PREDICTION OF THE DEVELOPMENT OF COVID-19 CASE IN INDONESIA BASED ON GOOGLE TREND ANALYSIS

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PREDICTION OF THE DEVELOPMENT OF COVID-19 CASE IN INDONESIA BASED ON GOOGLE TREND ANALYSIS

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ABSTRACT

The global outbreak of the coronavirus disease (COVID-19) has recently hit many countries around the world. Indonesia is one of the 10 most affected countries. Search engines such as Google provide data on search activity in a population, and this data may be useful for analyzing epidemics. Leveraging data mining methods on electronic resource data can provide better insights into the COVID-19 outbreak to manage health crises in every country and around the world. This study aims to predict the incidence of COVID-19 by utilizing data from the Covid 19 Task Force and the Google Trends website. Linear regression and long-term memory (LSTM) models were used to estimate the number of positive COVID-19 cases.

KEYWORDS

Covid-19, Long ShortTerm Memori, Google Trend



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INTRODUCTION

Long holidays often encourage people to travel, even though movement and crowds can have an impact on increasing Covid-19 cases (Wen, Kozak, Yang, & Liu, 2020). According to data from the Covid-19 Handling Task Force, there is always an upward trend of positive cases occurring every holiday period (Sharpe Jr, Kuszyk, & Mossa-Basha, 2021). Google Trends is a website owned by Google.Inc that contains trends in the use of keywords on the Google search engine website and trending news (Jun, Yoo, & Choi, 2018). One of the benefits of Google Trends is for research. RNN has been used for sequential time series applications with temporal dependencies.

RNN which has the ability to process the current data by using the previous data. Meanwhile, the RNN is problematic to train long-term dependency data, which is solved

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by one of the RNN variants. The LSTM was anticipated by Hochreiter and Schmidhuber, has been used as an advanced version of the RNN network and has overcome the limitations of RNN by using a hidden layer unit known as a memory cell. The memory cells are self-connected which store the temporal state of the network and are controlled through three named gates: input gate, output gate and forget gate (Gers & Schmidhuber, 2001).

The work of input and output gates is used to control the flow of input and output of memory cells throughout the network (Sak, Senior, & Beaufays, 2014). In addition, a forget gate has been added to the memory cell, which passes high-weighted output information from the previous neuron to the next neuron. Information residing in memory depends on high activation yield; if the input unit has high activation, the information is stored in the memory cell. In addition, if the output unit has a high activation, the information will be passed on to the next neuron (Shahid, Zameer, & Muneeb, 2020). Otherwise, the high-weighted input information resides in the memory cell.

This study analyzes the development of Covid-19 cases associated with several keywords on Google Trends. In this study, several algorithms were tested to analyze the development of Covid 19 cases associated with keywords in Google Trends (Pan, Nguyen, Abu-Gellban, & Zhang, 2020).

RESEARCH METHOD

In the first stage of this research, we will explore the data in Google Trends. The keywords used are 'covid 19', 'ppkm', 'lockdown', 'ptm', 'wfh', 'vaccination', 'cluster', 'coronavirus', 'psbb', 'delta variant'. With a period starting from 2020-01-01 to 2021-11-10. The study began by downloading data on the development of daily spread on the COVID-19 website. Data Repository by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University (https://github.com/CSSEGISandData/COVID-19) and from https://data.humdata.org/dataset/indonesia-covid-19-cases-recoveries-and-deaths-per-province

The data in Google Trends is a random sample of Google search data. This data is anonymized (identity not disclosed), classified (search query topics defined), and aggregated (grouped together). Google Trends data can be filtered in two ways: real time and non-real time. Real time refers to a random sample of searches from the previous seven days, while non-real time refers to a random sample of the entire Google dataset, which can range from 2004 to 36 hours ago (Pretorius, Kruger, & Bezuidenhout, 2022). Google Trends are two separate random samples, so the graph will show one or the other, but not both at the same time.

RESULT AND DISCUSSION

1. Analysis of Community Activities on Daily Cases of Confirmed Covid 19 and their Visualization.

This step begins with preparing the data to be used, namely daily case variables with global mobility, visualizing daily cases with each global mobility variable and analyzing their correlation.



Prediction of the Development of Covid-19 Case In Indonesia Based on Google Trend
Analysis 1.408

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```
for column in dataakhir.columns:
    print(column)

tanggal
Kasusharian
Totalkasus
retail.and_recreation
grocery_and_pharmacy
branatt_s_tations
workplaces
residential

[20]:

dataakhir3=dataakhir.loc[:,['tanggal','retail_and_recreation','Kasusharian']]
dataakhir5=dataakhir.loc[:,['tanggal','grocery_and_pharmacy','Kasusharian']]
dataakhir6=dataakhir.loc[:,['tanggal','retail_and_recreation','Kasusharian']]
dataakhir6=dataakhir.loc[:,['tanggal','grocery_and_pharmacy','Kasusharian']]
dataakhir6=dataakhir.loc[:,['tanggal','retail_and_recreation','Kasusharian']]
dataakhir6=dataakhir.loc[:,['tanggal','residential','Kasusharian']]
dataakhir9=dataakhir.loc[:,['tanggal','residential','Kasusharian']]
```

Figure 1.11 Preparing data for analysis of community activities and daily confirmed cases of covid 19

```
dataakhir3.plot.line(x="tanggal", title="retail_and_recreation Mobility and Confirm Positif Covid 19");
plot.show(block=True);
dataakhir4.plot.line(x="tanggal", title="grocery_and_pharmacy Mobility and Confirm Positif Covid 19");
plot.show(block=True);
dataakhir5.plot.line(x="tanggal", title="parks Mobility and Confirm Positif Covid 19");
plot.show(block=True);
dataakhir6.plot.line(x="tanggal", title="transit_stations Mobility and Confirm Positif Covid 19");
plot.show(block=True);
dataakhir8.plot.line(x="tanggal", title="workplaces Mobility and Confirm Positif Covid 19");
plot.show(block=True);
dataakhir9.plot.line(x="tanggal", title="residential Mobility and Confirm Positif Covid 19");
plot.show(block=True);
```

Figure 1. 12 Coding for visualization of daily cases and community activities retail_and_recreation Mobility and Confirm Positif Covid 19

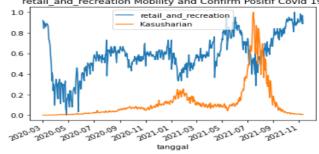


Figure 1.13 Visualization between Retail_and_recreation and daily cases
Figure 1.13 shows that if daily cases decrease then retail and recreation increases,
otherwise if daily cases increase then retail and recreation decreases.

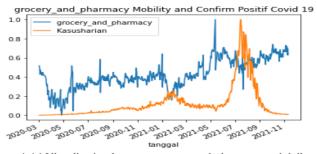


Figure 1.14 Visualization between grocery and pharmacy and daily cases

Figure 1.14 shows that grocery and pharmacy during the pandemic is consistently high, this shows that the public's need for medicines is quite high during the pandemic.

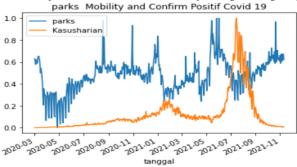


Figure 1.15 Visualization between parks and daily cases

Figure 1.15 shows that if daily cases decrease then activities in parks increase, on the contrary if daily cases increase then community activities in parks decrease.

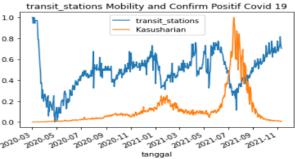


Figure 1.16 Visualization between transit_station and daily cases

Figure 1.16 shows that if daily cases decrease then activity at the station (transit_station) increases, otherwise if daily cases increase then activity at the station (transit_station) decreases.

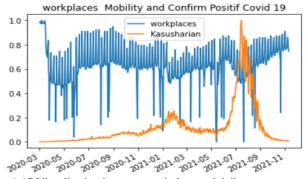


Figure 1.17 Visualization between workplace and daily cases

Figure 1.17 shows that in the early days of the pandemic work activities were quite high, but during high daily cases it can be seen that work activities fell drastically due to the lockdown.

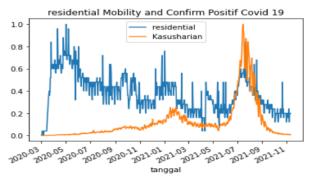


Figure 1.18 Visualization between residential and daily cases

Figure 1.18 shows that activity in housing is high at the beginning of the pandemic, daily cases are low, but at high daily cases activity in housing decreases but does not decrease at all.

```
import pandas as pd
from scipy.stats import pearson

#dataAkhi?adataAkhir.log(:,['date', 'retail_and_recreation', 'positir']]

# Convert dataFrame into series

list1 = dataAkhira['decusion']

# Apply the pearson(')

print('Pearsons correlation', '.37' \ corr)

# Convert dataFrame into series

list1 = dataAkhira['decusion', '.37' \ corr)

# Convert dataFrame into series

list1 = dataAkhira['decusion', '.37' \ corr)

## Apply the pearson(')

## Convert dataFrame into series

## Apply the pearson(')

## Appl
```

Figure 1.19 Pearson Correlation Analysis of daily cases and community activities Figure 1.19 shows the results of correlation analysis on daily cases and community activities using Pearson Correlation. From the picture, it can be seen that the negative correlations are retail and recreation, parks, transit station, workplaces retail and pharmacy meaning that if daily cases increase, these four activities will decrease and vice versa. While the correlation between daily cases and grocery and pharmacy, residential is positive even though the value is small.

```
x_simple = dataakhir3
my_r = x_simple.corr(method="spearman")
print((m)r)
x_simple = dataakhir4
my_r = x_simple.corr(method="spearman")
print(my_r)
print((m)r)
x_simple = dataakhir5
my_r = x_simple.corr(method="spearman")
print(my_r)
print('\n')
x_simple = dataakhir6
my_r = x_simple.corr(method="spearman")
print('\n')
x_simple = dataakhi
```

Figure 1.20 Spearman Correlation Analysis of daily cases and community activities

Figure 1.20 shows a correlation analysis using Spearman correlation, from the figure it shows that the positive correlations are retail_and_recreation, grocery_and_pharmacy, parks, transit_station. While the value is worksplaces and residential.

2. Analysis of Community Activities on the Total Confirmed Cases of Covid 19 and its Visualization.

This step begins with preparing the data to be used, namely the Total Case variable with global mobility, visualizing daily cases with each global mobility variable and analyzing the correlation.

```
dataakhir13=dataakhir.loc[:,['tanggal','retail_and_recreation','Totalkasus']]
dataakhir14-dataakhir.loc[:,['tanggal','grocery_and_pharmacy','Totalkasus']]
dataakhir16-dataakhir.loc[:,['tanggal','wrasit.stations','Totalkasus']]
dataakhir18-dataakhir.loc[:,['tanggal','workplaces','Totalkasus']]
dataakhir19-dataakhir.loc[:,['tanggal','residential','Totalkasus']]

dataakhir19-dataakhir.loc[:,['tanggal','residential','Totalkasus']]

[25]:

dataakhir13.plot.line(x='tanggal', title='retail_and_recreation Mobility and Total Cumalatif Confirm Posiplot.show(block=True);
dataakhir14.plot.line(x='tanggal', title='grocery_and_pharmacy Mobility and Total Cumalatif Confirm Posiplot.show(block=True);
dataakhir15.plot.line(x='tanggal', title='parks Mobility and Total Cumalatif Confirm Positif toplot.show(block=True);
dataakhir16.plot.line(x='tanggal', title='transit_stations Mobility and Total Cumalatif Confirm Positif toplot.show(block=True);
```

Figure 1.21 Preparation of data for analysis of total cases and community activities
Figure 1.21 shows the coding of data preparation for visualization between the total cases
and community activities.



Figure 1.22 Visualization between Retail and recreation and total cases Figure 1.22 shows a visualization between retail and recreation and total cases, where when the total cases are high, retail and recreation decreases.

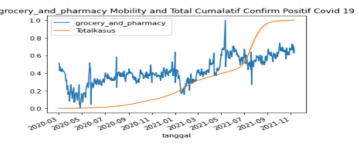


Figure 1.23 Visualization between grocery_and_pharmacy and total cases Figure 1.23 shows that grocery_and_pharmacy activities are quite stable, meaning that drug buying activities are quite stable, the increase occurs when the total number of cases is high.

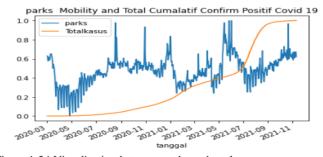


Figure 1.24 Visualization between parks and total cases
Figure 1.24 shows that community activities in parks at the beginning of the
pandemic were quite high, but when total cases were high, parks activities decreased.

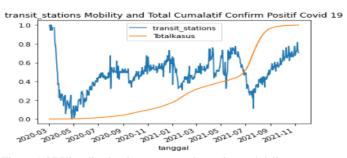


Figure 1.25 Visualization between transit_station and daily cases
Figure 1.25 shows that transit_station activity is high if daily cases are low. When

the daily case goes up, transit_station goes down.

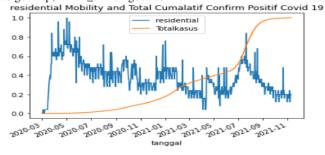


Figure 1.26 Visualization between residential and daily cases
Figure 1.26 shows that daily cases are low, so residential is high. However, when
daily cases are high, it can be seen that residential drops drastically.

```
# Apply the pearsonr()
corr, _ = pearsonr(list1, list2)
print('Pearsons correlation: %.3f' % corr)
print("\n")

Pearsons correlation: 0.593

Pearsons correlation: 0.748

Pearsons correlation: 0.325

Pearsons correlation: 0.304

Pearsons correlation: 0.144

Pearsons correlation: -0.401
```

Figure 1.27 Pearson correlation analysis between daily cases and community activities

Figure 1.27 shows the correlation value between daily cases and community mobility using the Pearson correlation formula. It can be seen that the highest correlation is 0.748, namely the correlation between daily cases and grocery_and_pharmacy, meaning that the correlation is quite high and positive.

Figure 1. 28 Spearman Correlation Analysis between total cases and community activities

Figure 1.28 shows the correlation analysis using Spearman, from the results obtained the highest correlation value is 0.784, namely the correlation between grocery_and_pharmacy and total cases. This means that the correlation is high and positive.

Analysis of Community Activities on the Total Confirmed Deaths of Covid 19 and its Visualization.

This step begins with preparing the data to be used, namely the Total Died variable with global mobility, visualizing daily cases with each global mobility variable and analyzing the correlation.

```
datashir23-datashir1.ole[;, 'senggal', 'retall_end_rerestion', 'totleminggal']]
datashir23-datashir1.ole[;, 'tanggal', 'gravel_end_rerestion', 'totleminggal']]
datashir23-datashir1.ole[;, 'tanggal', 'prike', 'totleminggal']]
datashir23-datashir1.ole[;, 'tanggal', 'prike', 'totleminggal']]
datashir23-datashir1.ole[;, 'tanggal', 'retalminggal']]
datashir23-datashir1.ole[;, 'tanggal', 'retalminggal']]

datashir23.plot,line(='tanggal', 'tile='retall_end_recreation Hobility and total meninggal Confirm Covid 10");

plot.show(block='frue);
datashir23.plot.line(='tanggal', tile='retall_end_recreation Hobility and total meninggal Confirm Covid 10");
plot.show(block='frue);
datashir24.plot.line(='tanggal', till="prike Hobility and total meninggal Confirm Covid 10");
plot.show(block='frue);
datashir24.plot.line(='tanggal', till="renami_stations Hobility and total meninggal Confirm Covid 10");
datashir24.plot.line(='tanggal', till="renami_stations Hobility and total meninggal Confirm Covid 10");
datashir24.plot.line(='tanggal', till="workplaces Hobility and total meninggal Confirm Covid 10");
datashir24.plot.line(='tanggal', till="workplaces Hobility and total meninggal Confirm Covid 10");
```

Figure 1.29 Data preparation for analysis of total deaths by community activities

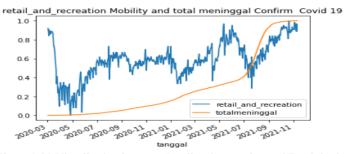


Figure 1.30 Visualization between Retail and recreation and Total death Figure 1.30 shows a visualization between retail and recreation and total deaths, where when total deaths are low, retai and recreation is high, on the other hand, total deaths are high, retail and recreation is low.



Figure 1.31 Visualization between grocery_and_pharmacy and Total died Figure 1.31 shows a visualization between grocery_and pharmacy and total deaths, where when the total deaths are low, grocery_and_pharmacy is high, on the other hand, the total deaths are high, so grocery and pharmacy is low.

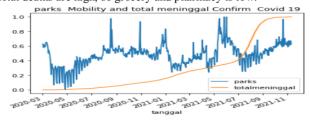


Figure 5.32 Visualization between Parks and Total died

Figure 1.32 shows a visualization between Parks and the total death toll, where when the total death toll is low, Parks is high, on the other hand, when the total death toll is high, Parks is low.



Figure 1.33 Visualization between transit_station and Total died

Figure 1.33 shows a visualization between transit_station and total deaths, where when the total death toll is low, the transit_station is high, otherwise the total death is high, the transit_station is low.

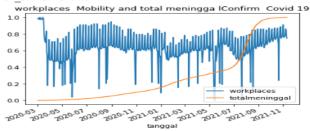


Figure 1.34 Visualization between workplaces and Total dies

Figure 1.34 shows a visualization between the workplaces and the total number of deaths, where when the total number of deaths is low, the workplaces are high, otherwise the total number of deaths is high, the workplaces are low.

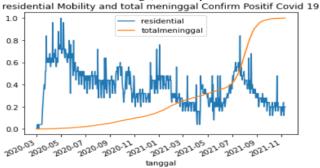


Figure 1.35 Visualization between Residential and Total Dies

Figure 1.35 shows a visualization between residential and total deaths, where when the total death toll is low, residential is high, on the other hand, the total death toll is high, the residential is low.

```
# Apply the pearson()
corr, _ = pearsons correlation: %.3f' % corr)
print('Pearsons correlation: %.3f' % corr)
print(in')
# Convert dataframe into series
list1 = dataskhir26['transit_stations']
# Apply the pearson()
corr, _ = pearsons correlation: %.3f' % corr)
print('Pearsons correlation: %.3f' % corr)
print('Pearsons correlation: %.3f' % corr)
print(in')

# Apply the pearson()
corr, _ = pearsonf(list1, list2)
corr, _ = pearsonf(list1, list2)
print('No')

# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('No')
# Convert dataframe into series
list1 = dataskhir29['residential']

# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('No')
# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('No')

# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('Pearsons correlation: %.3f' % corr)
print('Pearsons correlation: %.3f' % corr)

# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('Pearsons correlation: %.3f' % corr)

# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('Pearsons correlation: %.3f' % corr)

# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('Pearsons correlation: %.3f' % corr)
# Apply the pearsonf()
corr, _ = pearsonf(list1, list2)
print('Pearsons correlation: %.3f' % corr)
# Apply the pearsonf()
corr, _ = pearsons correlation: %.3f' % corr)
# Apply the pearsonf()
corr, _ = pearsons correlation: %.3f' % corr)
# Apply the pearsonf()
corr, _ = pearsons correlation: %.3f' % corr)
# Apply the pearsonf()
corr, _ = pearsons correlation: %.3f' % corr)
# Apply the pearsonf()
# Apply the
```

Figure 1.36 Pearson Correlation Analysis between total deaths and community activities

Figure 5.36 shows the Pearson correlation value between total deaths and community mobility using the Pearson correlation formula. It can be seen that the highest correlation is 0.733, which is the correlation between total deaths and grocery and pharmacy, meaning that the correlation is quite high and positive (Prawoto, Priyo Purnomo, & Az Zahra, 2020).

```
my.r = x_maple.corr(method="spearman")
print('\n')
pri
```

Figure 1.37 Spearman Correlation Analysis between total deaths and community activities

Figure 1.37 shows the correlation value between total deaths and community mobility using the Spearman correlation formula. It can be seen that the highest correlation is 0.784, which is the correlation between total deaths and grocery and pharmacy, meaning that the correlation is quite high and positive.

Prediction of Number of New Cases Per Day using Long Short Term Memory (LSTM)

At the initial stage, determine the dataset_train, namely the number of new_cases_per_day and as the dataset_test is the number of new_cases_per_days, the code is as follows:

```
dataset_train= pd.DataFrame (training_set, columns = ['Jumlah_Kasus_Baru_per_Hari'])
dataset_test= pd.DataFrame (test_set, columns =['Jumlah_Kasus_Baru_per_Hari'])
```

Figure 1.38 Setting training data and testing data for prediction

Then import the packages needed for prediction.

```
# Mengimpor library
import numpy as np
      import matplotlib.pyplot as plt
     import pandas as pd
from keras.layers import Dense,Dropout,SimpleRNN,LSTM
    # Proses feature scaling
from sklearn.preprocessing import MinMaxScaler
sc = MinMaxScaler(feature_range = (0, 1))
training_set_scaled = sc.fit_transform(training_set)
represer-7
     rentang=7
  rentang=/
# Membuat prediksi dengan 60 time-window (3 bulan)
X_train = []
y_train = []
for i in range(rentang, training_set.shape[0]):
    X_train.append(training_set_scaled[i-rentang:i, 0])
    y_train.append(training_set_scaled[i, 0])
X_train, y_train = np.array(X_train), np.array(y_train)
   # Reshaping
X_train = np.reshape(X_train, (X_train.shape[0], X_train.shape[1], 1))
# Mulai membuat RNN
Mesin_saham = Sequential()
# Menambah layer LSTM yang pertama dan Dropout regularisation
Mesin_saham.add(SimpleRNM(units = 50, return_sequences = True, input_shape = (X_train.shape[1], 1)))
Mesin_saham.add(Dropout(0.2))
# Menambah layer LSTM yang kedua dan Dropout regularisation
Mesin_saham.add(SimpleRNN(units = 50, return_sequences = True))
Mesin_saham.add(Dropout(0.2))
# Menambah layer LSTM yang ketiga dan Dropout regularisation
Mesin_saham.add(SimpleRNN(units = 50, return_sequences = True))
Mesin_saham.add(Dropout(0.2))
 Mesin_saham.add(Dropout(0.2))
 Mesin_saham.add(Dense(units = 1))
 # Melihat rancangan network LSTM kita
Mesin_saham.summary()
 Mesin_saham.compile(optimizer = 'adam', loss = 'mean_squared_error', metrics=["acc"])
 hist =Mesin_saham.fit(X_train, y_train, validation_split=0.3,epochs = 100, batch_size = 32,verbose=2
 # Mengimpor data saham sesungguhnya untuk Test set
  X_{test} = []
  \label{eq:formula} \textbf{for i in} \  \, \text{range}(\text{rentang, saham\_real.shape}[\theta] \, + \, \text{X\_train.shape}[1]) \colon \\
      X_test.append(inputs[i-rentang:i, 0])
  X_{test} = np.array(X_{test})
  X\_test = np.reshape(X\_test, (X\_test.shape[0], X\_test.shape[1], 1))
  predicted_stock_price = Mesin_saham.predict(X_test)
  predicted_stock_price = sc.inverse_transform(predicted_stock_price)
  # Visualisasi perbandingan hasil prediksi dan data sesunguhnya
  plt.plot(saham_real, color = 'red', label = 'Jumlah_Kasus_Baru_per_Hari sesungguhnya')
  plt.plot(predicted_stock_price, color = 'blue', label = 'Jumlah_Kasus_Baru_per_Hari prediksi')
  plt.title('Prediksi Jumlah_Kasus_Baru_per_Hari')
  plt.xlabel('Waktu')
  plt.ylabel('Prediksi Jumlah Kasus Baru Covid 1 per Hari')
```

Figure 1.39 Importing packages needed for prediction The next process is to build the model with epoch.

```
Total parems: 17,001
Trainable parems: 17,001
Trainable parems: 17,001

Sport 17/002

Sport 17/002
```

Figure 1.40 The process of building a predictive model

Prediction results are visualized as follows:

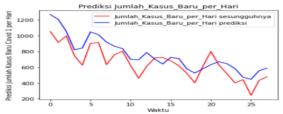


Figure 1.41 Predicted number of cases per day

After making predictions, then testing the models that have been obtained and making visualizations.

Figure 1. 42 Prediction Graph

From Figure 5.42 it can be seen that between the validation loss values (red) and train loss (blue), the graphs are close together. This shows that the predictions made are quite accurate and are also shown in Figure 5.43 with a value of RMSE = 145,135 which is quite small.

```
from sklearn.metrics import mean_squared_error
from math import sqrt
rmse = sqrt(mean_squared_error(saham_real, predicted_stock_price))
print('Test RMSE: %.3f' % rmse)
Test RMSE: 145.135
```

Figure 1.43 Evaluation Results using RMSE

5. Predicting the Number of Cumulative Cases using Long Short Term Memory (LSTM)

At the initial stage, determine the dataset_train, namely the number of new cases per day and as the dataset test is the Cumulative Number of Cases, the coding is as follows:

Figure 1.43 Setting training data and testing data for prediction

Then import the packages needed for prediction.

```
import numpy as np
import matplotlib.pyplot as plt
import matplotlib.pyplot as plt
import matplotlib.pyplot as plt
from keras.layers import Dense.Dropout.SimpleRNN.LSTM

# Proses resture scaling
from sklearn.preprocessing import MinMaxScaler
from the filter imports import MinMaxScaler
from tensor flow.
from tensor flow.keras.models import MinMaxScaled(1-eq);
from tensor flow.keras.models import Sequential
from tensor flow.keras.layers import Dense
from tensor flow.keras.layers import Sequential
from tensor flow.keras.layers import Dense
from tensor flow.keras.layers import Dense
from tensor flow.keras.layers import Sequential

### Meansman ### MinMax Sequenti
```

Figure 1. 44 Setting up the necessary packages for modeling The next step is to build a model using epochs and then visualize it.

```
Epoch 1/100
13/13 - 4s - loss: 0.1989 - acc: 0.0000e+00 - val_loss: 0.0651 - val_acc: 0.0057
Epoch 2/100
13/13 - 0s - loss: 0.0861 - acc: 0.0000e+00 - val_loss: 0.1286 - val_acc: 0.0000e+00
13/13 - 0s - loss: 0.0470 - acc: 0.0000e+00 - val_loss: 0.2368 - val_acc: 0.0000e+00
Epoch 3/100
13/13 - 0s - loss: 0.0470 - acc: 0.0000e+00 - val_loss: 0.2368 - val_acc: 0.0000e+00
Epoch 4/100
13/13 - 0s - loss: 0.0486 - acc: 0.0000e+00 - val_loss: 0.0290 - val_acc: 0.0057
Epoch 5/100
13/13 - 0s - loss: 0.0346 - acc: 0.0000e+00 - val_loss: 0.0480 - val_acc: 0.0057
Epoch 6/100
13/13 - 0s - loss: 0.0301 - acc: 0.0000e+00 - val_loss: 0.0480 - val_acc: 0.0057
Epoch 7/100
13/13 - 0s - loss: 0.0223 - acc: 0.0000e+00 - val_loss: 0.0207 - val_acc: 0.0057
Epoch 8/100
13/13 - 0s - loss: 0.0196 - acc: 0.0000e+00 - val_loss: 0.0207 - val_acc: 0.0057
Epoch 10/100
13/13 - 0s - loss: 0.0166 - acc: 0.0000e+00 - val_loss: 0.0344 - val_acc: 0.0057
Epoch 10/100
13/13 - 0s - loss: 0.0166 - acc: 0.0000e+00 - val_loss: 0.0344 - val_acc: 0.0057
Epoch 11/100
13/13 - 0s - loss: 0.0126 - acc: 0.0000e+00 - val_loss: 0.0542 - val_acc: 0.0057
Epoch 11/100
13/13 - 0s - loss: 0.0126 - acc: 0.0000e+00 - val_loss: 0.0542 - val_acc: 0.0057
Epoch 13/100
13/13 - 0s - loss: 0.0147 - acc: 0.0000e+00 - val_loss: 0.0552 - val_acc: 0.0057
Epoch 13/100
13/13 - 0s - loss: 0.0147 - acc: 0.0000e+00 - val_loss: 0.0244 - val_acc: 0.0057
Epoch 13/100
13/13 - 0s - loss: 0.0134 - acc: 0.0000e+00 - val_loss: 0.0244 - val_acc: 0.0057
Epoch 13/100
13/13 - 0s - loss: 0.0134 - acc: 0.0000e+00 - val_loss: 0.0244 - val_acc: 0.0057
Epoch 13/100
13/13 - 0s - loss: 0.0108 - acc: 0.0000e+00 - val_loss: 0.0044 - val_acc: 0.0057
Epoch 13/100
13/13 - 0s - loss: 0.0108 - acc: 0.0000e+00 - val_loss: 0.0044 - val_acc: 0.0057
Epoch 13/100
```

Figure 1.45 Model building process

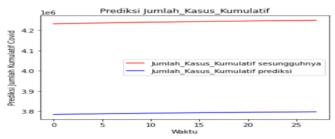


Figure 1. 46 Visualization of the built prediction model

```
fig, ax = plt.subplots(figsize=(14,6))
ax.plot(hist.history['loss'], 'b' ,label = 'train loss', linewidth=2)
ax.plot(hist.history['valloss'], 'r', label = 'Validation loss', linewidth=2)
ax.set_ritle('model loss')
ax.set_xlabel('mse')
ax.set_xlabel('epoch')
ax.legend()
plt.show()

Thouse ross

Validation loss

Validation loss
```

Figure 1.47 Prediction graph

From Figure 1.47 it can be seen that between the validation loss values (red) and the train loss (blue), the graphs are close together. This shows that the predictions made are quite accurate and are also shown in Figure 1.48 with a sufficient value of RMSE = 449516,694.

```
rmse = sqrt(mean_squared_error(saham_real, predicted_stock_price))
print('Test RMSE: %.3f' % rmse)
Test RMSE: 449516.694
```

Figure 5.48 Evaluation Results using RMSE

6. Prediction of the Number of Cumulative Death Cases using Long Short Term Memory (LSTM)

At the initial stage, the dataset_train is determined, namely the cumulative number of death cases and as the dataset_test is the Cumulative Number of Cases Death Cumulative, the coding is as follows:

Figure 1.49 Setting training data and testing data for prediction

Figure 1.50 Importing packages needed for prediction The next step is to build a model using epochs and then visualize it

Model: "sequential_2"

Layer (type)	Output Shape	Param #
simple_rnn_8 (SimpleRNN)	(None, 7, 50)	2600
dropout_8 (Dropout)	(None, 7, 50)	0
simple_rnn_9 (SimpleRNN)	(None, 7, 50)	5050
dropout_9 (Dropout)	(None, 7, 50)	0
simple_rnn_10 (SimpleRNN)	(None, 7, 50)	5050
dropout_10 (Dropout)	(None, 7, 50)	0
simple_rnn_11 (SimpleRNN)	(None, 50)	5050
dropout_11 (Dropout)	(None, 50)	0
dense_2 (Dense)	(None, 1)	51
Total params: 17,801 Trainable params: 17,801 Non-trainable params: 0		

Figure 1.51 The process of building a predictive model

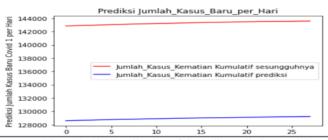


Figure 1. 52 Visualization of the built prediction model

```
fig, ax = plt.subplots(figsize=(14,6))
ax.plot(hist.history['loss'], 'b' ,label = 'train loss', linewidth=2)
ax.plot(hist.history['val_loss'], 'r', label ='Validation loss', linewidth=2)
ax.set_title('model loss')
ax.set_ylabel('mse')
ax.set_xlabel('epoch')
ax.legend()
plt.show()
```

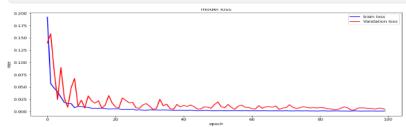


Figure 1.53 Prediction graph

From Figure 1.53, it can be seen that between the validation loss values (red) and the train loss (blue), the graphs are close together. This shows that the predictions made are quite accurate and are also shown in Figure 5.54 with a value of RMSE = 14331,656 which is quite small.

```
rmse = sqrt(mean_squared_error(saham_real, predicted_stock_price))
print('Test RMSE: %.3f' % rmse)
Test RMSE: 14331.656
```

Figure 1.54 Evaluation Results using RMSE

CONCLUSION

Based on the results of research that has been carried out on positive confirmed COVID-19 data downloaded from Google Trend from January 1, 2020 to November 10, 2021 with 617 records including daily case variables, total cases, total deaths with global mobility variables (community activities) including retail and recreation, grocery and pharmacy, parks, transit stations, workplaces, a model has been obtained to predict the number of cases per day, predict the number of cumulative cases, and the number of cumulative deaths.

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The best prediction result is the prediction of the number of cases per day with an RMSE = 145,135. Meanwhile, the highest correlation analysis is 0.784 between the total death variable and grocery and pharmacy.

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