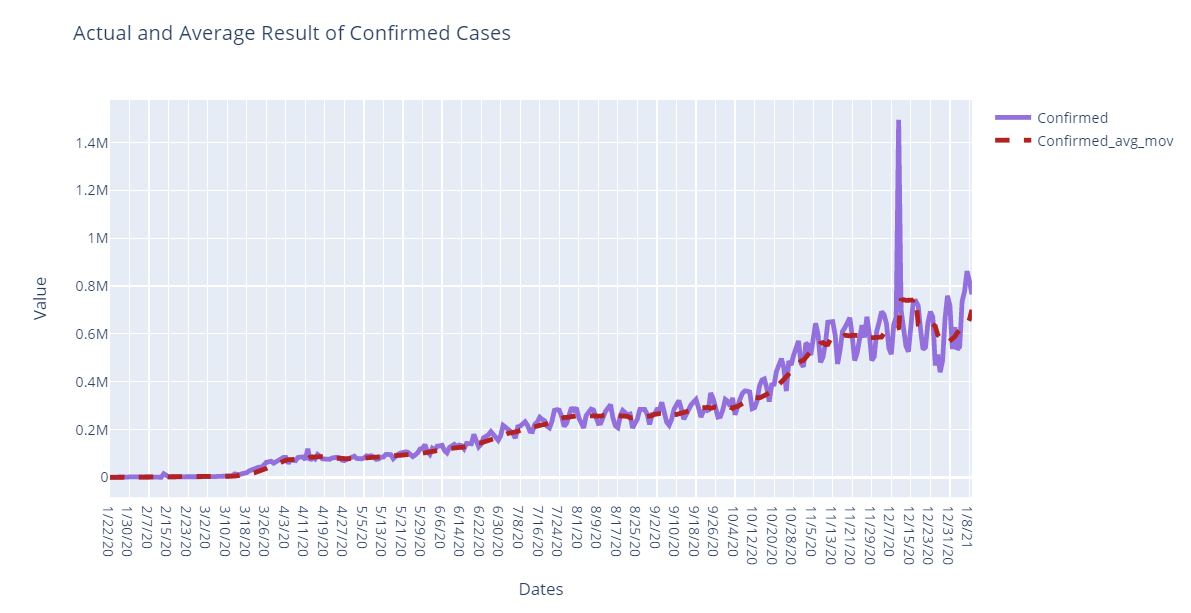
Figure 2.4.4 shows the bar chart that extract the 20 countries that have the highest number of confirmed cases recorded within COVID 19 epidemic. The result shows US lead the chart among other countries and followed by India that has total cases more than 10 million.

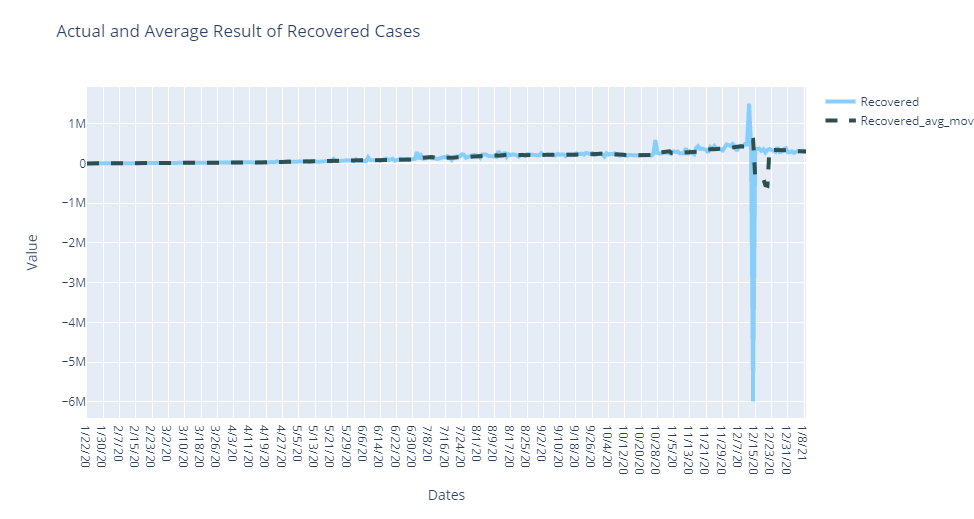
***Figure 2.4.5:*** *Bar Chart based on Top 20 Countries Deaths cases COVID-19*

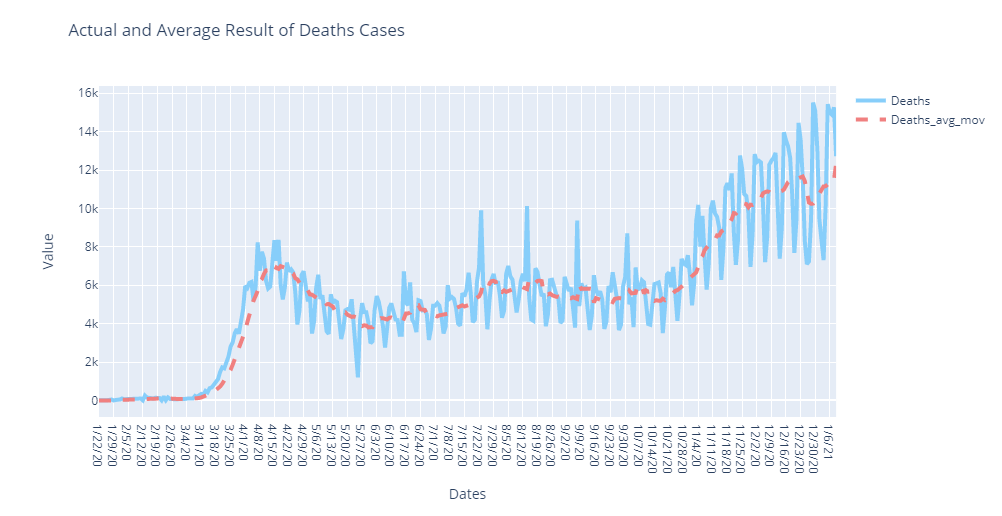
Figure 2.4.5 shows the bar chart that extract the 20 countries that have the highest number of deaths cases recorded within COVID 19 epidemic. The result shows the same as confirmed cases which US lead the chart with 384,764 cases among other countries and followed by India that has total cases 205, 964 cases recorded every day.



***Figure 2.4.3:*** *queries of computation the average of moving cases within 7 days*







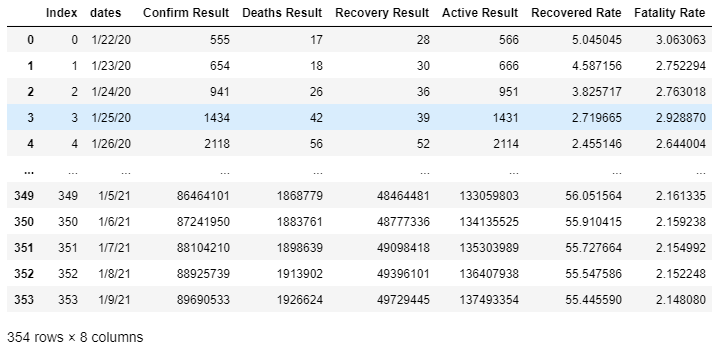
***Figure 2.4.4:*** *Actual and Average moving result of confirmed cases, deaths cases and recovered cases.*

In the graphs above, it is show the relationship of the total cases of confirmed, deaths and recovered with their average moving. A moving average implies that numbers take the previous days, take the average of those days, and graph it on the graph. It takes the last 7 days for a 7-day moving average, adds them up, and splits it by 7. The last 14 days will take an average of 14 days.

So, the datasets have data on COVID starting on 22nd February 2020 for the project planned. 7 days of COVID cases are required for the 7-day moving average. Between 22nd January 2020 with 30th February 2020, it added all the cases together, splitting them by 7. That point is then plotted. The cases started extremely growth is by the end of March where the virus start attack all over the world after its origin happen in Wuhan, China. Over time, it gives viewer an average line, and over a period of time it knocks out these wide peaks and valleys to the average.

# result and discussion

The considered COVID-19 datasets which are Confirmed Cases, Deaths Cases and Active Cases contains 272 records of cases all over the world. The datasets contain features of cases result such as Country/Region, Longitude, Latitude and dates. The data preprocessing and cleaning process remove the missing and outlier’s data values from the dataset. The resulted result dataset after processing is reduce to 191 rows and 357 columns required relevant features of the cases details.

The new data frame was created for prediction result that summarize the total cases of Confirmed, Deaths, Recovered and Active data frame with sum up cases of all country and divided by dates. From the summarize data frame, the new columns which are the percentage of Fatality Rate and Recovery rate are compute based on the calculation of (Recovered/Confirmed) x 100% for Recovery rate and (Deaths/Confirmed) x 100% for Fatality rate. The result of the new data frame is shown below.

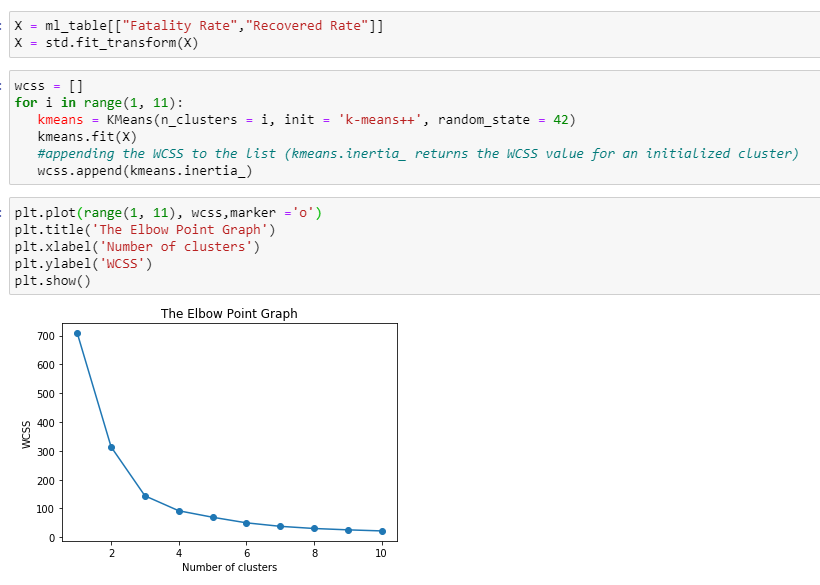
***Figure 3.0:*** *New Data Frame of cases with new columns*

## Clustering Method

An unsupervised machine learning task is cluster analysis, or clustering. It requires the discovery of natural grouping in data automatically. Clustering algorithms, unlike supervised learning (like predictive modelling), only interpret the input data and find natural groups or clusters in the feature space. There are many kinds of algorithms to cluster.

The most recognized clustering algorithm is K-Means Clustering, which involves assigning examples to clusters to minimize the variance within each cluster. To perform the K-Means Clustering, find the values of ‘K’ which the optimum number of cluster by using The Elbow Method and Silhouette value Method.

### The Elbow Method



***Figure 3.1.1:*** *The query of elbow method*

The variance (within-cluster sum of squares) decreases as the number of clusters increases. The elbow at 3 or 4 clusters is the most parsimonious balance between minimizing the number of clusters and minimizing the variance within each cluster, the k value can be 3 or 4.

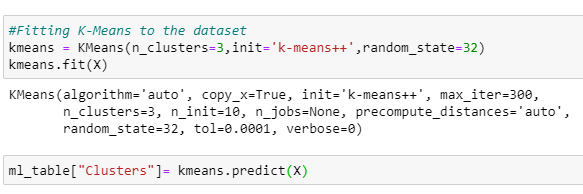
### Silhouette value Method



***Figure 3.1.2:*** *The query of Silhouette value Method*

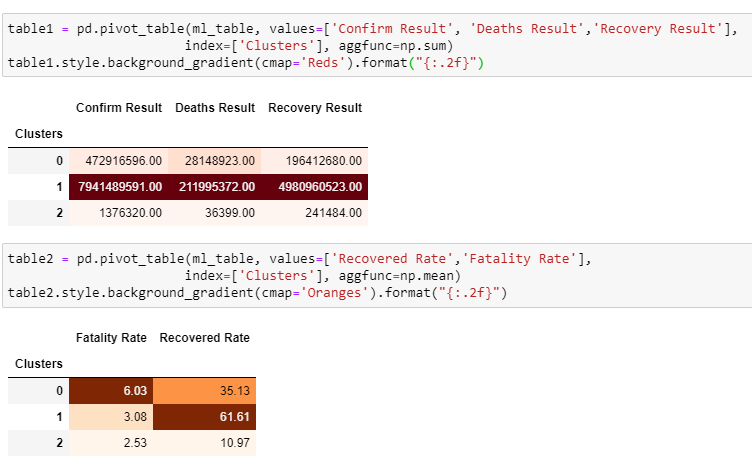
From observation, the optimum number of clusters at n = 3 and the result of the chosen for k values is 3.

### Fitting K Means algorithm

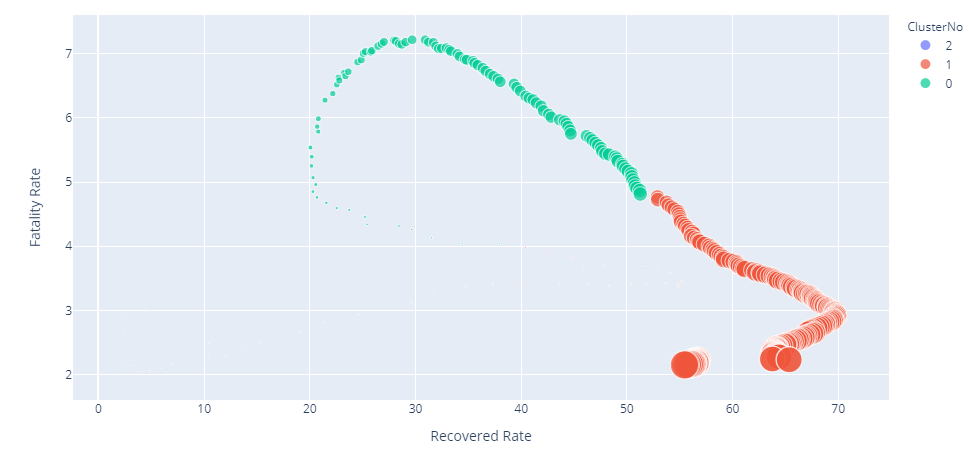


***Figure 3.1.3:*** *Fitting K Means algorithm for 3 clusters*

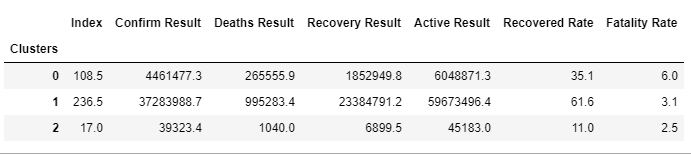
Next, all 3 clusters are group by into pivot table with Fatality rate and Recovery rate data and group by with Confirm cases, Deaths Cases and Recovered Cases with other pivot table.



***Figure 3.1.4:*** *Pivot tables with the clusters*



***Figure 3.1.5:*** *Scatter plot of the clusters*



***Figure 3.1.6:*** *Interpretation of the cluster summary*

The final analysis from clustering method are :

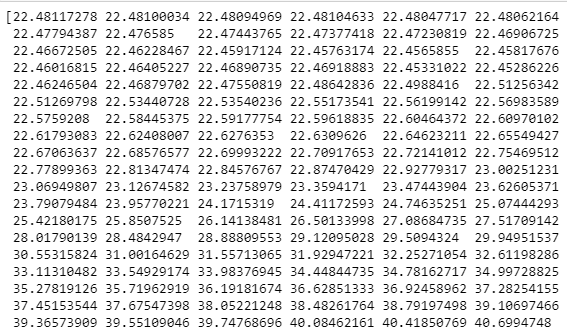
* **Cluster 0 :** The cases of all countries are around million cases with the average value of Recovered rate and Fatality rate and the second highest cases recorded.
* **Cluster 1 :** The total cases of the world are more than 20 million and the highest cases recorded.
* **Cluster 2 :** The total cases of the world are around thousands and the lowest cases recorded.

## Regression

In this project, the multiple variables or features are used to predict one output. The variables used are result of confirm, deaths, recovery and active. The output that will be predict using the model is Recovery rate.

### Multiple Linear Regression, MLR

## For the model, the data frame was created before that contains independent variables mark as ‘x’ and dependent variable state as ‘y’. The both was fitted by using the fit function and use the model to make predictions. The result of the prediction was shown in figure below.

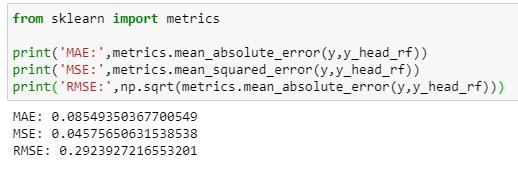


***Figure 4.2.1.1:*** *Prediction of the data*

The predict function, lm.predict(), predicts the y (dependent variable) using the linear model creation. The score of the model was **0.7974.** The model are compared with other model such as Random Forest and Decision Tree model.

### Random Forest

The variables are same as MLR model. The ‘RandomForestRegressor’ is used to solve regression problems on Random Forest. The important of the RandomForestRegressor is the n\_estimators parameter that defines the number of trees in random forest. The n\_estimator stated as 100 to see the algorithms performance. The score of the Random Forest is high than linear regression is **0.9999.**  The last step is to evaluate the performance of the algorithm. The metrics used to compute the mean absolute error, Mean Square Error and Root Mean Squared Error as shown in figure below.



***Figure 4.3.1.1:*** *Prediction of the data*

### Decision Tree

The object was created using DecisionTreeClassifier class and store the address in dt\_reg variable so it is easy to be access. The tree was fitted with selected features. The score of the Random Forest is high than linear regression is **0.9999** almost same as random forest.

# conclusion

The corresponding model is developed by analyzing the current Hubei epidemic situation data, and then the simulation is carried out. New COVID-19 cases were significantly predicted by the number of days and new cases all over the world, and the number of days and new recoveries significantly predicted new COVID-19 fatalities. According to this analysis, if the cases keep increasing within days, the predicted values of COVID-19 new cases and new deaths will be high as well. Among all the model, the best fitted model to predict the recovery rate and fatal rate was random forest and decision tree with score of 0.9999.

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