

Abstract:

The Sars Cov 2 Virus Which cause covid-19 (coronavirus Disease) . Most people got infected with this virus and die. This virus is deadly This virus mutated its body so many times. For scientists, it is tough to make medication and vaccination on this virus. But They finally Made a vaccination on this but if the virus mutated its body the vaccination work successfully or not. We decided to make research on it. By using a Machine learning SVM (Support Vector Machine) Algorithm. This Algorithm set of supervised learning methods is used for classification, regression, and outliers detection. The advantages of support vector machines are Effective in high dimensional spaces. Still effective in cases where several dimensions are greater than the number of samples.

*Keywords:*sars cov 2 virus ,mutation ,vccination,SVM(support vector machine).

Introduction:

The outbreak of Covid-19(coronavirus) has resulted in pandemic infection of over 5.9 million people [1]. This positive-strand RNA Syndrome in humans(Covid-19), With between December 2019 and May 30, 2020 [1] [2].To Combat this pandemic, the World Health Organization (WHO) is coordinating global surveillance,epidemiology.

In this pandemic, the scientists work on vaccination they successfully find the vaccination. But covid-19 Mutation it bodies it tough for them to find vaccination of different variant, let understand How vaccination work.

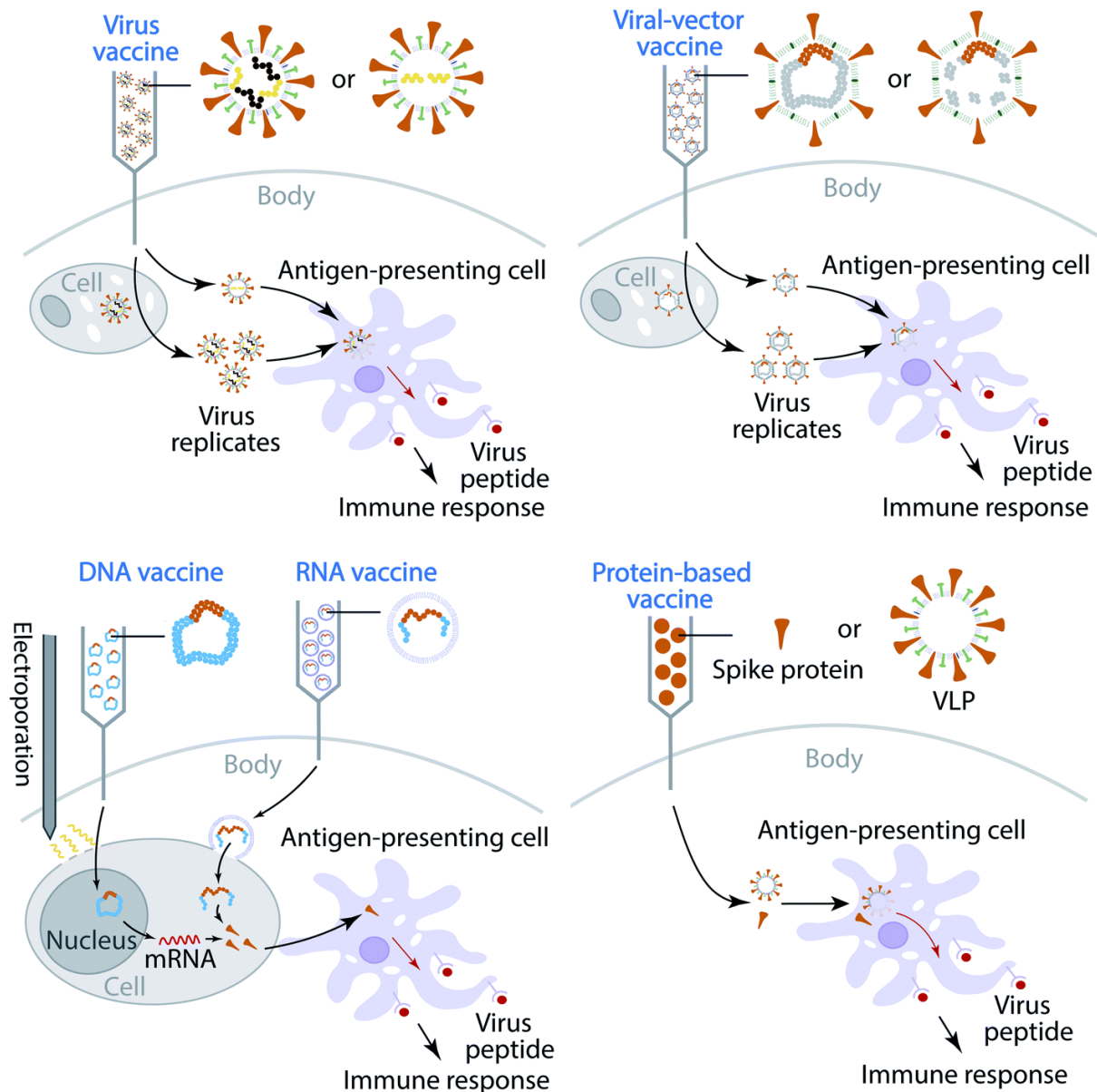


Fig-1[3]:-illustration of four types of COVID-19 vaccine that are currently in development.

There is no doubt it will going improvement in vaccination if the variant and body of the virus are changed. If it is not so many people die in this pandemic situation. With the help of AI and ML, we created a model of how vaccination affects our bodies if the virus mutated.

Our Model:

In this article, we integrated the epidemic prediction model with real global pandemic data and considered the Mutation of the virus and How it affects vaccination. The Dataset is taken from the [<https://www.gavi.org/>]. Let understand how growth of covid-19 affects the countries.

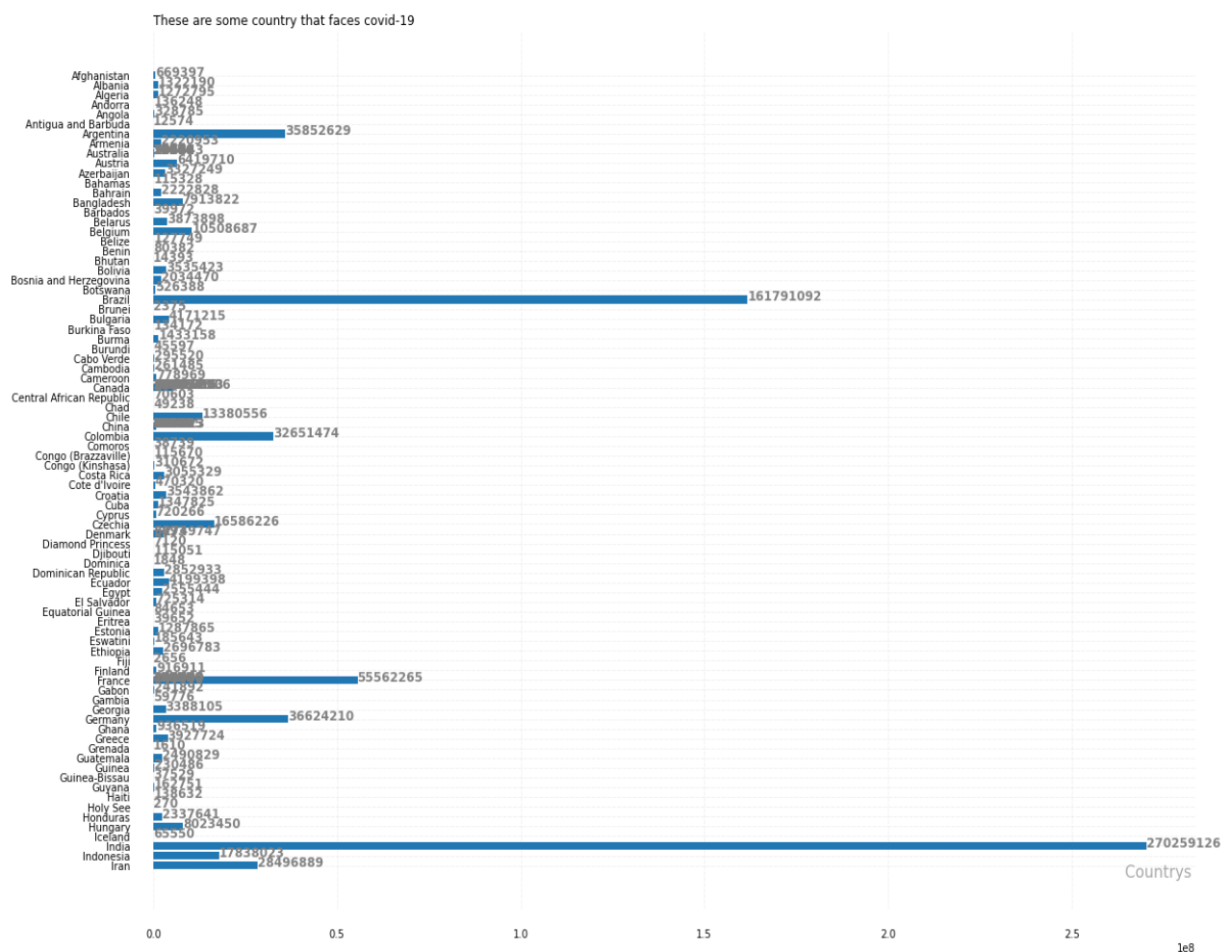
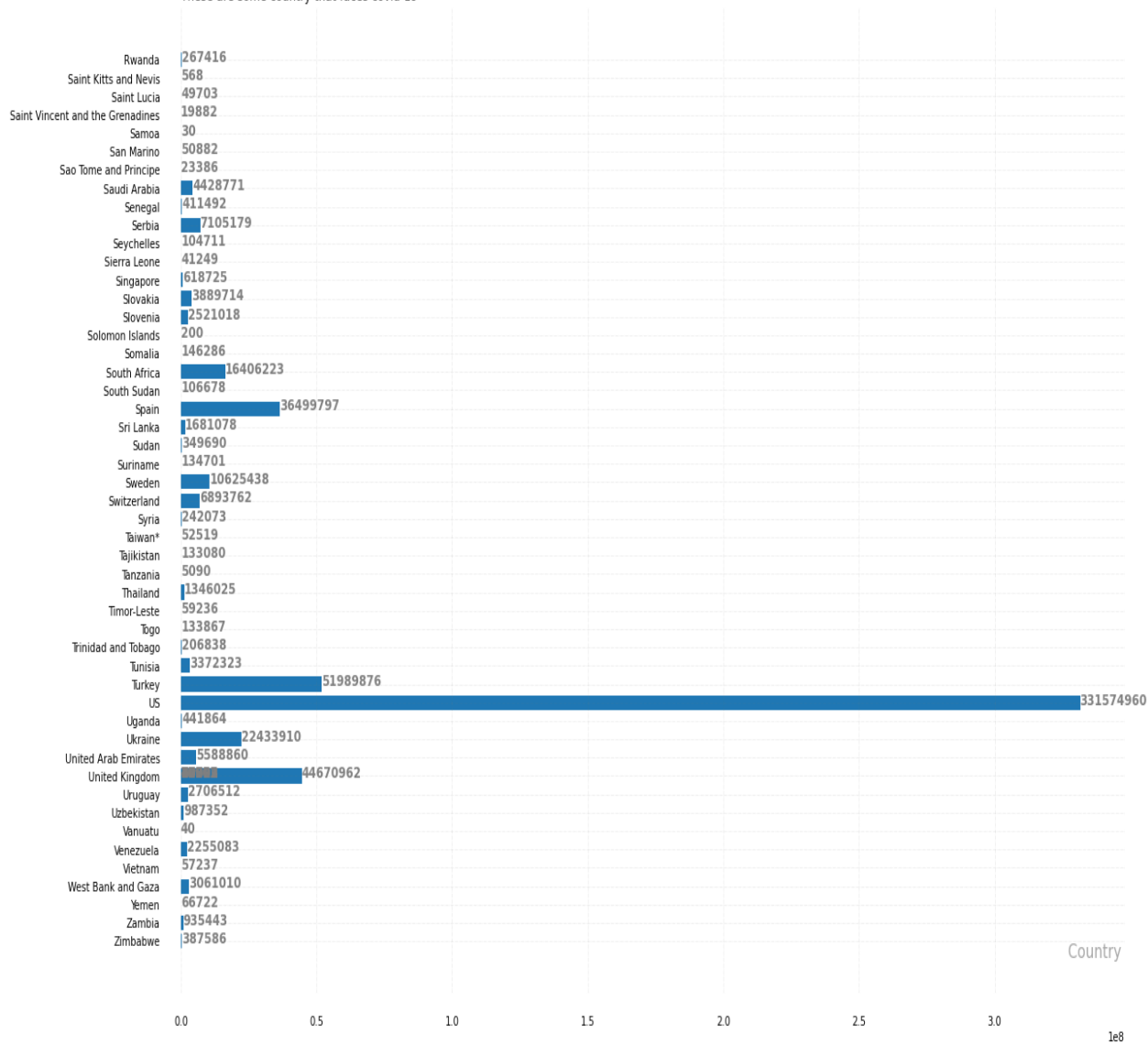


fig-2:-This data is 60 countries that affected due to covid-19

These are some country that faces covid-19



The Mutation of covid-19 and its growth in the third wave

	Actual	3 Waves
184	10.0	2.768527e+03
15	10166.0	2.874462e+04
55	70603.0	1.095083e+05
74	7424.0	7.417171e+03
188	993337.0	8.300916e+05
209	19286567.0	2.177564e+07
92	32651474.0	3.062212e+07
223	4428771.0	7.460642e+06
125	241000.0	1.846669e+05
190	706223.0	5.402453e+05
116	2696783.0	2.537099e+06
218	49703.0	2.761901e+04
244	5090.0	1.350136e+04
101	16586226.0	1.292774e+07
189	5174167.0	6.159557e+06
22	3873898.0	3.528713e+06
59	10934.0	1.782998e+04
156	7303393.0	4.873831e+06
270	2255083.0	2.245193e+06
8	1240.0	6.332910e+03
63	24828.0	1.444058e+04
152	8393650.0	8.922950e+06
194	16093.0	8.335266e+03

200	53869.0	5.631077e+04
12	7498.0	2.300823e+04
173	90.0	2.977163e+03
89	9826.0	2.002178e+04
122	117530.0	1.036090e+05
7	2220953.0	2.475221e+06
215	49564464.0	5.164327e+07
220	30.0	2.787411e+03
108	2852933.0	3.371014e+06
81	6246.0	9.264995e+03
261	42892.0	4.129800e+04
236	1681078.0	8.306415e+05

205	9069769.0	9.555314e+06
240	6893762.0	6.941936e+06
259	40626.0	4.616979e+04
109	4199398.0	4.371052e+06
27	3535423.0	3.699722e+06
118	916911.0	7.602495e+05
178	142481.0	1.282309e+05
187	541057.0	1.393063e+05
224	411492.0	4.570367e+05
37	261485.0	6.308734e+04
64	2936.0	3.026595e+03
154	480384.0	3.320051e+05

Fig 4:-The Mutation of covid-19

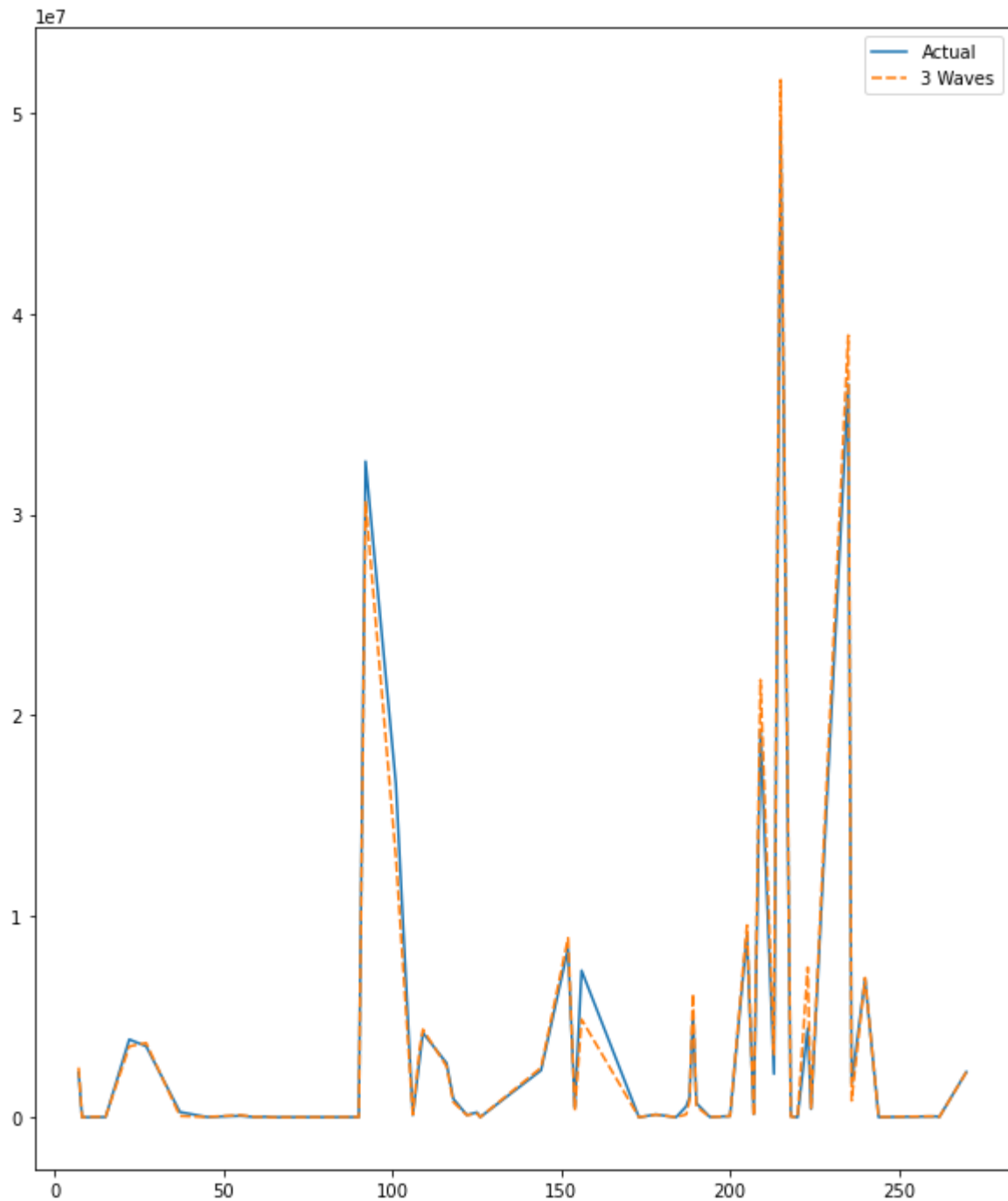


Fig-5 *The prediction growth of covid-19*

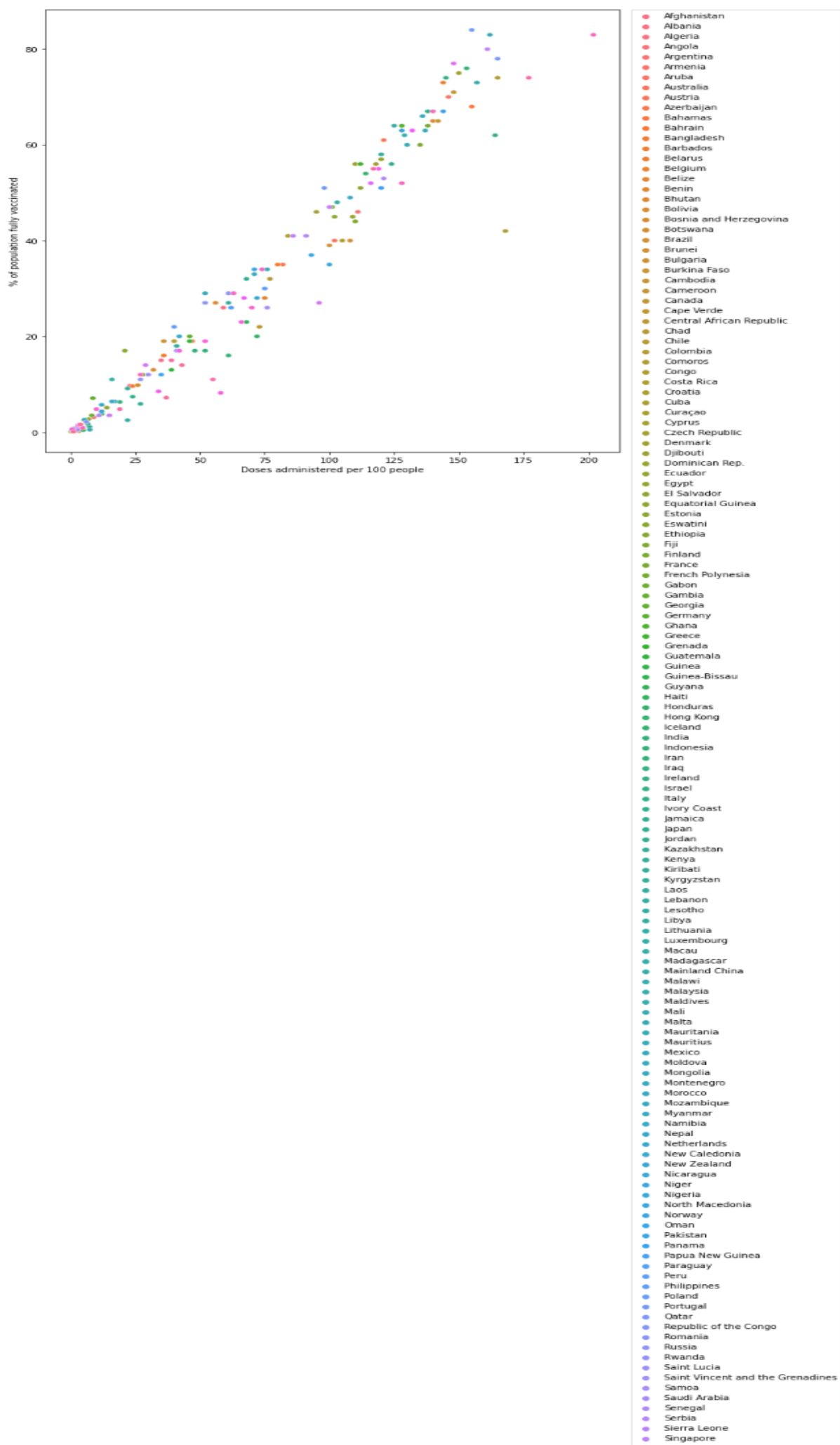
The Vaccination

Vaccines to prevent coronavirus disease 2019 (COVID-19) are perhaps the best hope for ending the pandemic. But as the U.S. Food and Drug Administration (FDA) continues approving or authorizing emergency use of COVID-19 vaccines [4].

	Country	Doses administered per 100 people	Total doses administered	% of population vaccinated	% of population fully vaccinated
0	Afghanistan	8.2	3133227.0	2.0	NaN
1	Albania	59.0	1674093.0	32.0	26.0
2	Algeria	23.0	9989662.0	14.0	9.7
3	Angola	8.9	2820134.0	5.8	3.1
4	Argentina	111.0	50010169.0	65.0	46.0
...
175	Vietnam	37.0	35675840.0	30.0	7.2
176	West Bank & Gaza	43.0	2001407.0	29.0	14.0
177	Yemen	1.1	322934.0	1.1	0.1
178	Zambia	3.8	670361.0	1.7	1.6
179	Zimbabwe	35.0	5166508.0	21.0	15.0

Fig:6 The Vaccination Dose In the world.

As of 29 June 2021, there had been more than 181 million reported infections with SARS-CoV-2 and nearly 4 million reported deaths from COVID-19. In May 2020, the 73rd World Health Assembly issued a resolution recognizing the role of extensive immunization as a global public-health goal for preventing, containing, and stopping transmission of SARS-CoV-2. Globally, there are now more than 125 vaccine candidates, 365 vaccine trials ongoing, and 18 vaccines against COVID-19 approved by at least one country. Published research carried out largely in high-income countries cites concerns about the safety of vaccines against COVID-19, including the rapid pace of vaccine development, as one of the primary reasons for hesitancy, but data from low- and middle-income countries (LMICs) have been limited. In this issue of *Nature Medicine*, Solis Arce et al. present data that begin to address this research gap.[5]



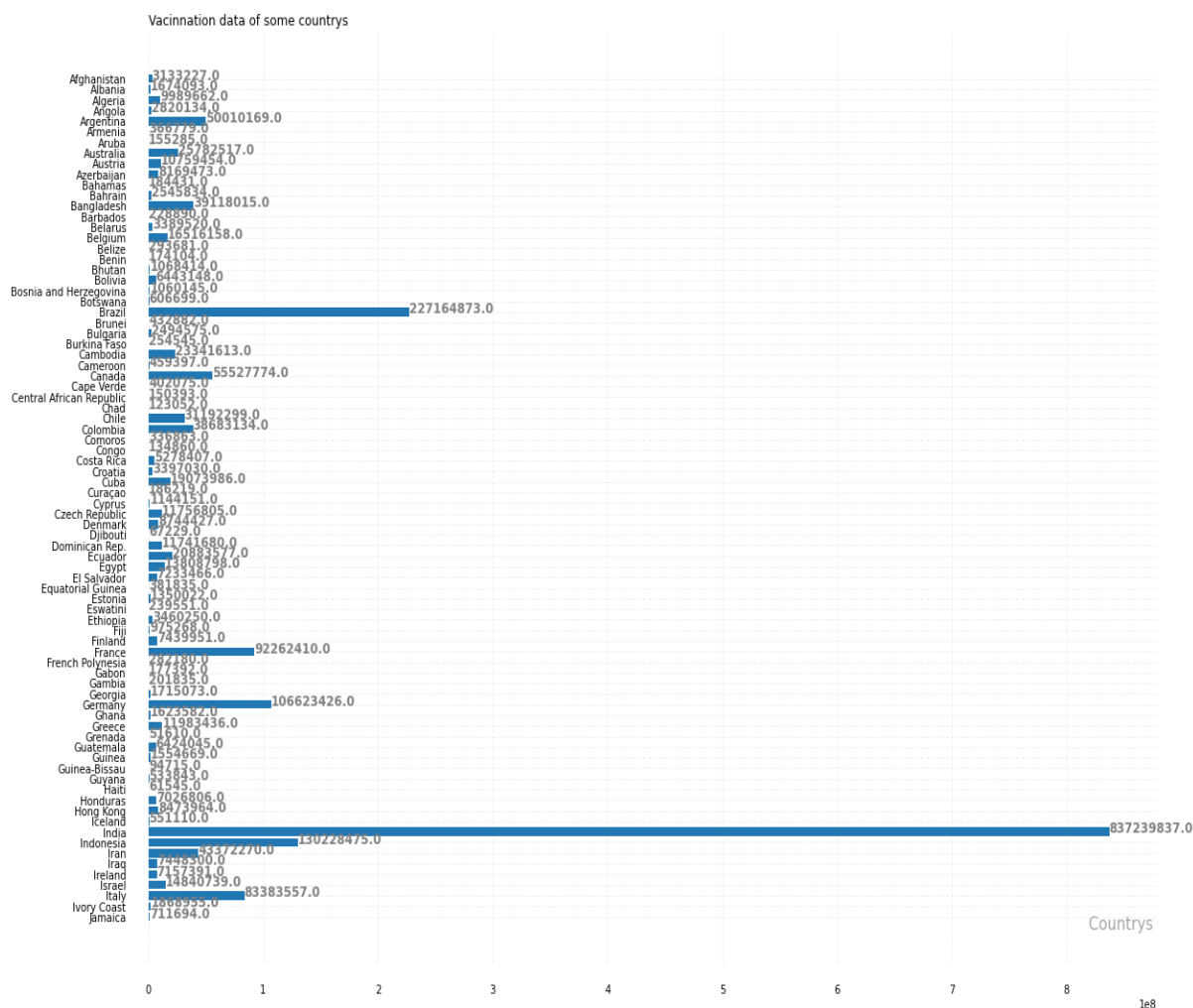


Fig-8 % of the population fully vaccinated

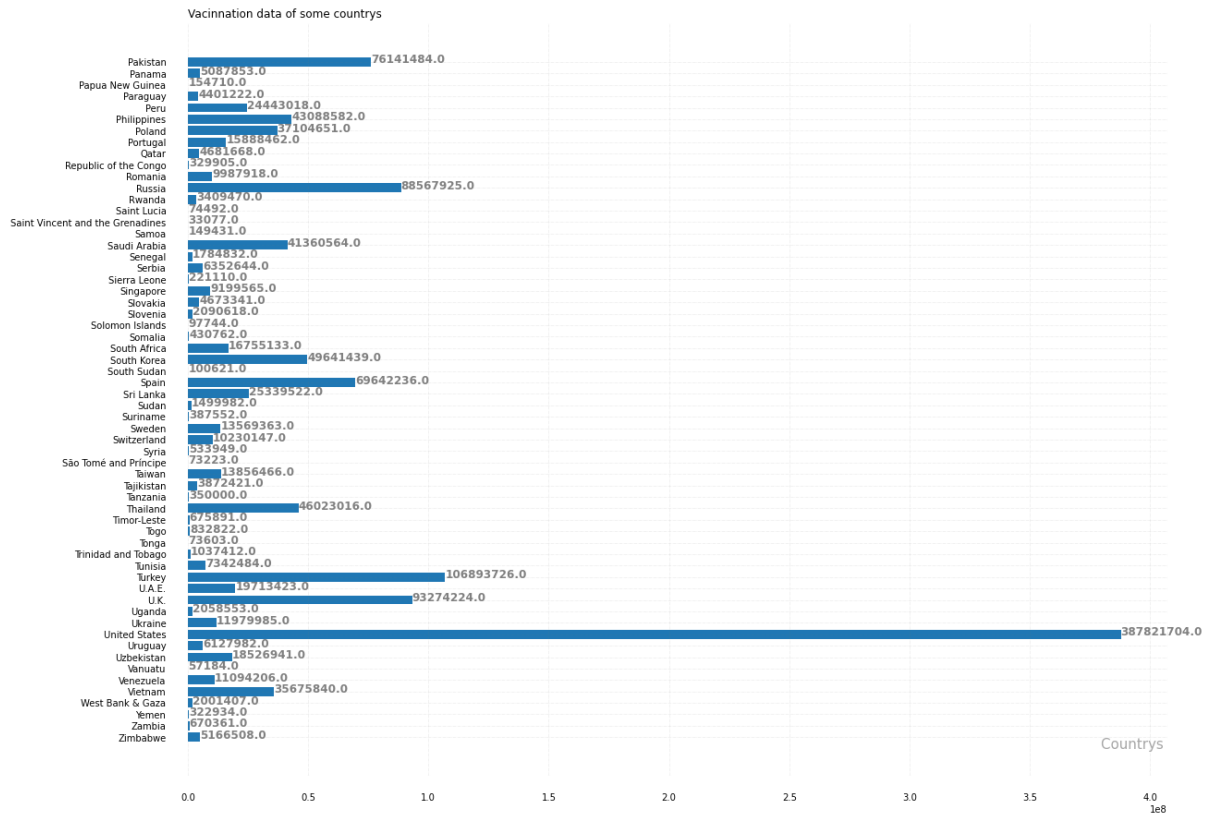


Fig-9 % of the population fully vaccinated

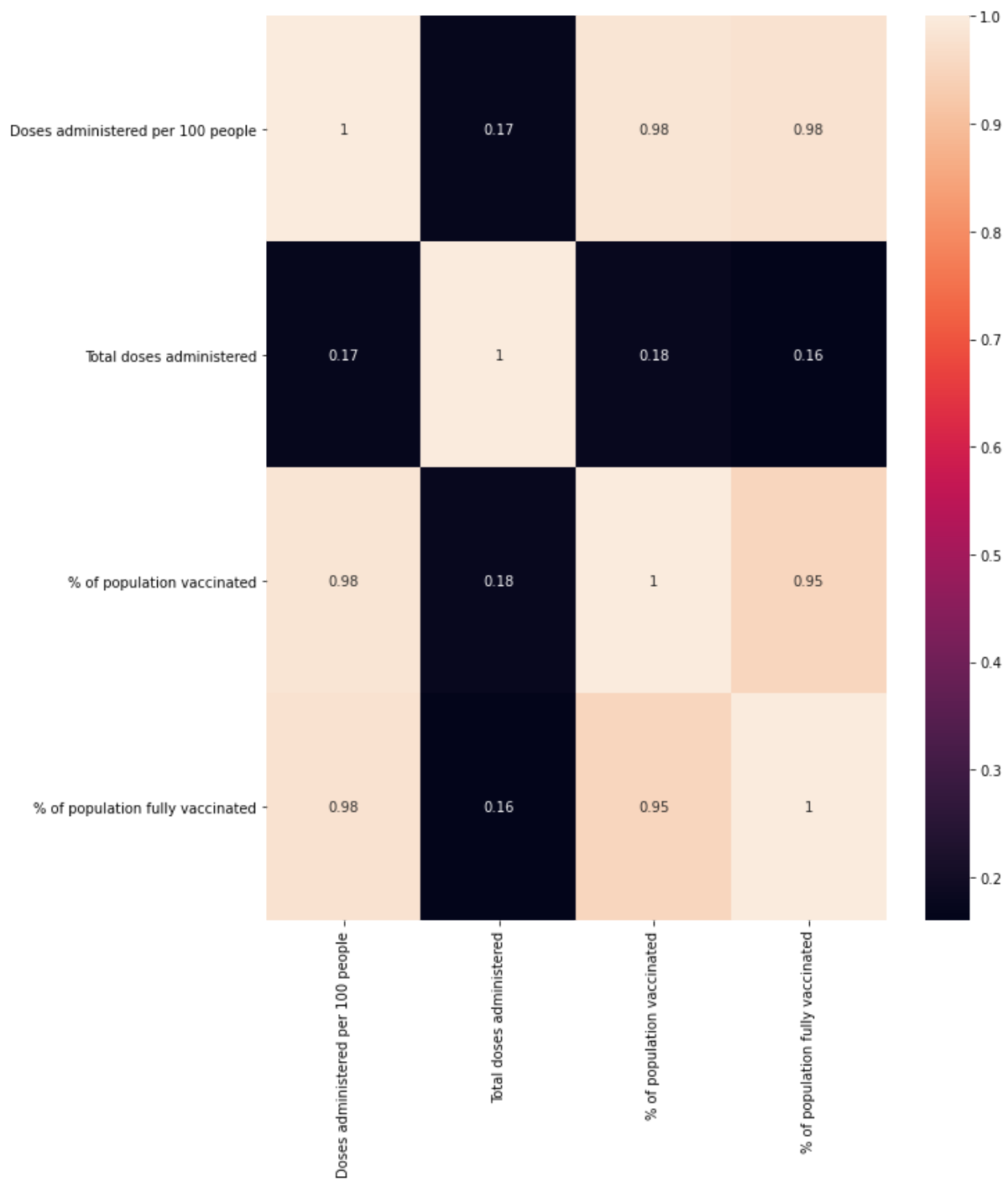


Fig-10 Area divided according to population and Administration

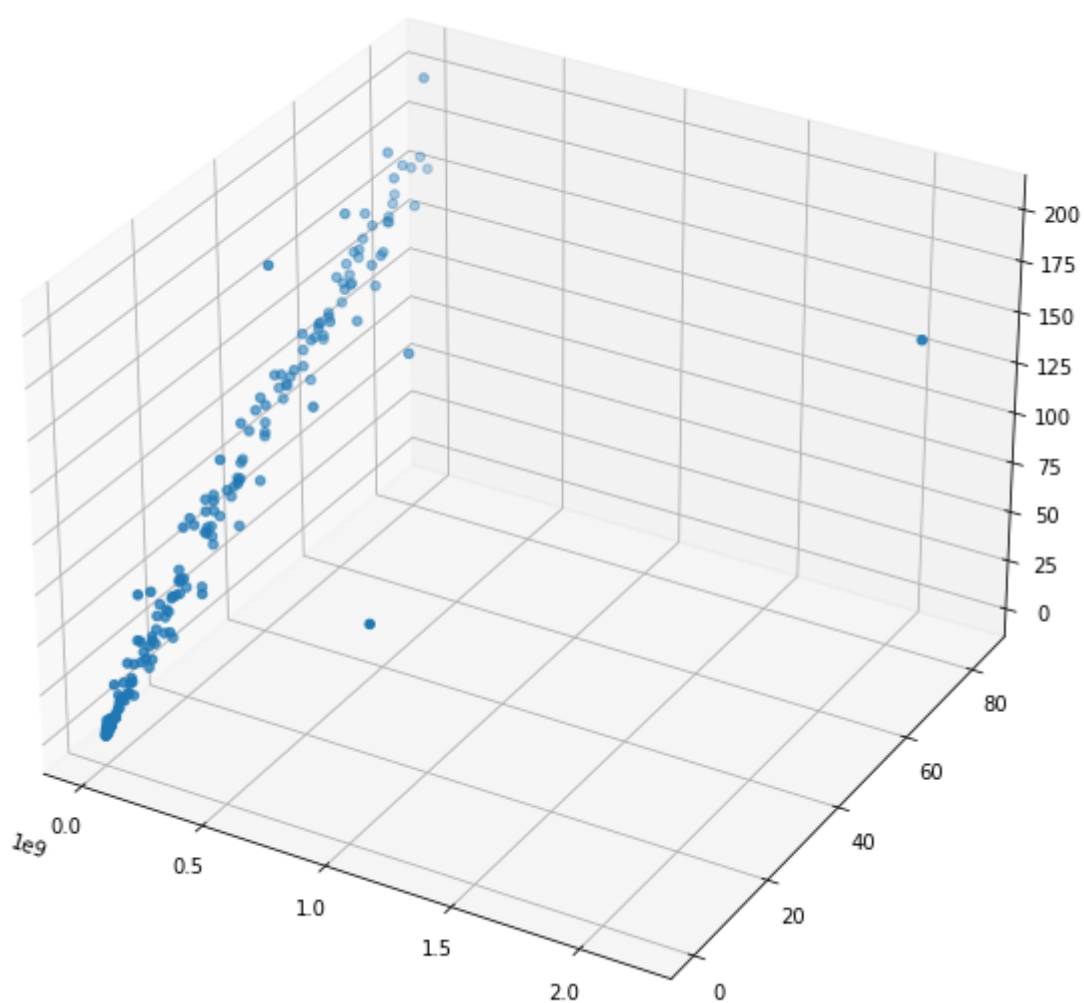


Fig-11 3d graph of % of the population fully vaccinated and total dose of administration

The Mutation of Covid-19 and Vaccination relation

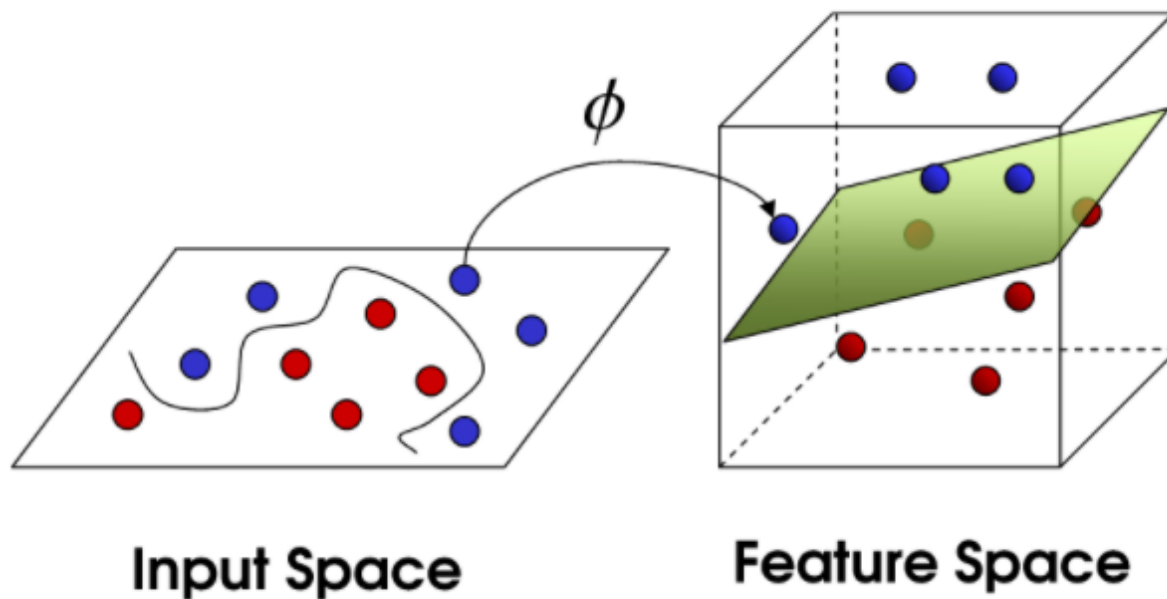


fig-12[6]-How prediction system work between vaccination and mutation in covid-19

```

| df = df.select_dtypes(exclude=['object'])
| df=df.fillna(df.mean())
| X = df.drop('Long',axis=1)
| y = df['Lat']

| from sklearn.model_selection import train_test_split
| X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=0)

| from sklearn.svm import SVR

| svr = SVR(kernel = 'linear',C = 1000)

| from sklearn.preprocessing import StandardScaler
| sc = StandardScaler()
| X_fit = sc.fit(X_test)
| X_std = X_fit.transform(X_test)

| X_std
: array([[ -0.43517467, -0.2024578 , -0.24620888, ..., -0.43534702,
          -0.43534221, -0.43535844],
        [ -0.43409457, -0.2024578 , -0.24620888, ..., -0.43427332,
          -0.43427124, -0.43428982],
        [ -0.42766704, -0.2024578 , -0.24620888, ..., -0.42786284,
          -0.42787493, -0.42790755],
        ...,
        [ -0.40736657, -0.2024578 , -0.24620888, ..., -0.40614056,
          -0.40557852, -0.40504161],
        [ -0.43486319, -0.21288932, -0.22175572, ..., -0.43505716,
          -0.43505716, -0.43505716]])
    
```

```
X_std.sum()
-2.7533531010703882e-14
```

```
type(y_test)
pd.core.frame.DataFrame
y_test.shape
(3, 1)
```

```
(3, 1)
```

```
type(X_test)
pd.core.frame.DataFrame
X_test.shape
(3, 1)
```

```
(3, 1)
```

```
X_train_std = sc.transform(X_train)
X_test_std = sc.transform(X_test)
X_test_std
sc = StandardScaler().fit(X_train)
```

Fig-13 how we implement the SVM

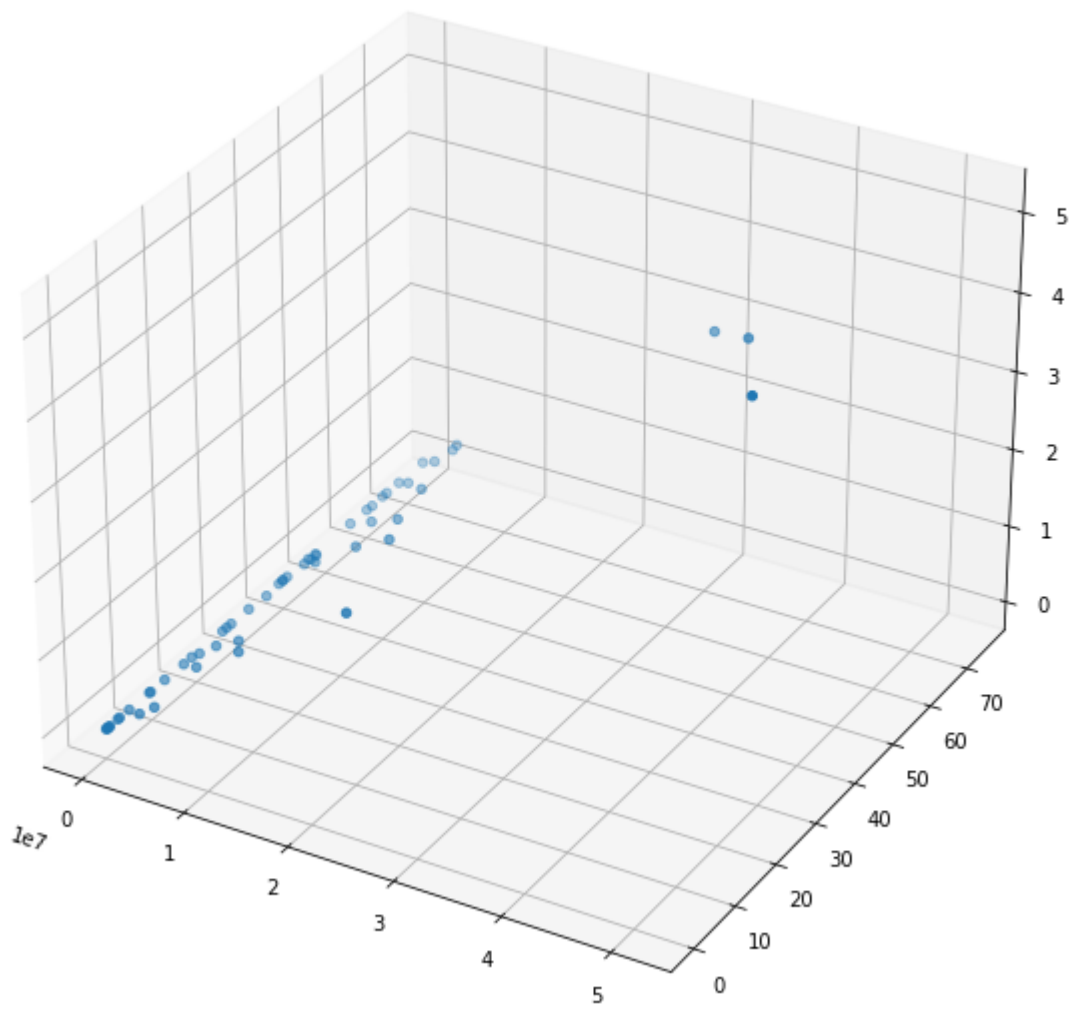


Fig-14 3-d graph the mutation and vaccination of covid-19

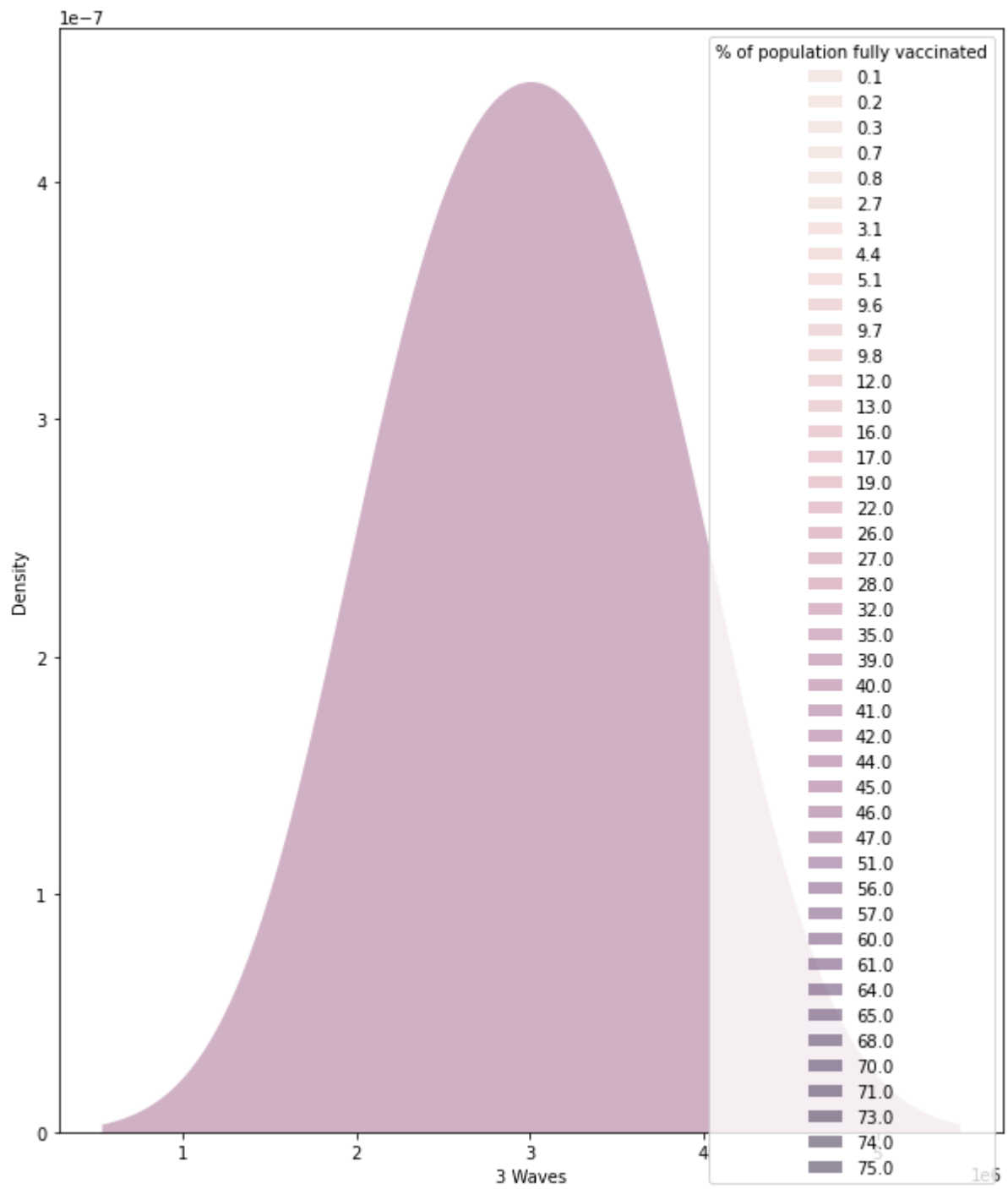


Fig-15 graph the mutation and vaccination of covid-19