

In [17]:

```
import pandas as pd
import numpy as np
from datetime import datetime
import matplotlib.pyplot as plt
```

In [18]:

```
df = pd.read_csv('D:\Covid\complete-Covid-19.csv')# read whole data
print('Covid Dataframe rows and col::\t',df.shape)
df2 = df[(df['Name of State / UT'] == 'West Bengal')] #read data of westbengal 89 rowa 10 c
print('WB dataframe rows and col::\t',df2.shape)
print('\nWB dataframe Description::\n\n',df2.describe())
```

Covid Dataframe rows and col:: (3056, 10)

WB dataframe rows and col:: (89, 10)

WB dataframe Description::

	Latitude	Longitude	Total Confirmed cases	Death \
count	8.900000e+01	8.900000e+01	89.000000	89.000000
mean	2.298680e+01	8.785500e+01	2508.404494	145.842697
std	4.287411e-14	2.000792e-13	3160.836194	157.859467
min	2.298680e+01	8.785500e+01	1.000000	0.000000
25%	2.298680e+01	8.785500e+01	116.000000	5.000000
50%	2.298680e+01	8.785500e+01	922.000000	33.000000
75%	2.298680e+01	8.785500e+01	3667.000000	272.000000
max	2.298680e+01	8.785500e+01	11494.000000	485.000000

	Cured/Discharged/Migrated	New cases	New deaths	New recovered
count	89.000000	89.000000	89.000000	89.000000
mean	939.595506	129.134831	5.449438	61.730337
std	1354.832720	147.579577	10.934735	95.343609
min	0.000000	0.000000	0.000000	0.000000
25%	16.000000	16.000000	0.000000	0.000000
50%	151.000000	58.000000	3.000000	9.000000
75%	1339.000000	183.000000	8.000000	90.000000
max	5494.000000	476.000000	98.000000	518.000000

In [19]:

```
df2.plot(x='Date', y='Total Confirmed cases', style='-')
plt.title('West Bengal--Date vs Total Confirmed cases')
plt.xlabel('Particular date')
plt.ylabel('Confirmed case')
plt.show() #plot date Vs confirm case graph
```

'''From the graph above, we can clearly see that there is a positive linear relation between the number of hours studied and percentage of score.'''

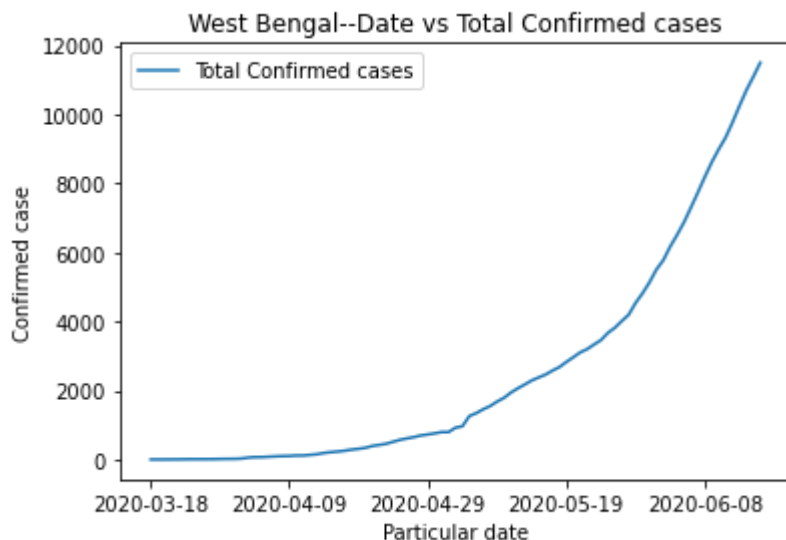
#Now we have an idea about statistical details of our data.

#The next step is to divide the data into "attributes" and "Labels".

'''Attributes are the independent variables(here it is X) while labels are dependent variables whose values are to be predicted.'''

#We want to predict the number of confirmed cases on a particular day.

#Therefore our attribute set will consist of the "Date" column, and the label will be the "



Out[19]:

'Attributes are the independent variables(here it is X) while labels are dependent variables(here it is Y) \nwhose values are to be predicted.'

In [20]:

```
# iloc is integer-location based indexing for selection by position.
#it will select till the second last column of the data frame instead of the last column

Z = df2.iloc[:,0].values # iloc is integer-location based indexing for selection by position
X2=[(datetime.strptime(t, '%Y-%m-%d').date()-datetime.strptime(Z[0], '%Y-%m-%d').date()).days
X=np.array(X2)
y = df2.iloc[:, 4].values
```

In [21]:

```
print(X) # Days of WB cases
print(y) # No of WB confirmed case on particular day
```

```
[ 0  1  2  3  4  5  6  7  8  9 10 11 12 13 16 17 18 19 20 21 22 23 24 25
26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49
50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73
74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90]
[   1   1   1   3   4   7   9   9  10  10  15  18
 19  26  53  69  69  80  91  99 103 116 116 134
152 190 213 231 255 287 310 339 392 423 456 514
571 611 649 697 725 758 795 795 922 963 1259 1344
1456 1548 1678 1786 1939 2063 2173 2290 2377 2461 2576 2677
2825 2961 3103 3197 3332 3459 3667 3816 4009 4192 4536 4813
5130 5501 5772 6168 6508 6876 7303 7738 8187 8613 8985 9328
9768 10244 10698 11087 11494]
```

In [22]:

```
#Now that we have our attributes and labels, the next step is to split this data into train
#We'll do this by using Scikit-Learn's built-in train_test_split() method:
```

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=0)
X_train=X_train.reshape(-1,1)
X_test=X_test.reshape(-1,1)
'''The above script splits 70% of the data to training set while 30% of the data to test se
The test_size variable is where we actually specify the proportion of test set.'''
```

Out[22]:

```
'The above script splits 70% of the data to training set while 30% of the da
ta to test set.\nThe test_size variable is where we actually specify the pro
portion of test set.'
```

In [23]:

```
#Train the Algorithm
```

```
'''With Scikit-Learn it is extremely straight forward to implement linear regression models
to do is import the LinearRegression class, instantiate it, and call the fit() method along
This is about as simple as it gets when using a machine learning library to train on your d
```

```
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_train, y_train)
```

```
'''The linear regression model basically finds the best value for the intercept and slope,
best fits the data.
To see the value of the intercept and slop calculated by the linear regression algorithm fo
execute the following code.'''
```

```
print("Regression intercept\t",regressor.intercept_) # To retrieve the intercept
print("Regression cofficient\t",regressor.coef_) # retrieving the slope (coefficient of x)
#y=mx+b
#This means that for every one day, the change in the confirmed cases is about approx 107.
```

```
Regression intercept      -2319.7978205731974
Regression cofficient     [106.4427851]
```

In [24]:

```
#Making Predictions
```

```
'''To do predictions, we will use our test data and see how accurately our algorithm predic
```

```
y_pred = regressor.predict(X_test)
```

```
# To compare the actual output values for X_test with the predicted values, execute the fol
```

```
df = pd.DataFrame({'Actual': y_test, 'Predicted': y_pred})
```

```
print('\n',df)
```

	Actual	Predicted
0	1	-2106.912250
1	26	-936.041614
2	2063	3534.555360
3	758	2257.241939
4	3667	4918.311566
5	310	1086.371303
6	963	2683.013079
7	795	2470.127509
8	7303	6195.624987
9	11494	7260.052838
10	9	-1574.698325
11	213	660.600162
12	423	1405.699658
13	3197	4598.983211
14	10	-1468.255540
15	69	-403.827689
16	152	447.714592
17	2377	3853.883715
18	6876	6089.182202
19	795	2363.684724
20	116	234.829022
21	9	-1681.141110
22	2961	4386.097641
23	1456	3002.341434
24	8187	6408.510557
25	2173	3640.998145
26	5501	5663.411062

In [25]:

```
#the predicted percentages are close to the actual ones.
```

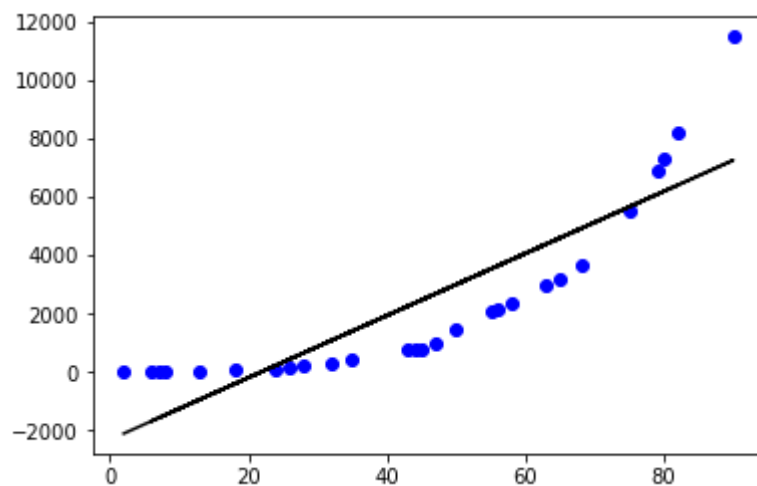
```
print(X_test.shape)
print(y_test.shape)
print(y_pred.shape)
print("\n Prediction V/s Actual \n")
plt.scatter(X_test, y_test, color='b')
plt.plot(X_test, y_pred, color='k')
plt.show()
```

(27, 1)

(27,)

(27,)

Prediction V/s Actual



In [26]:

#Evaluating the Algorithm

```
'''The final step is to evaluate the performance of algorithm.
This step is particularly important to compare how well different algorithms perform on a p
For regression algorithms, three evaluation metrics are commonly used:
    3.1 Mean Absolute Error (MAE) is the mean of the absolute value of the errors.
    3.2 Mean Squared Error (MSE) is the mean of the squared errors.
    3.3 Root Mean Squared Error (RMSE) is the square root of the mean of the squared error.

from sklearn import metrics
print('Mean Absolute Error:', metrics.mean_absolute_error(y_test, y_pred))
print('Mean Squared Error:', metrics.mean_squared_error(y_test, y_pred))
print('Root Mean Squared Error:', np.sqrt(metrics.mean_squared_error(y_test, y_pred)))

df2.describe()

print("""\n\nWe can see that the value of root mean squared error is 1526 , which is
about 50% of the mean value of the percentages of all the Confirmed cases i.e. 2508.40.
This means that our algorithm did a decent job.\n """)
```

```
Mean Absolute Error: 1314.4211134431707
Mean Squared Error: 2329697.2161476873
Root Mean Squared Error: 1526.3345688765905
```

We can see that the value of root mean squared error is 1526 , which is about 50% of the mean value of the percentages of all the Confirmed cases i.e. 2508.40.
This means that our algorithm did a decent job.

In []:

In []: