## Appendix D. Python code

The study considers Python as an application to establish the forecasting model and implement the external parameters such as vaccination rate, covid-variant, and control measures into the final model. The following figures describe the detailed code, explanation of the code is directly provided next to the coding sentences:

1. Import libraries into Python file:

```
1
2
       import pandas as pd
       import numpy as np
       import matplotlib.pyplot as plt
4
5
       from statsmodels.tsa.stattools import adfuller
       from statsmodels.tsa.arima.model import ARIMA
       from pandas import DataFrame
7
       from statsmodels.graphics.tsaplots import plot_acf, plot_pacf
8
       import warnings; warnings.filterwarnings(action='once')
9
       import math
10
```

2. Import excel file, this excel file contains the COVID-19 confirmed cases have been recorded daily from 27th April to 24th November 2021. After which, the data was plotted in order to visualize the distribution of the collected data

```
11
12  WB = pd.read_excel("Data import for ADF.xlsx")
13  # Plot data into a graph:
14  y = list(WB['Data value'])
15  x = list(WB['Date'])
16  plt.title('Recorded confirmed cases time series plot')
17  plt.plot(x, y)
18  plt.show()
```

3. The first step in model ARIMA development is considering whether the data is stationary. Hence, an ADF test is provided in the code. From which, d parameters will then be determined based on the stationarity of data in which differencing stage. The COVID-19 confirmed cases in this study is considered to be stationary on the first differencing, which means d = 1 in ARIMA model:

```
# ADF test
 21 😅 Considering the stationarity of the original data
 22
       X = WB['Data value'].values
       result = adfuller(X)
 23
       print('The result of ADF test of the original value is:')
 24
       print('ADF Statistic: %f' % result[0])
 25
 26
       print('p-value: %f' % result[1])
 27
       print('Critical Values:')
       for key, value in result[4].items():
 28
 29
           print(key, value)
 30
 31
       if result[0] < result[4]["5%"]:</pre>
           print("Reject Ho - Time Series is Stationary")
 32
 33
       else:
           print("Failed to Reject Ho - Time Series is Non-Stationary")
 34
35
       # Considering the 1st differencing
       WB['1st_Differencing'] = WB['Data value'] - WB['Data value'].shift(1)
36
37
       plt.plot(x, WB['1st_Differencing'])
       plt.title('First differencing of recorded confirmed cases')
38
39
       plt.show()
       result2 = adfuller(WB['1st_Differencing'].dropna())
41
       # Note we are dropping na values because the first value of the first difference is NA
       print('----')
42
       print('The result of ADF test of 1st Differencing is:')
43
44
       print('Critical Values:')
45
       for key, value in result2[4].items():
46
           print(key, value)
       print('ADF Statistic: %f' % result2[0])
47
       print('p-value: %f' % result2[1])
48
       if result2[0] < result2[4]["5%"]:</pre>
49
           print("Reject Ho - Time Series is Stationary")
50
51
       else:
52
           print("Failed to Reject Ho - Time Series is Non-Stationary")
53
```

4. AR(p) and MA(q) were determined using ACF and PACF method, the code is as following:

```
# ACF and PACF method
55
       WB['1st_Differencing'] = np.asarray(WB['1st_Differencing'])
       fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(16, 6), dpi= 80)
56
       ACF_plot = plot_acf(WB['1st_Differencing'].dropna(),ax=ax1, lags=60)
57
       PACF_plot = plot_pacf(WB['1st_Differencing'].dropna(),ax=ax2, lags=60, method='ywm')
58
59
      # Decoration:
      # 1.lighten the borders
       ax1.spines["top"].set_alpha(.3); ax2.spines["top"].set_alpha(.3)
       ax1.spines["bottom"].set_alpha(.3); ax2.spines["bottom"].set_alpha(.3)
62
       ax1.spines["right"].set_alpha(.3); ax2.spines["right"].set_alpha(.3)
63
       ax1.spines["left"].set_alpha(.3); ax2.spines["left"].set_alpha(.3)
64
       # 2.Font size of tick labels
       ax1.tick_params(axis='both', labelsize=12)
       ax2.tick_params(axis='both', labelsize=12)
67
       plt.show()
68
69
```

5. To implement the external parameters into the model, the study uses Pearson's correlation to determine the relationships between Confirmed cases and external data defined. However, since the correlation can easily be calculated through excel, the following code only considers when the Beta parameters of external data have been defined and implement these parameters into the ARIMA model for simplicity:

```
70
       # Implementing external parameters, the coefficients were performed in excel file
71
       WB2 = pd.read_excel("External Parameters.xlsx")
72
       Vaccination = WB2['Vaccination coverage (PPM)'].values
       Social_distance = WB2['Social distancing'].values
       Incubation = WB2['Incubation period'].values
74
75
       Generation_time = WB2['Generation time'].values
76
       beta_vaccination = 0.361
       beta_distancing = 0.577
78
       beta_incubation = -0.03
79
       beta_generation = 0.092
80
       X1 = beta_vaccination*Vaccination
81
       X2 = beta_distancing*Social_distance
82
       X3 = beta_incubation*Incubation
83
       X4 = beta_generation*Generation_time
       WB['External'] = X1 + X2 + X3 + X4
84
85
```

6. The ARIMA parameters and external parameters were used to forecast the COVID-19 confirmed cases in this stage, the following code provides a way to plot these data and include the real data for fitting. Noted that in this stage, a 95% confidence level is applied for COVID-19 confirmed cases collected

```
# ARIMA model: provide a forecasting model of ARIMA => ARIMA (1,1,0)
        model = ARIMA(WB['Data value'], order=(1,1,0))
 87
88
        RESULT = model.fit()
 89
        print(RESULT.summary())
90
 91
        fig2, ax = plt.subplots(1,2,figsize=(16,6), dpi= 80)
        residuals = DataFrame(RESULT.resid) # Line plot of residuals
 92
        residuals.plot(title="Residuals", ax=ax[0])
 93
        residuals.plot(title="Density", ax=ax[1], kind='kde') # Density plot of residuals
 94
        plt.show()
 95
96
97
        # Summary stats of residuals
 98
        print(residuals.describe())
        # Forecast ARIMA (1,1,0)
99
        WB['ARIMA(1,1,0)'] = RESULT.predict(start=0, end=207, dynamic=False)
        WB['Final Forecast result'] = WB['ARIMA(1,1,0)'] + WB['External']
103
        # 95% Confidence interval:
104
       CI = 1.96*np.std(WB['Data value'])/np.sqrt(len(x)) # since Confidence level = 95%, z-value = 1.96
105
       # Plot
106
       fig3, ax3 = plt.subplots(1,1)
107
        plt.title('Forecast of COVID-19 Confirmed Cases in HCMC', alpha = 0.5)
        plt.plot(x, WB['Data value'])
108
109
        plt.plot(x, WB['Final Forecast result'])
        ax3.fill_between(x, (WB['Data value'] - CI), (WB['Data value'] + CI), color = "b", alpha = 0.2)
110
        plt.legend(["COVID-19 Confirmed Cases", "ARIMA(1,1,0) Forecast"])
112 plt.show()
```

7. After providing forecast results, the study continued with calculating and plotting a 7-day moving average between forecasting results and real data. MSE and RMSE calculation was also included in this part for error comparison in this study:

```
114
       🗦 # Calculate the 7-day moving average of both confirmed cases and forecasting results:
115
       # For confirmed cases:
        window_size1 = 7
117
        i = 0
118
        moving_avg1 = [] # Provide an empty list to store moving averages
        # Loop through the array to consider every window of size 7:
119
        while i < len(WB['Data value']) - window_size1 + 1:</pre>
120
121
            list1 = list(WB['Data value'])
122
            window1 = list1[i:i+window_size1] # Store elements from i to i+window_size in list to get the current window
            window_average1 = round(sum(window1) / window_size1, 2) # Calculate the average of current window
123
            moving_avg1.append(window_average1) # Store the average of current window in moving average list
124
125
            i = i+1
127
        # for forecasting results:
        window_size2 = 7
128
129
        j = 0
        moving_avg2 = [] # Provide an empty list to store moving averages
130
        # Loop through the array to consider every window of size 7:
132
        while j < len(WB['Final Forecast result']) - window_size2 + 1:</pre>
            list2 = list(WB['Final Forecast result'])
133
134
            window2 = list2[j:j+window_size2] # Store elements from i to i+window_size in list to get the current window
            window_average2 = round(sum(window2) / window_size2, 2) # Calculate the average of current window
135
            moving_avg2.append(window_average2) # Store the average of current window in moving average list
136
137
            j = j+1
138
        x1 = np.linspace(0,201,201) # Specify the number of observation in 7-day moving average => 201 observations
139
140
        plt.plot(x1, moving_avg1)
        plt.plot(x1, moving_avg2)
141
        plt.legend(["COVID-19 Confirmed Cases","ARIMA(1,1,0) Forecast"])
142
        plt.title('7-day moving average of Covid-19 confirmed cases & forecasting result', alpha=0.5)
143
144
        plt.show()
146
        # RMSE Calculation:
147
        MSE = np.square(np.subtract(WB['Data value'], WB['Final Forecast result'])).mean()
        RMSE = math.sqrt(MSE)
148
149
        print("Mean Square Error: ", MSE)
        print("Root Mean Square Error: ", RMSE)
150
```