COVID Vaccines Analysis

Aim –

This Project mainly aims to find out the trend of the vaccinations around the world for the prevention of the Covid 19 pandemic and how much has been achieved so far.

Introduction –

The COVID 19 pandemic caused due to the Corona virus devastated the world by causing several fatalities around the world. This virus originated in Wuhan, China in 2019 and was later spread throughout the world due to human contact in one way or the other. The disease showed symptoms as basic as mild fever and cold but also caused life threatening symptoms like breathing problems caused by damage to the lungs. As this virus was new to the world and there was no vaccine or cure to it at the initial period there were several deaths around the world. The countries around the world were forced to shut themselves to others in order to avoid the further spread of the virus and people were stuck inside their houses and faced many issues with their finances, mental health etc., and felt like animals in a cage. An effort was made to find a cure or vaccine by several health organizations to bring a stop to this pandemic.

In later stages of 2020 several experimental vaccines were developed and was administered to humans. The efforts were successful as the vaccines were helpful in reducing the affects the virus and even if people were infected, they were not in any life threating situation and escaped the illness having only minor symptoms.

Many countries later developed their own vaccines and also helped other countries without the resources by providing them with vaccines developed.

Problem Statements –

- 1. In this project we have analyzed the top 10 fully vaccinated countries
- 2. We have analyzed the top 5 vaccinated countries
- 3. We have analyzed the top 5 daily vaccinating countries
- 4. We have analyzed the total number of daily vaccinations, people who have fully vaccinated, people who is vaccinated
- 5. We have analyzed year wise daily vaccinating details, fully vaccinated people details, vaccinated people details
- 6. We have analyzed the country wise vaccines and iso code details.

Methodology -

Step 1 –

Data Importing –

In power BI desktop with the help of the get data option import the CSV data which is named as country vaccinations and clicked load option.

Step 2 –

Data Cleaning –

After loading the data and after analyzing the data | understood that there are 86512 rows and 15 columns. And in that some of the columns contained null values I have replaced the null values by 0 with the use of replace functions and started working on the data.

Step 3 –

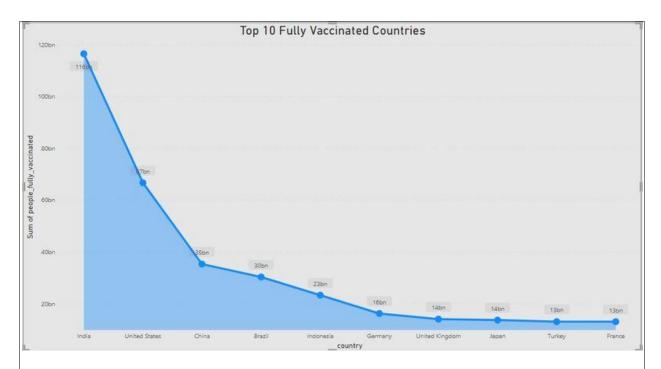
Visualizations –

In visualization part with the help of power BI desktop software I have used different kinds of charts, graphs, cards and table to display the data in the format which will be easy to understand.

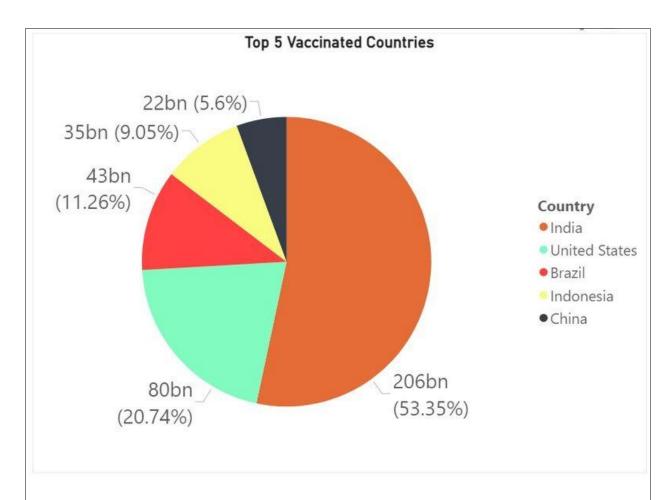
Analysis –

In the analysis part first | have analyzed the top 10 fully vaccinated countries by using area chart and have used the filter option to find the top countries and the result obtained as below,

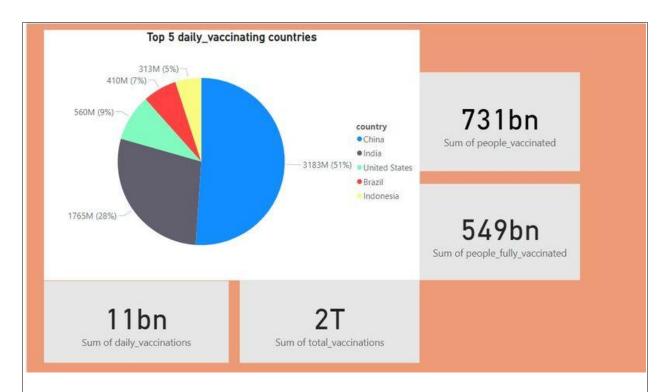
From the below image we can able to come to know that India is the top country in terms full vaccination with 116 billion, followed by united states of America and china with 67 billion and 35 billion respectively.



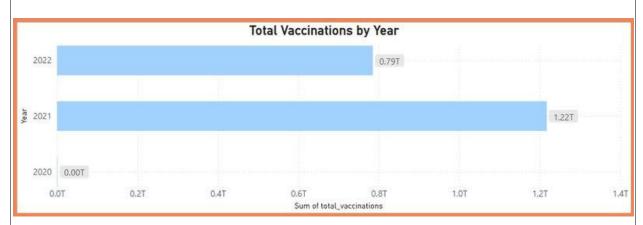
In the second analysis we have analyzed the top 5 vaccinated countries with the help of pie chart and used filter option to find the top countries and with that we came to know that India is the top country with more number of vaccinated peoples followed by United States of America and Brazil.



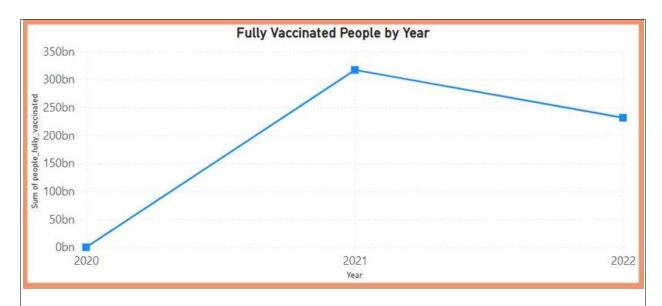
In the third analysis we have analyzed the top 5 country with daily vaccinations with the help of pie chart and used filter option to find the top country with daily vaccinations and with that we came to know that China is the top country with more number of vaccinations followed by India and United States of America.



From the above images we have came to know about the statics of daily vaccination, people who have fully vaccinated and people who is vaccinated and I used cards for this to display the value.



Above image have shown that daily vaccinating details year wise and here we can conclude that 2021 is a year which having maximum daily vaccinating details and I have used bar chart.



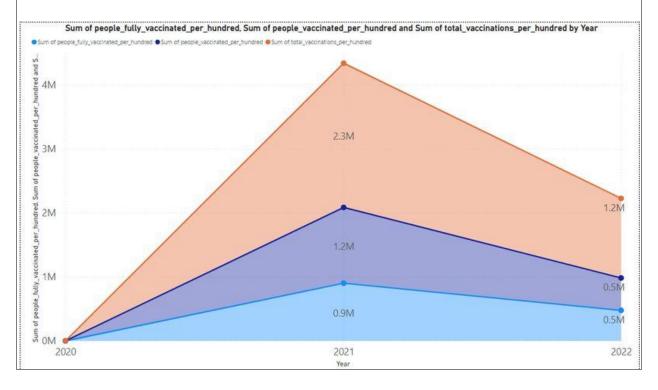
Above image have shown that fully vaccinating details year wise and here we can conclude that 2021 is a year which is having maximum number fully vaccinated peoples and I have used Line chart.

country	vaccines	iso_code
Afghanistan	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing	AFG
Albania	Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik V	ALB
Algeria	Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik V	DZA
Andorra	Moderna, Oxford/AstraZeneca, Pfizer/BioNTech	AND
Angola	Oxford/AstraZeneca	AGO
Anguilla	Oxford/AstraZeneca, Pfizer/BioNTech	AIA
Antigua and Barbuda	Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik V	ATG
Argentina	CanSino, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik V	ARG
Armenia	Moderna, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik V	ARM
Aruba	Pfizer/BioNTech	ABW
Australia	Moderna, Oxford/AstraZeneca, Pfizer/BioNTech	AUS
Austria	Johnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech	AUT
Azerbaijan	Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik V	AZE
Bahamas	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech	BHS
Bahrain	Johnson & Johnson, Moderna, Oxford/Astra Zeneca, Pfizer/Bio NTech, Sinopharm/Beijing, Sputnik Light, Sputnik V	BHR
Bangladesh	Johnson & Johnson, Moderna, Oxford/Astra Zeneca, Pfizer/Bio NTech, Sinopharm/Beijing, Sinovac	BGD
Barbados	Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing	BRB
Belarus	Sinopharm/Beijing, Sputnik V	BLR
Belgium	Johnson & Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech	BEL
Belize	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing	BLZ
Benin	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac	BEN
Bermuda	Oxford/AstraZeneca, Pfizer/BioNTech	BMU
Bhutan	Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing	BTN

From the above chart I have used table to understand the vaccine details and iso_code details country wise.



From the above images we have analyzed the daily vaccinations per million, people who have fully vaccinated per hundred, people who have vaccinated per hundred, total vaccinations per hundred.



From the above image we came to know about people who have vaccinated per hundred, people who have fully vaccinated per hundred, total vaccinated per hundred and we came. to know about that 2021 was the peak year in all the 3 cases.

Insights –

- Here we analyzed the top 10 fully vaccinated countries in which India tops the list which indicates that people in the country where showing lots of interests to get vaccinated.
- And also analyzed top 5 vaccinated countries here also India tops the list.
- And then analyzed top 5 daily vaccinating countries and here China tops the list.
- And also we analyse the sum of daily vaccinating details, fully vaccinating and vaccinating people details.
- And our year wise analyse shows that 2021 was the peak year for every vaccination details.

Recommendations -

- We should collect day to day reports and we should update our records daily to get more accurate details.
- So that we can move forward with more vaccination to the right country which needs the most

Title: Analyzing the Effectiveness and Impact of COVID-19 Vaccination Programs

Abstract: The global response to the COVID-19 pandemic has been characterized by the rapid development and distribution of vaccines. These vaccines have played a crucial role in controlling the spread of the virus, reducing severe cases, and ultimately saving lives. This comprehensive analysis focuses on COVID-19 vaccination programs, offering insights into their development, distribution, effectiveness, and broader societal implications.

Vaccine Development to COVID-19 Vaccines, Vaccine Platforms (mRNA, Viral Vector, Protein Subunit), Mechanisms of Action, Efficacy and Variants, Safety Profiles and Adverse Events

- Vaccine Distribution Global Distribution Challenges, Supply Chain Logistics, Vaccine Hesitancy and Public Perception, Equity in Access, International Collaboration
- ➤ Effectiveness and Impact, Reduction in COVID-19 Cases, Impact on Hospitalizations and Mortality, Long-term Public Health Implications, Emerging Variants and Vaccine Adaptation
- Societal and Economic Implications Economic Recovery, Social Reintegration, Psychological and Mental Health Effects, Vaccine Passports and Privacy Concerns
- Lessons Learned and Future Outlook Key Takeaways from COVID-19 Vaccination Programs, Preparedness for Future Pandemics, Ethical Considerations, Conclusion and Policy Recommendations

This analysis provides a comprehensive overview of the development, distribution, effectiveness, and societal impacts of COVID-19 vaccines, offering valuable insights for policymakers, healthcare professionals, and the general public as we continue to navigate the evolving landscape of the pandemic.

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import os
for dirname, _, filenames in os.walk('/input'):
    for filename in filenames:
        print(os.path.join(dirname, filename))

/opt/conda/lib/python3.10/site-packages/scipy/__init__.py:146: UserWarning: A Num
Py version >=1.16.5 and <1.23.0 is required for this version of SciPy (detected v
ersion 1.23.5</pre>
```

warnings.warn(f"A NumPy version >={np_minversion} and <{np_maxversion}"

/input/covid-world-vaccination-progress/country_vaccinations_by_manufacturer.csv

/input/covid-world-vaccination-progress/country_vaccinations.csv

In [2]:

data = pd.read_csv("/input/covid-world-vaccination-progress/country_vaccinations.csv")</pre>

data = pd.read_csv(/input/covid-world-vaccination-progress/country_vaccinations.csv)
data.head()

Out[2]:

1	co un try	is o - c o d e	d a t e	total _va ccin atio ns	peo ple_ vacc inat ed	peopl e_full y_vac cinate d	daily _vacc inatio ns_ra w	dail y_v acci nati ons	total_v accinati ons_per _hundr ed	people_ vaccina ted_per _hundr ed	people_f ully_vac cinated_ per_hund red	daily_v accinati ons_pe r_milli on	vacc ines	so urc e_ na me	sourc e_we bsite
	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 2	0.0	0.0	NaN	NaN	Na N	0.0	0.0	NaN	NaN	John son &Jo hnso n, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
1 3 1 3	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 3	Na N	Na N	NaN	NaN	136 7.0	NaN	NaN	NaN	34.0	John son & Johnson, Oxfo rd/A straZ enec a, Pfize	W orl d He alt h Or ga niz ati	https: //covi d19. who.i nt/

													r/Bi	on	
2	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 4	Na N	Na N	NaN	NaN	136 7.0	NaN	NaN	NaN	34.0	John son & Johnson, Oxford/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
3	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 5	Na N	Na N	NaN	NaN	136 7.0	NaN	NaN	NaN	34.0	John son & Johnson, Oxford/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
4	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 6	Na N	Na N	NaN	NaN	136 7.0	NaN	NaN	NaN	34.0	John son & Johnson, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/

In [3]:

data.describe()

Out[3]:

	[[3]:			1					
	total_v accinat ions	people _vacci nated	people_fu lly_vacci nated	daily_vac cinations _raw	daily_v accinat ions	total_vaccin ations_per_h undred	people_vacci nated_per_h undred	people_fully_v accinated_per_ hundred	daily_vaccin ations_per_ million
c o u nt	4.3607 00e+0 4	4.1294 00e+04	3.880200 e+04	3.536200 e+04	8.6213 00e+04	43607.00000	41294.00000	38802.000000	86213.0000 00
m ea n	4.5929 64e+0 7	1.7705 08e+07	1.413830 e+07	2.705996 e+05	1.3130 55e+05	80.188543	40.927317	35.523243	3257.04915 7
st d	2.2460 04e+0 8	7.0787 31e+07	5.713920 e+07	1.212427 e+06	7.6823 88e+05	67.913577	29.290759	28.376252	3934.31244 0
m in	0.0000 00e+0 0	0.0000 00e+00	1.000000 e+00	0.000000 e+00	0.0000 00e+00	0.000000	0.000000	0.000000	0.000000
2 5 %	5.2641 00e+0 5	3.4946 42e+05	2.439622 e+05	4.668000 e+03	9.0000 00e+02	16.050000	11.370000	7.020000	636.000000
5 0 %	3.5900 96e+0 6	2.1873 10e+06	1.722140 e+06	2.530900 e+04	7.3430 00e+03	67.520000	41.435000	31.750000	2050.00000
7 5 %	1.7012 30e+0 7	9.1525 20e+06	7.559870 e+06	1.234925 e+05	4.4098 00e+04	132.735000	67.910000	62.080000	4682.00000 0

m a x	3.2631 29e+0 9	1.2755 41e+09	1.240777 e+09	2.474100 e+07	2.2424 29e+07	345.370000	124.760000	122.370000	117497.000 000	
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In [4]:

pd.to_datetime(data.date)

data.country.value_counts()

Out[4]:

Norway	482
Latvia	480
Denmark	476
United States	471
Russia	470
	• • •
Bonaire Sint Eustatius and Saba	146
Tokelau	114
Saint Helena	92
Pitcairn	85
Falkland Islands	67

Name: country, Length: 223, dtype: int64

In [5]:

```
data = data[data.country.apply(lambda x: x not in ["England", "Scotland", "Wales", "N
orthern Ireland"])]
```

data.country.value_counts()

Out[5]:

Norway	482
Latvia	480
Denmark	476
United States	471
Canada	470
	• • •
Bonaire Sint Eustatius and Saba	146
Tokelau	114
Saint Helena	92
Pitcairn	85
Falkland Islands	67
Names according Langeths 210 dt., acc	:

Name: country, Length: 219, dtype: int64

In [6]:

data.vaccines.value_counts()

Out[6]:

Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech 7608

Oxford/AstraZeneca 6022

Oxford/AstraZeneca, Pfizer/BioNTech 4629

Moderna, Oxford/AstraZeneca, Pfizer/BioNTech 4491

Johnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech 3564

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. . .
```

```
Johnson&Johnson, Oxford/AstraZeneca, Sinovac
    312

Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik V
    311

Johnson&Johnson, Moderna
    251

Johnson&Johnson, Pfizer/BioNTech, Sinopharm/Beijing
    228

EpiVacCorona, Oxford/AstraZeneca, QazVac, Sinopharm/Beijing, Sputnik V, ZF2001
    190

Name: vaccines, Length: 84, dtype: int64

In [7]:

df = data[["vaccines", "country"]]

df.head()
```

Out[7]:

	vaccines	country
0	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi	Afghanistan
1	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi	Afghanistan
2	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi	Afghanistan
3	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi	Afghanistan

4

```
In [8]:
dict_ = {}
for i in df.vaccines.unique():
 dict_[i] = [df["country"][j] for j in df[df["vaccines"]==i].index]
vaccines = {}
for key, value in dict_.items():
 vaccines[key] = set(value)
for i, j in vaccines.items():
 print(f"{i}:>>{j}")
Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing:>>{'Camer
oon', 'Afghanistan', 'Belize', 'Namibia', 'Trinidad and Tobago'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik V:>>{'Oman', 'Bosnia and He
rzegovina', 'Albania', 'Azerbaijan'}
Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik V:>>{'Algeria', 'Zimbabwe
'}
Moderna, Oxford/AstraZeneca, Pfizer/BioNTech:>>{'Guernsey', 'United Kingdom', 'Fi
ji', 'Sweden', 'Australia', 'Jersey', 'Sint Maarten (Dutch part)', 'Finland', 'An
dorra', 'Japan', 'Isle of Man'}
Oxford/AstraZeneca:>>{'Montserrat', 'Kiribati', 'Saint Helena', 'Saint Vincent an
d the Grenadines', 'Liberia', 'Falkland Islands', 'Solomon Islands', 'Tuvalu', 'V
anuatu', 'Democratic Republic of Congo', 'Pitcairn', 'Mali', 'Papua New Guinea',
'Nigeria', 'Samoa', 'Nauru', 'Togo', 'Angola', 'Tonga', 'Sao Tome and Principe'}
Oxford/AstraZeneca, Pfizer/BioNTech:>>{ 'Anguilla', 'Saudi Arabia', 'Gibraltar', '
Saint Lucia', 'Cayman Islands', 'New Zealand', 'Saint Kitts and Nevis', 'Panama',
 'Bermuda', 'Costa Rica', 'Kosovo'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik V:>>{'Antigua and Barbuda'}
```

```
CanSino, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik
V:>>{'Argentina'}
Moderna, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik V:>>{'Armenia'}
Pfizer/BioNTech:>>{'Monaco', 'Tokelau', 'Cook Islands', 'New Caledonia', 'Turks a
nd Caicos Islands', 'Niue', 'Aruba'}
Johnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech:>>{'Czechi
a', 'Slovenia', 'Netherlands', 'Germany', 'Austria', 'South Korea', 'Lithuania',
'Italy'}
Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech:>>{'Bahamas', 'Eswatini', 'G
renada'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,
 Sputnik Light, Sputnik V:>>{'Bahrain'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,
 Sinovac:>>{'Bangladesh'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing:>>{'Maldives', 'Peru', 'Su
riname', 'Barbados', 'Dominica'}
Sinopharm/Beijing, Sputnik V:>>{'Belarus', 'Kyrgyzstan'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech:>>{'Cyprus', 'Portu
gal', 'Iceland', 'Malta', 'Belgium', 'Croatia', 'Jamaica', 'Luxembourg', 'Poland',
 'France', 'Greece', 'Spain', 'Romania', 'Bulgaria', 'Estonia', 'Ireland', 'Canad
a'}
Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac:>>{ 'Benin', 'Brazil
'}
Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing:>>{'Cape Verde',
'Bhutan'}
Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik
V:>>{'Moldova', "Cote d'Ivoire", 'Morocco', 'Bolivia'}
Moderna, Pfizer/BioNTech:>>{'Faeroe Islands', 'Norway', 'Bonaire Sint Eustatius a
nd Saba', 'Curacao', 'Qatar', 'Israel'}
Covaxin, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinova
c:>>{'Botswana'}
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Johnson&Johnson, Oxford/AstraZeneca:>>{'British Virgin Islands', 'South Sudan', '
Malawi'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijin
g:>>{'Nepal', 'Brunei', 'Kenya', 'Kuwait'}
Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing:>>{'Gambia', 'Mozambique',
 'Madagascar', 'Senegal', 'Lesotho', 'Zambia', 'Burkina Faso'}
Sinopharm/Beijing:>>{'Equatorial Guinea', 'Burundi', 'Chad'}
Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac:>>{'Somalia', 'Ca
mbodia'}
Covaxin, Oxford/AstraZeneca:>>{'Central African Republic'}
CanSino, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac:>>{'Chile', 'Ecuador'}
CanSino, Sinopharm/Beijing, Sinopharm/Wuhan, Sinovac, ZF2001:>>{'China'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac:>>{'Uganda
', 'Ukraine', 'Colombia'}
Covaxin, Oxford/AstraZeneca, Sinopharm/Beijing:>>{'Mauritius', 'Comoros'}
Moderna, Oxford/AstraZeneca, Sinopharm/Beijing, Sputnik V:>>{'Congo'}
Abdala, Soberana Plus, Soberana02:>>{'Cuba'}
Johnson&Johnson, Moderna, Pfizer/BioNTech:>>{'United States', 'Liechtenstein', 'D
enmark', 'Switzerland'}
Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac,
Sputnik V:>>{'Egypt', 'Djibouti', 'Guinea'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac:>>{'Dominican Rep
ublic', 'Georgia', 'El Salvador'}
Covaxin, Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac:>>{'Ethi
opia'}
Johnson&Johnson, Pfizer/BioNTech:>>{'South Africa', 'French Polynesia'}
Pfizer/BioNTech, Sinopharm/Beijing, Sputnik V:>>{'Gabon'}
Oxford/AstraZeneca, Sputnik V:>>{'Ghana'}
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Moderna:>>{'Greenland', 'Wallis and Futuna'}
Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik V:>>{'Guatemala'}
Oxford/AstraZeneca, Sinopharm/Beijing:>>{'Niger', 'Myanmar', 'Mauritania', 'Sierr
a Leone', 'Guinea-Bissau'}
Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik V:>>{'Sr
i Lanka', 'Guyana'}
Johnson&Johnson, Moderna:>>{'Haiti'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik V:>>{'Hond
uras'}
Pfizer/BioNTech, Sinovac:>>{'Hong Kong'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,
Sputnik V:>>{'Hungary', 'Jordan'}
Covaxin, Oxford/AstraZeneca, Sputnik V:>>{'India'}
Johnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm
/Beijing, Sinovac:>>{'Indonesia'}
COVIran Barekat, Covaxin, FAKHRAVAC, Oxford/AstraZeneca, Razi Cov Pars, Sinopharm
/Beijing, Soberana02, SpikoGen, Sputnik V:>>{'Iran'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik V:>>{'Lebanon', '
Iraq', 'Montenegro', 'Mongolia', 'Serbia'}
QazVac, Sinopharm/Beijing, Sputnik V:>>{'Kazakhstan'}
Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac,
Sputnik Light, Sputnik V:>>{'Laos'}
Johnson&Johnson, Moderna, Novavax, Pfizer/BioNTech:>>{'Latvia'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik V:>>{'No
rth Macedonia', 'Libya'}
Pfizer/BioNTech, Sinopharm/Beijing:>>{'Macao'}
CanSino, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac:>>{'Mala
ysia'}
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CanSino, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac,
Sputnik V:>>{'Mexico'}
Abdala, Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Soberana02, Sputnik
Light, Sputnik V:>>{'Nicaragua'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac:>>{'Northern Cyprus', 'Timor', 'Urug
uay'}
CanSino, Covaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,
 Sinovac, Sputnik V:>>{'Pakistan'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,
 Sinovac, Sputnik Light, Sputnik V:>>{'Palestine', 'Philippines'}
Covaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac,
 Sputnik V:>>{'Paraguay'}
EpiVacCorona, Sputnik V:>>{'Russia'}
Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,
Sinovac, Sputnik V:>>{'Tunisia', 'Rwanda'}
Pfizer/BioNTech, Sputnik V:>>{'San Marino'}
Oxford/AstraZeneca, Sinopharm/Beijing, Sputnik V:>>{'Seychelles'}
Moderna, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac:>>{'Singapore'}
Johnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik
V:>>{'Slovakia'}
Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinova
c:>>{'Sudan'}
Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik Light, S
putnik V:>>{'Syria'}
Medigen, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech:>>{'Taiwan'}
Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik V:>>{'Tajikistan'}
Johnson&Johnson, Pfizer/BioNTech, Sinopharm/Beijing:>>{'Tanzania'}
Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac:>>{'Thai
land'}
```

```
Pfizer/BioNTech, Sinovac, Turkovac:>>{'Turkey'}
EpiVacCorona, Oxford/AstraZeneca, QazVac, Sinopharm/Beijing, Sputnik V, ZF2001:>>
{'Turkmenistan'}
Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinopharm/Wuhan, Sputnik
V:>>{'United Arab Emirates'}
Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik Light, Sputnik V,
ZF2001:>>{'Uzbekistan'}
Abdala, Sinopharm/Beijing, Sinovac, Soberana02, Sputnik Light, Sputnik V:>>{'Vene
zuela'}
Abdala, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik
V:>>{'Vietnam'}
Johnson&Johnson, Oxford/AstraZeneca, Sinovac:>>{'Yemen'}
In [9]:
import plotly.express as px
import plotly.offline as py
vaccine_map = px.choropleth(data, locations = 'iso_code', color = 'vaccines')
vaccine_map.update_layout(height=300, margin={"r":0,"t":0,"l":0,"b":0})
vaccine_map.show()
```

Machine Learning and Deep Learning Based Time Series Prediction and Forecasting of Ten Nations' COVID-19 Pandemic

- Country this is the country for which the vaccination information is provided;
- Country ISO Code ISO code for the country:

- **Date-** date for the data entry; for some of the dates we have only the daily vaccinations, for others, only the (cumulative) total;
- **Total number of vaccin ations** this is the absolute number of total immunizations in the country;
- **Total number of people vaccinated** a person, depending on the immunization scheme, will receive one or more (typically 2) vaccines; at a certain moment, the number of vaccination might be larger than the number of people;
- Total number of people fully vaccinated this is the number of people that received the entire set of immunization according to the immunization scheme (typically 2); at a certain moment in time, there might be a certain number of people that received one vaccine and another number (smaller) of people that received all vaccines in the scheme;
- **Daily vaccinations (raw)** for a certain data entry, the number of vaccination for that date/country;
- Daily vaccinations for a certain data entry, the number of vaccination for that date/country;
- **Total vaccinations per hundred** ratio (in percent) between vaccination number and total population up to the date in the country;
- **Total number of people vaccinated per hundred** ratio (in percent) between population immunized and total population up to the date in the country;
- Total number of people fully vaccinated per hundred ratio (in percent) between population fully immunized and total population up to the date in the country;
- Number of vaccinations per day number of daily vaccination for that day and country;
- **Daily vaccinations per million** ratio (in ppm) between vaccination number and total population for the current date in the country;
- Vaccines used in the country total number of vaccines used in the country (up to date);
- **Source name** source of the information (national authority, international organization, local organization etc.);
- **Source website** website of the source of information;

Content:

- Missing Data
- <u>Data Visualization</u>

In [1]:

- # This Python 3 environment comes with many helpful analytics libraries installed
- # It is defined by the /python Docker image/docker-python
- # For example, here's several helpful packages to load

```
import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
# Input data files are available in the read-only "../input/" directory
# For example, running this (by clicking run or pressing Shift+Enter) will list all f
iles under the input directory
import matplotlib.pyplot as plt
# plotly
# import plotly.plotly as py
from plotly.offline import init_notebook_mode, iplot, plot
import plotly.express as px
import plotly as py
init_notebook_mode(connected=True)
import plotly.graph_objs as go
from pandas_profiling import ProfileReport
import scipy
# seaborn library
import seaborn as sns
# word cloud library
```

```
from wordcloud import WordCloud
```

```
import os

for dirname, _, filenames in os.walk('/input'):
    for filename in filenames:
        print(os.path.join(dirname, filename))

# You can write up to 206B to the current directory (/working/) that gets preserved a s output when you create a version using "Save & Run All"

# You can also write temporary files to /temp/, but they won't be saved outside of the current session

/input/covid-world-vaccination-progress/country_vaccinations_by_manufacturer.csv
/input/covid-world-vaccination-progress/country_vaccinations.csv

In [2]:

data = pd.read_csv("/input/covid-world-vaccination-progress/country_vaccinations.csv")

data.head()
```

Out[2]:

	co un try	is o — c o d e	d a t e	total _va ccin atio ns	peo ple_ vacc inat ed	peopl e_full y_vac cinate d	daily _vacc inatio ns_ra w	dail y_v acci nati ons	total_v accinati ons_per _hundr ed	people_ vaccina ted_per _hundr ed	people_f ully_vac cinated_ per_hund red	daily_v accinati ons_pe r_milli on	vacc	so urc e_ na me	sourc e_we bsite
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2	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 4	Na N	Na N	NaN	NaN	136 7.0	NaN	NaN	NaN	34.0	John son &Jo hnso n, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
3	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 5	Na N	Na N	NaN	NaN	136 7.0	NaN	NaN	NaN	34.0	John son &Jo hnso n, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati	https: //covi d19. who.i nt/

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4	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 6	Na N	Na N	NaN	NaN	136 7.0	NaN	NaN	NaN	34.0	John son & Jo hnso n, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/

In [3]:

report = ProfileReport(data)

report

Abstract

In the paper, the authors investigated and predicted the future environmental circumstances of a COVID-19 to minimize its efects using artificial intelligence techniques. The experimental investigation of COVID-19 instances has been performed in ten countries, including India, the United States, Russia, Argentina, Brazil, Colombia, Italy, Turkey, Germany, and France

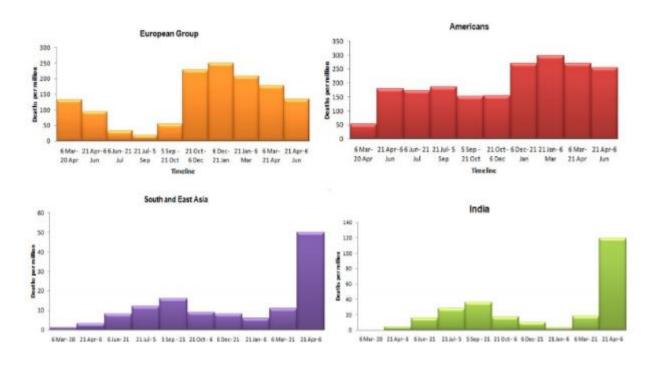
using machine learning, deep learning, and time series models. The confrmed, deceased, and recovered datasets from January 22, 2020, to May 29, 2021, of Novel COVID-19 cases were considered from the Kaggle COVID dataset repository. The country-wise Exploratory Data Analysis visually represents the active, recovered, closed, and death cases from March 2020 to May 2021. The data are pre-processed and scaled using a MinMax scaler to extract and normalize the features to obtain an accurate prediction rate. The proposed methodology employs Random Forest Regressor, Decision Tree Regressor, K Nearest Regressor, Lasso Regression, Linear Regression, Bayesian Regression, Theilsen Regression, Kernel Ridge Regressor, RANSAC Regressor, XG Boost, Elastic Net Regressor, Facebook Prophet Model, Holt Model, Stacked Long Short-Term Memory, and Stacked Gated Recurrent Units to predict active COVID-19 confrmed, death, and recovered cases. Out of diferent machine learning, deep learning, and time series models, Random Forest Regressor, Facebook Prophet, and Stacked LSTM outperformed to predict the best results for COVID-19 instances with the lowest root-mean-square and highest *R*₂ score values.

Keywords COVID-19 · Prediction · XG Boost · Facebook Prophet · Holt model · Stacked gated recurrent units · RANSAC regressor · Random forest regressor · Stacked long short-term memory

Introduction

Throughout history, the world has confronted several major pandemic and epidemic problems. The frst recorded pandemic occurred in Athens during the Peloponnesian War in 430 BC, followed by the Antonine Plague in 165 A.D., in 250 A.D.—the Cyprian Plague, in 541 A.D.—the Justinian Plague, in the eleventh century—leprosy, in 1350—The Black Death, in 1492—The Columbian Exchange, in 1665— The Great Plague of London, in 1817—The First Cholera Pandemic, in 1855—The Third Plague Pandemic, in 1875— Fiji Measles Pandemic, in 1889—Russian Flu, in 1918— Spanish Flu, in 1957—Asian Flu, in 1981—HIV/AIDS, in 2003-SARS, and 2019—COVID-19 [1]. While still a public health concern, Coronavirus 19 (also known as COVID-19) is an infectious sickness that occurred by the severe acute respiratory syndrome coronavirus 2. The frst recorded case of SARS (severe acute respiratory syndrome) was identifed in December of 2019 in Wuhan, China. The disease has since spread to many other nations and healthcare systems worldwide. At the same time, humans inhale contaminated air, including airborne droplets and particles that are smaller than 0.1 microns, and COVID-19 spreads [2]. Inhalation of these particles is more dangerous when people are closely packed together; nevertheless, they can be inhaled further apart, especially indoors. Infected fuids sprayed on the skin, in the eyes, nose, or mouth, or on surfaces contaminated with them may result in transmission. Someone can carry and spread the disease for up to 20 days even if they have no symptoms. During COVID19, afrstwavebeganinthe spring, which

receded significantly throughout the summer, and a second wave appeared in the fall of 2020. The initial wave of the epidemic devastated several nations, and many patients perished. The severity of this early phase was exacerbated by a lack of specialist equipment and a lack of understanding of the disease [4]. We all learned from our mistakes during the frst wave of the pandemic, and as a result, our confidence in being able to handle the second wave much better was strong. Despite this, the second wave had considerably greater infection rates, more patients in ICUs, and, in certain countries, more fatalities [5]. Figure 1 depicts the death rates from March 6, 2020, to June 6, 2021, with Europe and the Americas having the most significant mortality rates compared to India and South and East Asia. Europe had 1,172,912 death cases, the Americas had 1,926,520, South and East Asia had 739,802 death cases, and India had 402,728 COVID death cases as of July



Related Work

Since 2020, researchers have made significant attempts to anticipate the onset of COVID illness in people or the end of the disease around the globe. Keeping this in mind, Shastri et al. [1] suggested a deep learning-based model, such as a recurrent neural network, to forecast the future circumstances of new coronaviruses by studying instances from India and the United States. Ten different nations with the most significant number of verified cases were investigated. It was shown that the predictive accuracy of a range of six separate time series modeling approaches for coronavirus epidemic detection varied by Papastefanopoulos et al. [2]. Using an LSTM model, Chimmula et al. [3] predicted the end of the COVID-19 pandemic and worldwide epidemics due to antiviral drugs and improved access to healthcare. Indicating the date of the pandemic's demise, the writers anticipate that it will be fnished by June of 2020. Using a deep learning model, Togacar et al. [4] identified coronavirus in datasets containing

instances of pneumonia, as well as standard X-ray imaging data. The COVID-19 disease can diagnosedwith 99.27% accuracy with the model that the authors used. COVID-19 drug and vaccine research achievements were evaluated using artifcial intelligence techniques in a recent study by Arshadi et al. [5]. In addition, the scientists gave information about the compounds, peptides, and epitopes in the CoronaDB-Al library, which were discovered both in silico and in vitro. Categorizing chest X-rays into two groups was proposed by the researchers led by Elaziz et al. [6]. The accuracy percentage for the frst and second datasets was 96.09% and 98.09%, respectively. Alimadadi et al. [7] presented a Alaska et al. [9] evaluated the efcacy of deep learning models in predicting COVID-19 illness using laboratory data from 600 patients and got 91.89% accuracy. Their approach was also utilized to help medical professionals validate test data and for clinical prediction research. The Johns Hopkins dashboard data, which were the primary source of the Punn et al.'s [10] research, were utilized with machine learning and deep learning models. The team's goal was to grasp the exponential growth of the COVID-19 and then make predictions about how widespread it may become across the country. Table 1 on the left shows the researchers who worked on the forecast and detection of COVID-19

Contribution Outline

The overall goal of this research is to build models that can calculate two necessary evaluative measures: RMSE and R_2 Score for confrmed, death, and recovered cases from ten different nations to help future forecasts. The steps are as follows:

Step 1: Initially, data are pre-processed to capture char acteristics utilizing various variables, such as active cases, recovered cases, and COVID-19 fatality cases.

Step 2: Exploratory Data Analysis of COVID-19's active cases, closed cases, confrmed cases, recovered cases, and death cases are calculated to summarize or interpret the information that is hidden in rows or columns, and scaling techniques such as Min–Max have been applied to normalize each feature that is obtained from these attributes.

Step 3: Later, utilizing confrmed cases, recovered cases, and death cases from 10 different nations, the gathered data

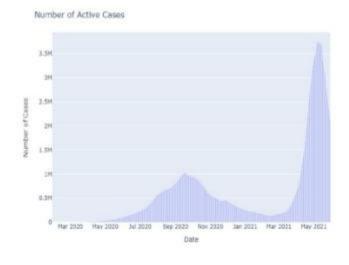
were used to anticipate the future conditions of a new Coro
naVirus. To get the fndings, several machine learning mod
els, time series models, and deep learning models were used,

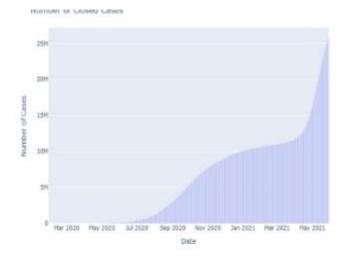
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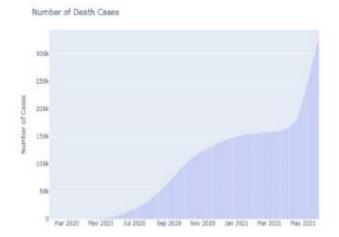
me	Dataset	Technique	Results	Limitations
Ē	1065 CT pathogenic images	Transfer Learning Model, CNN, Graph- Net	Accuracy: 89.5% Specificity: 0.88 Sensitivity: 0.87	Factors such as low signal-to-noise ratio and complex data integration led to reducing the efficacy of deep learning models
hyay et al. [12]	Data collected from WHO (Jan. 16-20,2020)	Long Short-Term Memory, Gated Recurrent Unit	Accuracy: 87%	The model failed to represent the spatio temporal components of the LSTM network
al. [4]	Data collected from Qatar University	Stacking Technique, Fuzzy Color, Deep Learning Model	Classification accuracy: 99.27%	Publications of COVID-19 images were limited. The system did not work with the low resolution and different size input images
Ξ	Dataset was sourced from the Ministry of Health and Family Welfare	Deep Neural Network, Long Short-Term Memory, Recurrent Neural Network, Polynomial Regression	Accuracy ConvLSTM: 98%	The comparative analysis had been per- formed only for two countries
al. [13]	COVID-19 chest X-ray dataset	Bayesian Deep Leaming	Accuracy: 80%	After reviewing the data, it was impossible to conclude anything regarding markers for imaging, discoveries concerning improved diagnosis and therapy for COVID-19
[01]	Data collected from Jan 22, 2020 to Apr 1 2020 at Johns Hopkins University	Support Vector Machine, Deep Neural Network, Long Short-Term Memory, Polynomial Regression	RMSE confirmed: 455.92 Death: 117.94 Recovered: 809.71	The study needed to work on more algorithms to enhance the RMSE score
[6] T	Samples collected from the Albert Einstein Israelite Hospital in Sao Paulo, Brazil	Artificial Neural Network, Convolution Neural Network, Long Short-Term Memory	Accuracy: 86.66% F1 Score: 91.89% Recall: 99.42% AUC: 62.50% Precision: 86.75%	The primary disadvantage of the study was the sheer amount of data. To increase the number of patients for whom the lab findings could not be assessed, the proce- dure was applied on 600 patients
. [14]	180 COVID-19 and 200 chest X-ray images	CNN model, SVM, ResNet50	Accuracy: 91.6%	The study needed to incorporate work on different imagistic patterns of COVID-19
1. [15]	337 patient images from real-world data	Deep learning, nCOVnet	Accuracy: 97.62%	The system worked on a small dataset
[9]	Dataset collected from Joseph Paul Cohen and Paul Morrison Lan Dao	Manta Ray Foraging Optimization, Fractional Multichannel Exponent Moments	Accuracy: 96.09% Accuracy: 98.09%	The system dealt with resource limitations and high CPU time

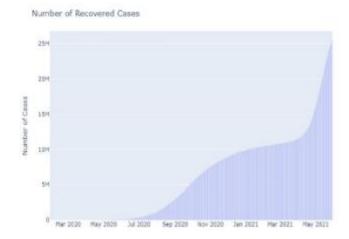
Table 2 Analysis of COVID-19 cases among the top ten countries

Countries	Confirmed cases	Death cases	Recovered cases
India	27,894,800	325,972	25,454,320
USA	33,251,939	594,306	-
Russia	4,995,613	118,781	46,16,422
Argentina	3,732,263	77,108	3,288,467
Brazil	16,471,600	461,057	14,496,224
Colombia	3,363,061	87,747	3,141,549
Italy	4,213,055	126,002	3,845,087
Turkey	33,251,939	47,271	5,094,279
Russia	4,995,613	118,781	4,616,422
Germany	3,684,672	88,413	3,479,700





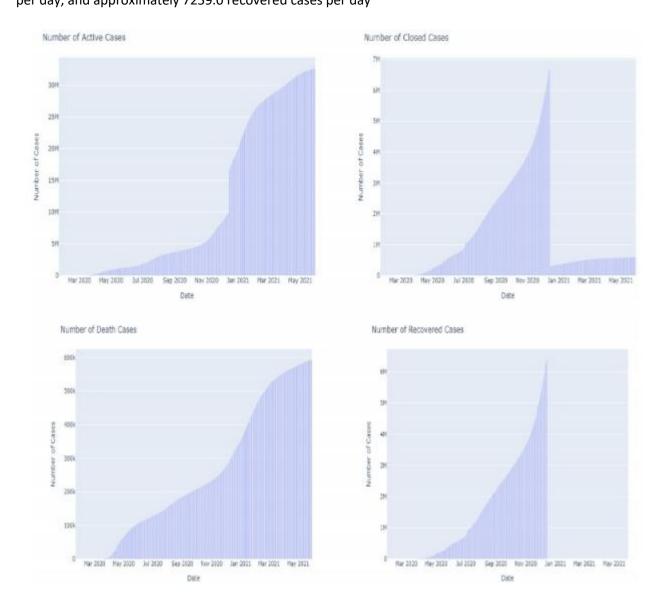




This work employed an exploratory analysis of ten different countries after pre-processing to assess its features via statistical graphs. Figures shown below depicts the graphical analysis of active cases, death cases, closed points, and recovered cases that have been recorded from Jan 2020 to May 2021. It wasdetermined in Fig. 3 that 27,894,800 cases had been confrmed, 2,114,508 were still active, 325,972 had died, 25,780,292 had been closed, and 25,454,320 people had been recovered from Jan 2020 to 29 May 2021. Additionally, the numbers of confrmed cases, deaths, and recovered cases each day were, respectively, 57,397, 671, and 52,375. According to Fig. 4, it has been discovered that US has 3,325,189,940 instances with high certainty, 3,266,576,333 cases with moderate certainty, 594,306 cases with low certainty, and 0 cases with a medium certainty which were seen from January 1st, 2020 to May 29th, 2021. Additionally, the daily average of confrmed cases was reported as 673,128, while the daily average of deaths was recorded as 12,030. Finally, the daily average of recovered cases was recorded as 0. As demonstrated in Fig. 5, the numbers of confrmed, active, and death cases have been as follows: 49,956,313.0, 260,410.0, 118,781.0, 47,352,203.0, and 46,164,322.0 from January 1, 2020 to May 29, 2021. Finally, the total number of confrmed cases was 10,300. The number of death cases was 245, and the total number of recovered cases was 9518. In Fig. 6, it was discovered that Argentina has reported 373,263.0 total cases, with 366,688.0 currently active

cases,77,108currentlyknowndeathcases,336,575 previously known to be closed cases, and 328,467 previously known recovered cases from January 1st, 2020 to May 31st, 2021.

In addition to this, there were around 8239.0 confrmed cases of the disease each day, approximately 170.0 deaths per day, and approximately 7259.0 recovered cases per day



COVID 19 WORLD VACCINATION PROGRESS

The data contains the following information:

- Country this is the country for which the vaccination information is provided;
- Country ISO Code ISO code for the country;
- **Date-** date for the data entry; for some of the dates we have only the daily vaccinations, for others, only the (cumulative) total;
- **Total number of vaccin ations** this is the absolute number of total immunizations in the country;
- **Total number of people vaccinated** a person, depending on the immunization scheme, will receive one or more (typically 2) vaccines; at a certain moment, the number of vaccination might be larger than the number of people;
- Total number of people fully vaccinated this is the number of people that received the entire set of immunization according to the immunization scheme (typically 2); at a certain moment in time, there might be a certain number of people that received one vaccine and another number (smaller) of people that received all vaccines in the scheme;
- **Daily vaccinations (raw)** for a certain data entry, the number of vaccination for that date/country;
- Daily vaccinations for a certain data entry, the number of vaccination for that date/country;
- **Total vaccinations per hundred** ratio (in percent) between vaccination number and total population up to the date in the country;
- **Total number of people vaccinated per hundred** ratio (in percent) between population immunized and total population up to the date in the country;
- Total number of people fully vaccinated per hundred ratio (in percent) between population fully immunized and total population up to the date in the country;
- Number of vaccinations per day number of daily vaccination for that day and country;
- **Daily vaccinations per million** ratio (in ppm) between vaccination number and total population for the current date in the country;
- Vaccines used in the country total number of vaccines used in the country (up to date);
- **Source name** source of the information (national authority, international organization, local organization etc.);
- Source website website of the source of information;

Content:

- Missing Data
- Data Visualization

```
In [1]:
# This Python 3 environment comes with many helpful analytics libraries installed
# It is defined by the /python Docker image: https://github.com/docker-python
# For example, here's several helpful packages to load
import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
# Input data files are available in the read-only "../input/" directory
# For example, running this (by clicking run or pressing Shift+Enter) will list all f
iles under the input directory
import matplotlib.pyplot as plt
# plotly
# import plotly.plotly as py
from plotly.offline import init_notebook_mode, iplot, plot
import plotly.express as px
import plotly as py
init_notebook_mode(connected=True)
import plotly.graph objs as go
from pandas profiling import ProfileReport
import scipy
```

```
# seaborn library
import seaborn as sns
# word cloud library
from wordcloud import WordCloud
import os
for dirname, _, filenames in os.walk('/input'):
    for filename in filenames:
        print(os.path.join(dirname, filename))
# You can write up to 20GB to the current directory (/working/) that gets preserved a
s output when you create a version using "Save & Run All"
# You can also write temporary files to /temp/, but they won't be saved outside of th
e current session
/input/covid-world-vaccination-progress/country_vaccinations_by_manufacturer.csv
/input/covid-world-vaccination-progress/country_vaccinations.csv
In [2]:
data = pd.read_csv("/input/covid-world-vaccination-progress/country_vaccinations.csv")
data.head()
Out[2]:
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                                daily
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In [3]:

report = ProfileReport(data)

report

Summarize dataset: 100%

106/106 [00:26<00:00, 3.82it/s, Completed]

Generate report structure: 100%

1/1 [00:07<00:00, 7.25s/it]

Render HTML: 100%

1/1 [00:03<00:00, 3.96s/it]

Out[3]:

Missing Data

We will fix some shortcomings to make data visualization easier and more understandable.

```
In [4]:
data.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 86512 entries, 0 to 86511
Data columns (total 15 columns):
    Column
                                        Non-Null Count Dtype
    country
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    iso_code
                                        86512 non-null object
 2
    date
                                        86512 non-null object
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    people_vaccinated
                                        41294 non-null float64
 5
    people_fully_vaccinated
                                        38802 non-null float64
    daily_vaccinations_raw
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    daily_vaccinations
                                        86213 non-null float64
 8
    total_vaccinations_per_hundred
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                                       41294 non-null float64
 10 people_fully_vaccinated_per_hundred 38802 non-null float64
 11 daily_vaccinations_per_million
                                        86213 non-null float64
```

12 vaccines	86512	non-null	object
13 source_name	86512	non-null	object
14 source_website	86512	non-null	object
<pre>dtypes: float64(9), object(6)</pre>			
memory usage: 9.9+ MB			
In [5]:			
data.shape			
Out[5]:			
(86512, 15)			
In [6]:			
<pre>data.isna().sum()</pre>			
Out[6]:			
country	0		
iso_code	0		
date	0		
total_vaccinations	42905		
people_vaccinated	45218		
people_fully_vaccinated	47710		
daily_vaccinations_raw	51150		
daily_vaccinations	299		
total_vaccinations_per_hundred	42905		
<pre>people_vaccinated_per_hundred</pre>	45218		
<pre>people_fully_vaccinated_per_hundred</pre>	47710		

daily_vaccinations_per_million	299
vaccines	0
source_name	0
source_website	0
dtype: int64	

As can be seen, there is quite much missing data.

Drop the total_vaccinations column from these deficiencies first

```
In [7]:
data = data.drop(data[data.total_vaccinations.isna()].index)
In [8]:
data.isna().sum()
Out[8]:
country
                                           0
iso_code
date
                                           0
total_vaccinations
                                           0
people_vaccinated
                                        2717
people_fully_vaccinated
                                        5097
daily_vaccinations_raw
                                        8245
daily_vaccinations
                                         223
total_vaccinations_per_hundred
                                           0
people_vaccinated_per_hundred
                                        2717
people_fully_vaccinated_per_hundred
                                        5097
```

daily_vaccinations_per_million 223
vaccines 0
source_name 0
source_website 0

dtype: int64

As you can see the missing data in the total_vaccinations column has been removed.

Now let's remove the missing data from the people_vaccinated column

In [9]:
check_data = data.drop(data[data.people_vaccinated.isna()].index)
In [10]:
check_data.head()

Out[10]:

	co un try	is o - c o d e	d a t e	total _va ccin atio ns	peo ple_ vac cina ted	peopl e_full y_vac cinate d	daily _vacc inatio ns_ra w	dail y_v acci nati ons	total_v accinati ons_pe r_hund red	people_ vaccina ted_per _hundr ed	people_f ully_vac cinated_ per_hund red	daily_v accinati ons_pe r_milli on	vacc ines	so urc e_ na me	sourc e_we bsite
0	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 2	0.0	0.0	NaN	NaN	Na N	0.00	0.00	NaN	NaN	John son &Jo hnso n, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/

6	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 8	820 0.0	820 0.0	NaN	NaN	136 7.0	0.02	0.02	NaN	34.0	John son & Johnson, Oxford/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
2 2	Af gh an ist an	A F G	2 0 2 1 - 0 3 - 1 6	540 00.0	540 00.0	NaN	NaN	286 2.0	0.14	0.14	NaN	72.0	John son & Johnson, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
4 4	Af gh an ist an	A F G	2 0 2 1 - 0 4 - 0 7	120 000. 0	120 000. 0	NaN	NaN	300 0.0	0.30	0.30	NaN	75.0	John son & Johnson, Oxford/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
5 9	Af gh an ist	A F G	2 0 2 1	240 000. 0	240 000. 0	NaN	NaN	800 0.0	0.60	0.60	NaN	201.0	John son &Jo hnso n,	W orl d He alt	https: //covi d19. who.i

an	0					Oxfo	h	nt/
	4					rd/A	Or	
	-					straZ	ga	
	2					enec	niz	
	2					a,	ati	
						Pfize	on	
						r/Bi		

As you can see the missing data in the people_vaccinated column has been removed.

Let's look at the values between the columns by looking at the correlation map

```
In [11]:
plt.subplots(figsize = (10,10))
sns.heatmap(data.corr(), annot = True, square = True)
plt.show()
```

people vaccinated and people vaccinated per hundred

The data of the total_vaccinations column and the people_vaccinated column look almost the same.

As can bee seen from the heatmap, these features have almost ideal correlation.

```
In [12]:
diff = check_data.total_vaccinations.mean() - check_data.people_vaccinated.mean()
diff_per_hundred = check_data.total_vaccinations_per_hundred.mean() - check_data.people_vaccinated_per_hundred.mean()

data.people_vaccinated = data.people_vaccinated.fillna(data.total_vaccinations - diff)
data.people_vaccinated_per_hundred = data.people_vaccinated_per_hundred.fillna(data.total_vaccinations_per_hundred - diff_per_hundred)
```

Let's check if everything ok.

In [13]: data.isna().sum() Out[13]: country 0 iso_code date total_vaccinations people_vaccinated 0 people_fully_vaccinated 5097 daily_vaccinations_raw 8245 daily_vaccinations 223 total_vaccinations_per_hundred people_vaccinated_per_hundred people_fully_vaccinated_per_hundred 5097 daily_vaccinations_per_million 223 vaccines 0 source_name source_website 0 dtype: int64

daily_vaccinations_raw and daily_vaccinations

The data of the daily_vaccinations column and the daily_vaccinations_raw column look almost the same.

As can bee seen from the heatmap, these features have almost ideal correlation.

```
In [14]:
diff = check_data.daily_vaccinations.mean() - check_data.daily_vaccinations_raw.mean()
data.daily_vaccinations_raw = data.daily_vaccinations_raw.fillna(data.daily_vaccinati
ons - diff)
In [15]:
data.isna().sum()
Out[15]:
country
                                           0
iso_code
date
                                           0
total_vaccinations
people_vaccinated
people_fully_vaccinated
                                        5097
daily vaccinations raw
                                         223
daily vaccinations
                                         223
total_vaccinations_per_hundred
people_vaccinated_per_hundred
                                           0
people_fully_vaccinated_per_hundred
                                        5097
daily_vaccinations_per_million
                                         223
vaccines
                                           0
source_name
source_website
                                           0
```

dtype: int64

```
In [16]:
data.daily_vaccinations = data.daily_vaccinations.fillna(0)
data.daily_vaccinations_raw = data.daily_vaccinations_raw.fillna(0)
In [17]:
data.isna().sum()
Out[17]:
country
iso_code
                                           0
date
total_vaccinations
people_vaccinated
                                           0
people_fully_vaccinated
                                        5097
daily_vaccinations_raw
                                           0
daily_vaccinations
total_vaccinations_per_hundred
people_vaccinated_per_hundred
                                           0
people_fully_vaccinated_per_hundred
                                        5097
daily_vaccinations_per_million
                                         223
vaccines
source_name
source_website
                                           0
```

The data of the total_vaccinations column and the people_fully_vaccinated column look almost the same.

dtype: int64

As can bee seen from the heatmap, these features have almost ideal correlation.

people_fully_vaccinated

```
In [18]:
diff = check_data.total_vaccinations.mean() - check_data.people_fully_vaccinated.mean
data.people_fully_vaccinated = data.people_fully_vaccinated.fillna(data.total_vaccina
tions - diff)
In [19]:
data.isna().sum()
Out[19]:
                                           0
country
iso_code
                                           0
date
total_vaccinations
                                           0
people_vaccinated
people_fully_vaccinated
daily_vaccinations_raw
                                           0
daily_vaccinations
total_vaccinations_per_hundred
                                           0
people_vaccinated_per_hundred
people_fully_vaccinated_per_hundred
                                        5097
daily_vaccinations_per_million
                                         223
vaccines
                                           0
```

```
source_name 0
source_website 0
dtype: int64
```

people fully vaccinated per hundred

The data of the total_vaccinations_per_hundred column and the people_fully_vaccinated_per_hundred column look almost the same.

As can bee seen from the heatmap, these features have almost ideal correlation.

```
In [20]:
diff = check_data.total_vaccinations_per_hundred.mean() - check_data.people_fully_vac
cinated_per_hundred.mean()
data.people_fully_vaccinated_per_hundred = data.people_fully_vaccinated_per_hundred.f
illna(data.total vaccinations per hundred - diff)
In [21]:
data.isna().sum()
Out[21]:
country
                                           0
iso_code
                                           0
date
                                           0
total_vaccinations
                                           0
people vaccinated
                                           0
people_fully_vaccinated
                                           0
daily_vaccinations_raw
                                           0
daily_vaccinations
                                           0
total_vaccinations_per_hundred
```

Since there is not much similarity between them and the others in the daily_vaccinations_per_million correlation map, we will assign the value 0 instead of the missing data.

```
In [22]:
data.daily_vaccinations_per_million = data.daily_vaccinations_per_million.fillna(0)
In [23]:
data.isna().sum()
Out[23]:
country
                                         0
iso_code
                                         0
date
                                         0
total_vaccinations
                                         0
people_vaccinated
                                         0
people_fully_vaccinated
                                         0
daily_vaccinations_raw
                                         0
daily_vaccinations
                                         0
total_vaccinations_per_hundred
                                         0
```

```
people_vaccinated_per_hundred 0
people_fully_vaccinated_per_hundred 0
daily_vaccinations_per_million 0
vaccines 0
source_name 0
dtype: int64
```

There is no missing data in our columns.

Our missing data has been cleared.

Everything worked fine

iso code

Let's see if there is any missing data in iso_code

```
In [24]:
data[data.iso_code.isna()].country.unique()
Out[24]:
array([], dtype=object)
```

Thats the iso-codes which are used for these countries: GB-ENG for England, NC for Northern Cyprus, GB-NIR for Northern Ireland, GB-SCT for Scotland, GB-WLS for Wales.

We will fill missing iso-codes with appropriate ones.

```
In [25]:
data[data.country == 'England'] = data[data.country == 'England'].fillna('GB-ENG')
data[data.country == 'Northern Ireland'] == data[data.country == 'Northern Ireland'].fillna('GB-NIR')
data[data.country == 'Scotland'] = data[data.country == 'Scotland'].fillna('GB-SCT')
```

```
data[data.country == 'Wales'] = data[data.country == 'Wales'].fillna('GB-WLS')
data = data.fillna('NC')
In [26]:
data.isna().sum()
Out[26]:
country
                                        0
iso_code
                                        0
date
total_vaccinations
                                        0
people_vaccinated
                                        0
people_fully_vaccinated
daily_vaccinations_raw
                                        0
daily_vaccinations
total_vaccinations_per_hundred
                                        0
people_vaccinated_per_hundred
people_fully_vaccinated_per_hundred
daily_vaccinations_per_million
                                        0
vaccines
                                        0
source_name
                                        0
source_website
dtype: int64
```

Finally we managed to organize the lost data

Lets transform date column

```
In [27]:
data["date"] = pd.to_datetime(data["date"])
data = data.sort_values("date", ascending = True )
data["date"] = data["date"].dt.strftime("%Y-%m-%d")
In [28]:
unique_dates = data["date"].unique()
In [29]:
data.head()
```

Out[29]:

	c o u n tr y	is o — c o d e	d a t e	tota l_v acci nati ons	peo ple_ vac cina ted	peopl e_ful ly_va ccina ted	daily _vac cinat ions_ raw	dail y_v acci nati ons	total_v accinat ions_p er_hun dred	people _vacci nated_ per_hu ndred	people_f ully_vac cinated_ per_hun dred	daily_ vaccin ations_ per_mi llion	vac cine s	so ur ce _n am e	source_we bsite
5 8 5 1 7	N o r w a y	N O R	2 0 2 0 - 1 2 - 0 2	0.0	0.0	- 1.834 694e +07	0.0	0.0	0.0	0.0	- 44.7191 68	0.0	Mo der na, Pfiz er/B ioN Tec h	No rw egi an Ins tit ute of Pu bli c He alt h	https://git hub.com/f olkehelsei nstituttet/s urve
5 8 5 1	N o r w a	N O R	2 0 2 0 -	0.0	0.0	- 1.834 694e +07	0.0	0.0	0.0	0.0	- 44.7191 68	0.0	Mo der na, Pfiz er/B	No rw egi an Ins	https://git hub.com/f olkehelsei nstituttet/s

8	у		1 2 - 0 3										ioN Tec h	tit ute of Pu bli c He alt	urve
4 3 1 1 7	L at v ia	L V A	2 0 2 0 - 1 2 - 0 4	1.0	1.0	- 1.834 693e +07	0.0	0.0	0.0	0.0	- 44.7191 68	0.0	Joh nso n&J ohn son, Mo der na, Nov ava x, Pfiz er/B ioN.	Na tio nal He alt h Se rvi ce	https://dat a.gov.lv/d ati/eng/dat aset/covid 19-v
5 8 5 1 9	N o r w a y	N O R	2 0 2 0 - 1 2 - 0 4	0.0	0.0	- 1.834 694e +07	0.0	0.0	0.0	0.0	- 44.7191 68	0.0	Mo der na, Pfiz er/B ioN Tec h	No rw egi an Ins tit ute of Pu bli c He alt h	https://git hub.com/f olkehelsei nstituttet/s urve
5 8 5 2 0	N o r w a	N O R	2 0 2 0 - 1 2	0.0	0.0	- 1.834 694e +07	0.0	0.0	0.0	0.0	- 44.7191 68	0.0	Mo der na, Pfiz er/B ioN Tec	No rw egi an Ins tit ute	https://git hub.com/f olkehelsei nstituttet/s urve

y	1					h	of	
	0						Pu	
	5						bli	
							c	
							Не	
							alt	
							h	

our dates are listed

Data Visualization

First, let's watch total_vaccinations and daily_vaccinations animatedly on the world map.

Let's take a look at the rate of vaccination in countries

```
In [30]:
fig = px.choropleth(
    data,
    locations="iso_code",
    color="total_vaccinations",
    title='Number of people vaccinated',
    color_continuous_scale='viridis',
    animation_frame="date",
    projection = "natural earth",
    range_color = [0,5000000],
)
date=2020-12-022020-12-022021-03-292021-07-242021-11-182022-03-1501M2M3M4M5Mtotal_vaccinationsNumber of people vaccinated▶■
```

Animated world map by date of total vaccinations by country

```
In [31]:
fig = px.choropleth(
    data,
    locations= "iso_code",
    color = "daily_vaccinations",
    animation_frame = "date",
    color_continuous_scale= "viridis",
    projection= "natural earth",
    range_color= [0,1000000] ,
    title = "Number of daily vaccinations"
)
fig.show()
date=2020-12-022020-12-022021-03-292021-07-242021-11-182022-03-1500.2M0.4M0.6M0.8M1Mdaily_vaccinationsNumber of daily vaccinations>.
```

Animated world map by date of daily vaccinations by country

First we will create a new table by selecting the columns we will use.

```
In [32]:
columns = ["country","iso_code","total_vaccinations","people_vaccinated","total_vacci
nations_per_hundred","vaccines","daily_vaccinations"]

vacc_data = data[columns].groupby("country").max().sort_values("total_vaccinations",
ascending = True)

In [33]:
vacc_data.head()
```

Out[33]:

Out[33]	•					
	iso_co de	total_vaccinati	people_vaccina ted	total_vaccinations_per_hu ndred	vaccines	daily_vaccinati ons
country						
Pitcairn	PCN	94.0	47.0	200.00	Oxford/AstraZen eca	1.0
Tokelau	TKL	1936.0	968.0	141.52	Pfizer/BioNTech	23.0
Niue	NIU	4161.0	1650.0	257.81	Pfizer/BioNTech	87.0
Montserr at	MSR	4211.0	1897.0	84.54	Oxford/AstraZen eca	53.0
Falkland Islands	FLK	4407.0	2632.0	124.91	Oxford/AstraZen eca	189.0

223 non-null object

In [34]:

0 iso_code

```
vacc_data.info()

<class 'pandas.core.frame.DataFrame'>

Index: 223 entries, Pitcairn to China

Data columns (total 6 columns):

# Column Non-Null Count Dtype
--- --- ---- -----
```

```
total_vaccinations 223 non-null
                                                    float64
                                   223 non-null
    people_vaccinated
                                                    float64
    total_vaccinations_per_hundred 223 non-null
                                                    float64
 4 vaccines
                                    223 non-null
                                                    object
    daily_vaccinations
                                   223 non-null
                                                    float64
dtypes: float64(4), object(2)
memory usage: 12.2+ KB
In [35]:
fig = px.choropleth(
   vacc_data,
   locations= "iso_code",
   color = "total_vaccinations_per_hundred",
   title = "Number of total vaccinations per hunderd",
   color_continuous_scale= "rainbow"
)
fig.show("notebook")
```

50100150200250300total_vaccinations_per_hundredNumber of total vaccinations per hunderd

As can be seen on the map, countries have vaccination percentages.

Let's draw a map to see which countries these vaccines are used in.

```
In [36]:
fig = px.choropleth(
    locations = vacc_data.iso_code,
    color = vacc_data.vaccines,
```

```
title = "name of the vaccine",
color_continuous_scale= "rainbow"
)
```

colorOxford/AstraZenecaPfizer/BioNTechSinopharm/BeijingModernaOxford/AstraZeneca, Pfizer/BioNTechModerna, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZenecaModerna, Oxford/AstraZeneca, Pfizer/BioNTechOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingPfizer/BioNTech, Sputnik VJohnson&Johnson, Moderna, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTechOxford/AstraZeneca, Pfizer/BioNTech, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VOxford/AstraZeneca, Sinopharm/Beijing, Sputnik VJohnson&Johnson, ModernaJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/BeijingJohnson&Johnson, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingPfizer/BioNTech, Sinopharm/Beijing, Sputnik VOxford/AstraZeneca, Sinopharm/BeijingOxford/AstraZeneca, Pfizer/BioNTech, SinovacCovaxin, Oxford/AstraZeneca, Sinopharm/BeijingOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, SinovacModerna, Oxford/AstraZeneca, Sinopharm/Beijing, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VCovaxin, Oxford/AstraZenecaJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingPfizer/BioNTech, Sinopharm/BeijingOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VModerna, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, SinovacCovaxin, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacJohnson&Johnson, Moderna, Novavax, Pfizer/BioNTechSinopharm/Beijing, Sputnik VJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik Light, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VJohnson&Johnson, Pfizer/BioNTech, Sinopharm/BeijingJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VEpiVacCorona, Oxford/AstraZeneca, QazVac, Sinopharm/Beijing, Sputnik V, ZF2001Covaxin, Moderna,

Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VOxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik VAbdala, Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Soberana02, Sputnik Light, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VOxford/AstraZeneca, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VModerna, Pfizer/BioNTech, Sinopharm/Beijing, SinovacPfizer/BioNTech, SinovacModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacQazVac, Sinopharm/Beijing, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinopharm/Wuhan, Sputnik VCovaxin, Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, SinovacCanSino, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacAbdala, Soberana Plus, Soberana02Abdala, Sinopharm/Beijing, Sinovac, Soberana02, Sputnik Light, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik Light, Sputnik V, ZF2001Medigen, Moderna, Oxford/AstraZeneca, Pfizer/BioNTechCanSino, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCanSino, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCOVIran Barekat, Covaxin, FAKHRAVAC, Oxford/AstraZeneca, Razi Cov Pars, Sinopharm/Beijing, Soberana02, SpikoGen, Sputnik VPfizer/BioNTech, Sinovac, TurkovacEpiVacCorona, Sputnik VCanSino, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VAbdala, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VCanSino, Covaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCovaxin, Oxford/AstraZeneca, Sputnik VCanSino, Sinopharm/Beijing, Sinopharm/Wuhan, Sinovac, ZF2001name of the vaccine

Vaccine types can be seen on the sides according to the colors of the countries. By looking at this map, it can be seen which country has which vaccine.

How many people have been vaccinated

First, let's look at the statistics of the countries, then let's show these countries on the world map.

```
In [37]:
vacc_country = data.groupby(["country", "iso_code", "vaccines"])['total_vaccinations',
```

```
'total_vaccina
tions_per_hundred',
                                                                      'daily_vaccinat
ions',
                                                                      'daily_vaccinat
ions_per_million',
                                                                      'people_vaccina
ted',
                                                                      'people_vaccina
ted_per_hundred',
                                                                       'people_fully_
vaccinated', 'people_fully_vaccinated_per_hundred'
                                                                      ].max().reset_i
ndex()
/opt/conda/lib/python3.7/site-packages/ipykernel_launcher.py:1: FutureWarning:
Indexing with multiple keys (implicitly converted to a tuple of keys) will be dep
recated, use a list instead.
In [38]:
def trace_bar(data, feature, title, xlab, ylab,color):
    data = data.sort_values(feature, ascending=False)
    trace = go.Bar(
            x = data['country'],
            y = data[feature],
            marker=dict(color=color),
            text=data['country']
        )
```

```
data = [trace]
layout = dict(title = title,
       xaxis = dict(
            title = xlab,
            showticklabels=True,
            tickangle=45,
            zeroline=True,
            zerolinewidth=1,
            zerolinecolor='grey',
            showline=True,
            linewidth=2,
            linecolor='black',
            mirror=True,
            tickfont=dict(
                    size=10,
                    color='black'),),
       yaxis = dict(
            title = ylab,
            gridcolor='lightgrey',
            zeroline=True,
            zerolinewidth=1,
```

ChinaIndiaUnited

StatesBrazilIndonesiaJapanBangladeshPakistanVietnamMexicoGermanyRussiaPhilippines TurkeyIranFranceUnited KingdomItalyThailandSouth

KoreaEnglandArgentinaSpainCanadaColombiaEgyptMalaysiaPeruSaudi

ArabiaAustraliaMoroccoPolandChileTaiwanMyanmarUzbekistanNepalSri

LankaVenezuelaCambodiaCubaNetherlandsSouth

AfricaEcuadorUkraineNigeriaEthiopiaMozambiqueBelgiumUnited Arab

EmiratesPortugalSwedenKazakhstanGreeceRwandaAustriaIsraelIraqAngolaCzechiaKeny aUgandaRomaniaHungaryGuatemalaSwitzerlandDominican RepublicHong

KongSingaporeAlgeriaAzerbaijanDenmarkTunisiaBoliviaGhanaScotlandHondurasFinlandB elarusNorwayNew ZealandIrelandTajikistanEl SalvadorCote d'IvoireCosta

IrelandPalestineBahrainLibyaZambiaSyriaBeninSloveniaKyrgyzstanLatviaGeorgiaTogoAlb aniaNigerMauritaniaBotswanaSenegalMauritiusSomaliaBurkina FasoSierra LeoneMoldovaArmeniaMalawiEstoniaBosnia and HerzegovinaMaliNorth

MacedoniaKosovoBhutanCyprusCameroonTrinidad and

TobagoJamaicaFijiMadagascarTimorLuxembourgMaltaMacaoLiberiaBruneiCentral African RepublicDemocratic Republic of

Congo Maldives Les otho Guyana Namibia Congo Yemen Iceland Cape

VerdeMontenegroComorosNorthern CyprusSouth SudanPapua New GuineaGuinea-BissauGabonEswatiniSurinameEquatorial GuineaBelizeChadNew CaledoniaFrench PolynesiaGambiaSolomon IslandsBahamasBarbadosSamoaHaitiCuracaoJerseySao Tome and PrincipeSeychellesVanuatuIsle of ManArubaDjiboutiAndorraGuernseyTongaCayman IslandsKiribatiBermudaAntigua and BarbudaSaint LuciaGibraltarFaeroe IslandsGrenadaGreenlandLiechtensteinSaint Vincent and the GrenadinesTurks and Caicos IslandsSan

MarinoChinaBangladeshPhilippinesThailandColombiaMoroccoNepalSouth AfricaBelgiumRwandaKenyaDominican RepublicTunisiaBelarusCote d'IvoireSerbiaSlovakiaAfghanistanBulgariaSyriaTogoMauritiusMalawiBhutanMadagascarB runeiNamibiaComorosEswatiniFrench PolynesiaHaitiIsle of ManCayman IslandsFaeroe IslandsSan MarinoBonaire Sint Eustatius and SabaTuvaluPitcairn00.5B1B1.5B2B2.5B3B

Vaccination total per countryCountryVaccination total

```
In [40]:
```

trace_bar(vacc_country, 'daily_vaccinations', 'Daily vaccinations per country', 'Coun
try', 'Daily vaccinations', "red")

ChinaIndiaBangladeshUnited

States Pakistan Japan Brazil Indonesia Vietnam Mexico Philippines Ethiopia Turkey Iran Germany Russia United Kingdom South

KoreaThailandEnglandFranceSpainItalyEgyptTaiwanMalaysiaCanadaSri

LankaVenezuelaArgentinaGhanaIraqUzbekistanColombiaSaudi

ArabiaMoroccoPeruNepalMyanmarEcuadorKazakhstanPolandMozambiqueCubaRwandaNi caraguaChileAustraliaUgandaCambodiaNetherlandsNigeriaAlgeriaUkraineKenyaSouth AfricaIsraelDominican RepublicAngolaGuineaTunisiaSudanUnited Arab

EmiratesBelgiumHondurasPortugalLaosBotswanaDenmarkRomaniaHungaryMongoliaCote d'IvoireAustriaSwedenGreeceBoliviaZimbabweCzechiaGuatemalaSwitzerlandBhutanIrela ndParaguaySyriaJordanCosta RicaHong KongAzerbaijanNigerTajikistanNew ZealandSingaporeBelarusEl

Salvador Finland Panama Scotland Afghanistan Norway Mauritania Serbia Wales Oman Benin Senegal Mali Lebanon Uruguay Cameroon Palestine Turkmenistan Kuwait Burkina

Faso Liberia Togo Tanzania Qatar Zambia Slovakia Croatia Somalia Sierra

LeoneGeorgiaLithuaniaBulgariaDemocratic Republic of CongoLibyaNorthern

IrelandKyrgyzstanBahrainMalawiMauritiusKosovoSloveniaCentral African

RepublicArmeniaTrinidad and TobagoAlbaniaGuinea-BissauLatviaNamibiaMoldovaNorth MacedoniaLesothoFijiBosnia and

Herzegovina Jamaica Cyprus Montene gro Timor Eswatini Madagas car Yemen Estonia South Sudan Guyana Malta Brune i Maldives I celand Luxembourg Gabon Congo Macao Cape

VerdeSurinameIsle of ManNorthern CyprusComorosPapua New GuineaNew CaledoniaEquatorial

GuineaBelizeBarbadosHaitiChadGambiaCuracaoBahamasSamoaFrench PolynesiaArubaSao Tome and PrincipeSeychellesSolomon IslandsTongaVanuatuSaint LuciaAndorraGuernseyKiribatiAntigua and

BarbudaJerseyBermudaGreenlandGibraltarCayman IslandsFaeroe IslandsSint Maarten (Dutch part)DominicaSaint Vincent and the GrenadinesCook IslandsDjiboutiSaint Kitts and NevisLiechtensteinSan MarinoGrenadaNauruTurks and Caicos IslandsChinaBrazilTurkeyThailandTaiwanGhanaPeruMozambiqueUgandaKenyaTunisiaLao sCote d'IvoireCzechiaSyriaTajikistanPanamaWalesUruguayLiberiaCroatiaDemocratic Republic of CongoMauritiusAlbaniaLesothoTimorGuyanaGabonNorthern CyprusBarbadosSamoaTongaAntigua and BarbudaFaeroe IslandsSaint Kitts and NevisBurundiBonaire Sint Eustatius and SabaPitcairn05M10M15M20M

Daily vaccinations per countryCountryDaily vaccinations

In [41]:

trace_bar(vacc_country, 'daily_vaccinations_per_million', 'Daily vaccinations per million per country', 'Country', 'Daily vaccinations per million', "magenta")

BhutanIsle of ManBotswanaNiueFalkland IslandsNauruNicaraguaCook
IslandsMongoliaGibraltarWallis and FutunaCubaGuernseySaint
HelenaArubaTaiwanSeychellesRwandaDenmarkAndorraBangladeshSri
LankaAnguillaPitcairnCuracaoSan MarinoSint Maarten (Dutch part)IcelandFaeroe
IslandsTuvaluIsraelGreenlandEcuadorBermudaLaosWalesLiechtensteinTongaKazakhstan
IrelandMontenegroBruneiNew CaledoniaSouth KoreaVietnamCambodiaCosta
RicaDominican RepublicTokelauMalaysiaPanamaSamoaVenezuelaMauritiusNew
ZealandMaltaJapanFijiNetherlandsBarbadosNorthern CyprusChileUnited Arab
EmiratesChinaCayman IslandsHondurasSaint Kitts and
NevisTurkeyPortugalEnglandPhilippinesMaldiyesIranMauritaniaSingaporeSpainCanadaUn

NevisTurkeyPortugalEnglandPhilippinesMaldivesIranMauritaniaSingaporeSpainCanadaUn ited KingdomUruguayTunisiaCyprusGhanaGermanyBelgiumTurks and Caicos IslandsNepalHungaryScotlandFinlandAntigua and BarbudaGuineaTrinidad and TobagoMexicoNorwayBahrainQatarUzbekistanParaguayNorthern

IrelandAustriaKiribatiJerseyThailandCape

VerdeEthiopiaDominicaOmanPeruGuyanaAustraliaHong KongSaudi ArabiaFrench PolynesiaLuxembourgBelizeBritish Virgin IslandsEl SalvadorItalySao Tome and PrincipeSwedenFranceKosovoMoroccoMontserratSwitzerlandUnited StatesSurinameGreeceIraqSaint

Lucia Macao Mozambi que Lithuania Argentina Pakistan Eswatini Kuwa it Serbia Slovenia Czechia Latvia Bolivia Brazil Poland Guinea-

BissauCroatiaJordanTajikistanAzerbaijanBahamasTimorLebanonBelarusColombiaPalestin eMonacoEstoniaLiberiaSaint Vincent and the GrenadinesNorth MacedoniaBonaire Sint Eustatius and

SabaGeorgiaIndiaRussiaRomaniaZimbabweIndonesiaTurkmenistanLesothoSlovakiaMyan

marNamibiaVanuatuUgandaComorosAlbaniaArmeniaEgyptUkraineAlgeriaAngolaGuatem alaGrenadaSyriaBeninJamaicaTogoCote d'IvoireKenyaBosnia and

HerzegovinaMoldovaSierra LeoneBulgariaCentral African RepublicSouth

AfricaSudanEquatorial GuineaKyrgyzstanSenegalLibyaNigerSolomon

IslandsGabonMaliSomaliaZambiaBurkina

FasoAfghanistanCameroonGambiaCongoNigeriaMalawiSouth

SudanDjiboutiTanzaniaPapua New GuineaHaitiMadagascarYemenDemocratic Republic of CongoChadBurundiBhutanNicaraguaGuernseyDenmarkCuracaoIsraelLiechtensteinNew CaledoniaTokelauNew ZealandNorthern CyprusSaint Kitts and NevisIranUruguayTurks and Caicos

Is lands Guinea Uzbekistan Thail and Guyana Belize France Surina me Lithuania Slovenia Guinea Lithuania Guinea Lithu

Bissau Timor Estonia India Lesotho Comoros Angola Togo Bulgaria Senegal Somalia Congo Papua New Guinea Burundi 020k 40k 60k 80k 100k 120k

Daily vaccinations per million per countryCountryDaily vaccinations per million

```
In [42]:
```

```
trace_bar(vacc_country, 'people_vaccinated', 'People vaccinated per country', 'Countr
y','People vaccinated', "lightblue" )
```

ChinaIndiaBrazilUnited

States Indonesia Japan Vietnam Russia Philippines Pakistan Bangladesh Mexico Spain Iran Germany Turkey Thailand Colombia France United Kingdom Italy South

KoreaEgyptEnglandArgentinaCanadaPolandPeruMalaysiaMyanmarSaudi

A rabia Uzbekistan Morocco Nepal Ethiopia Australia Venezuela Nigeria South

AfricaCubaTaiwanChileSri

Lanka Ukraine Ecuador Cambodia Mozambi que Uganda Netherlands Kenya Angola Iraq United Arab

Kong Hungary Switzerland Belarus Laos Nicaragua Romania Honduras Azerbaijan Sudan Tajiki stan Afghanistan Zimbabwe Singapore Denmark Jordan El

SalvadorFinlandScotlandTurkmenistanCosta RicaNorwayNew

FasoPalestineLithuaniaSomaliaBulgariaSierra

Leone Georgia Malawi Togo Mauritania Kyrgyzstan Senegal Botswana Northern

IrelandCameroonLatviaMaliAlbaniaSloveniaBahrainArmeniaMadagascarLiberiaMauritiusMoldovaBosnia and HerzegovinaCentral African RepublicKosovoEstoniaDemocratic

Republic of CongoNorth MacedoniaLesothoJamaicaTrinidad and

TobagoTimorCyprusBhutanCongoFijiYemenMacaoSouth SudanGuinea-

BissauLuxembourgMaltaGuyanaNamibiaBruneiMaldivesEswatiniCape

VerdeComorosGambiaPapua New GuineaIcelandGabonMontenegroNorthern

CyprusChadSurinameEquatorial GuineaBelizeSolomon IslandsNew CaledoniaFrench PolynesiaSamoaBahamasHaitiBarbadosDjiboutiVanuatuSao Tome and PrincipeCuracaoArubaSeychellesJerseyKiribatiTongaIsle of ManChinaVietnamSpainFranceArgentinaSaudi ArabiaVenezuelaSri LankaNetherlandsKazakhstanGuatemalaCzechiaBelarusSudanJordanNorwayPanamaSlov akiaSyriaLithuaniaTogoCameroonArmeniaCentral African RepublicJamaicaFijiMaltaCape VerdeMontenegroSolomon IslandsBarbadosSeychellesCayman IslandsGibraltarSaint Kitts and NevisBonaire Sint Eustatius and SabaTuvaluPitcairn00.5B1B1.5B2B2.5B3B

People vaccinated per countryCountryPeople vaccinated

In [43]:

trace_bar(vacc_country, 'people_vaccinated_per_hundred', 'People vaccinated per hundred', 'Country', 'People vaccinated per hundred', "orange")

GibraltarCubaUnited Arab

MarinoNiuePitcairnColombiaPhilippinesPanamaNepalChileBruneiSingaporeMaltaCayman IslandsBhutanArgentinaMacaoSouth KoreaCambodiaSamoaAustraliaHong KongCanadaCook IslandsUruguaySeychellesPeruFaeroe IslandsCosta RicaNauruItalyIcelandMalaysiaNew

ZealandDenmarkPolandEcuadorNicaraguaArubaJerseyIrelandScotlandTaiwanAntigua and BarbudaTurks and Caicos IslandsFranceWalesCroatiaBelgiumNorwayThailandSri LankaUzbekistanWallis and FutunaMauritiusEnglandNetherlandsHondurasUnited KingdomVenezuelaSwedenUnited

StatesBangladeshGreeceGermanyLaosLuxembourgFijiAustriaIranNorthern

IrelandAndorraFalkland IslandsOmanNorthern CyprusSaudi

ArabiaMaldivesMonacoGreenlandLithuaniaBonaire Sint Eustatius and

SabaIsraelTongaLatviaRussiaSaint HelenaTurkmenistanIndonesiaTokelauIndiaBahrainEl SalvadorAnguillaLiechtensteinSwitzerlandMongoliaTurkeyBelarusRwandaBritish Virgin IslandsMoroccoHungaryFrench PolynesiaMexicoDominican RepublicCuracaoNew CaledoniaEstoniaCzechiaKiribatiSint Maarten (Dutch part)Cape

VerdeSloveniaBotswanaTunisiaSaint Kitts and

NevisGuyanaBoliviaKosovoBelizePakistanBarbadosAzerbaijanParaguayTrinidad and TobagoTuvaluTajikistanTimorSlovakiaSao Tome and

 $\label{thm:principe} Principe Kazakh stan Serbia Romania Myanmar Montenegro Jordan Surina me Dominica Albania Mozambique Egypt Guatemala Bahamas North$

MacedoniaGeorgiaComorosGrenadaPalestineMontserratArmeniaLebanonVanuatuLesotho UkraineSouth AfricaSolomon IslandsZimbabweAngolaEswatiniMauritaniaSaint Vincent and the GrenadinesLibyaSaint LuciaUgandaBosnia and HerzegovinaGhanaBulgariaCote d'IvoireJamaicaGuinea-BissauGuineaIraqBeninMoldovaKyrgyzstanKenyaSierra LeoneEthiopiaLiberiaCentral African RepublicTogoEquatorial

Guinea Namibia Algeria Djibouti Gambia Zambia Gabon Afghanistan Syria Congo Somalia Sudara Gambia Congo Somalia Congo Somalia

nNigeriaBurkina FasoNigerSenegalMalawiTanzaniaMaliCameroonSouth SudanMadagascarPapua New GuineaYemenChadHaitiDemocratic Republic of CongoBurundiGibraltarPortugalJapanPitcairnBruneiMacaoCanadaCosta RicaDenmarkIrelandWalesUzbekistanUnited KingdomGermanyNorthern IrelandMaldivesTongaTokelauSwitzerlandMoroccoNew CaledoniaSloveniaKosovoTrinidad and TobagoKazakhstanSurinameBahamasMontserratSouth AfricaSaint Vincent and the GrenadinesBulgariaBeninLiberiaDjiboutiCongoSenegalMadagascarBurundi050100150200 250300

People vaccinated per hundred per countryCountryPeople vaccinated per hundred

```
In [44]:
def plot_scatter(data, x, y, size, color, hover_name, title):
    fig = px.scatter(data, x=x, y=y, size=size, color=color, hover_name=hover_name, t
itle=title, size_max=80)
   fig.update_layout({"legend_orientation":"h"})
   fig.update_layout(legend = dict(yanchor ="top", y = -0.2))
   fig.update_layout({"legend_title":"Vaccine scheme"})
   fig.update_layout({"plot_bgcolor":"rgba(0,0,0,0)","paper_bgcolor":"rgba(0,0,0,0)
"})
   fig.update_xaxes(showline=True, linewidth=2, linecolor='black', mirror=True)
   fig.update_yaxes(showline=True, linewidth=2, linecolor='black', mirror=True)
   fig.update_xaxes(zeroline=True, zerolinewidth=1, zerolinecolor='grey')
   fig.update_yaxes(zeroline=True, zerolinewidth=1, zerolinecolor='grey')
   fig.update_xaxes(showgrid=True, gridwidth=1, gridcolor='lightgrey')
   fig.update_yaxes(showgrid=True, gridwidth=1, gridcolor='lightgrey')
   fig.show()
In [45]:
plot_scatter(vacc_country, x = "total_vaccinations", y = "daily_vaccinations",
            size = "total_vaccinations", color="vaccines",
```

hover_name = "country", title = "Vaccinations (Total vs. Daily) grouped p
er country and vaccines")

01B2B3B4B050M

Vaccine schemeJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingOxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VOxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTechOxford/AstraZenecaOxford/AstraZeneca, Pfizer/BioNTechOxford/AstraZeneca, Pfizer/BioNTech, Sputnik VCanSino, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VModerna, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik VPfizer/BioNTechJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik Light, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingSinopharm/Beijing, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VModerna, Pfizer/BioNTechCovaxin, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacJohnson&Johnson, Oxford/AstraZenecaJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/BeijingSinopharm/BeijingJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, SinovacCovaxin, Oxford/AstraZenecaCanSino, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacCanSino, Sinopharm/Beijing, Sinopharm/Wuhan, Sinovac, ZF2001Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacCovaxin, Oxford/AstraZeneca, Sinopharm/BeijingModerna, Oxford/AstraZeneca, Sinopharm/Beijing, Sputnik VAbdala, Soberana Plus, Soberana02Johnson&Johnson, Moderna, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCovaxin, Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, SinovacJohnson&Johnson, Pfizer/BioNTechPfizer/BioNTech, Sinopharm/Beijing, Sputnik VOxford/AstraZeneca, Sputnik VModernaModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VOxford/AstraZeneca, Sinopharm/BeijingModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VJohnson&Johnson, ModernaJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VPfizer/BioNTech, SinovacJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VCovaxin, Oxford/AstraZeneca, Sputnik VJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCOVIran Barekat, Covaxin, FAKHRAVAC, Oxford/AstraZeneca, Razi Cov Pars, Sinopharm/Beijing,

Soberana02, SpikoGen, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VQazVac, Sinopharm/Beijing, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VJohnson&Johnson, Moderna, Novavax, Pfizer/BioNTechOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VPfizer/BioNTech, Sinopharm/BeijingCanSino, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCanSino, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VAbdala, Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Soberana02, Sputnik Light, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, SinovacCanSino, Covaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VCovaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VEpiVacCorona, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VPfizer/BioNTech, Sputnik VOxford/AstraZeneca, Sinopharm/Beijing, Sputnik VModerna, Pfizer/BioNTech, Sinopharm/Beijing, SinovacJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VMedigen, Moderna, Oxford/AstraZeneca, Pfizer/BioNTechModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VJohnson&Johnson, Pfizer/BioNTech, Sinopharm/BeijingModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacPfizer/BioNTech, Sinovac, TurkovacEpiVacCorona, Oxford/AstraZeneca, QazVac, Sinopharm/Beijing, Sputnik V, ZF2001Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinopharm/Wuhan, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik Light, Sputnik V, ZF2001Abdala, Sinopharm/Beijing, Sinovac, Soberana02, Sputnik Light, Sputnik VAbdala, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, SinovacVaccinations (Total vs. Daily) grouped per country and vaccinestotal_vaccinationsdaily_vaccinations

In [46]:

plot_scatter(vacc_country,x = "people_vaccinated", y = "daily_vaccinations_per_millio
n", size = "total_vaccinations", color = "vaccines", hover_name = "country",title = "
Vaccinations (daily / million vs. iso_code) grouped per country and vaccines")

01B2B3B4B-100k0100k

Vaccine schemeJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingOxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VOxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTechOxford/AstraZeneca, Pfizer/BioNTechOxford/AstraZeneca, Pfizer/BioNTech, Sputnik VCanSino, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VModerna,

Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik VPfizer/BioNTechJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik Light, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingSinopharm/Beijing, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VModerna, Pfizer/BioNTechCovaxin, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacJohnson&Johnson, Oxford/AstraZenecaJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/BeijingJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/BeijingSinopharm/BeijingJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, SinovacCovaxin, Oxford/AstraZenecaCanSino, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacCanSino, Sinopharm/Beijing, Sinopharm/Wuhan, Sinovac, ZF2001Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, SinovacCovaxin, Oxford/AstraZeneca, Sinopharm/BeijingModerna, Oxford/AstraZeneca, Sinopharm/Beijing, Sputnik VAbdala, Soberana Plus, Soberana02Johnson&Johnson, Moderna, Pfizer/BioNTechJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCovaxin, Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, SinovacJohnson&Johnson, Pfizer/BioNTechPfizer/BioNTech, Sinopharm/Beijing, Sputnik VOxford/AstraZeneca, Sputnik VModernaModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VOxford/AstraZeneca, Sinopharm/BeijingModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VJohnson&Johnson, ModernaJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VPfizer/BioNTech, SinovacJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VCovaxin, Oxford/AstraZeneca, Sputnik VJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCOVIran Barekat, Covaxin, FAKHRAVAC, Oxford/AstraZeneca, Razi Cov Pars, Sinopharm/Beijing, Soberana02, SpikoGen, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VQazVac, Sinopharm/Beijing, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VJohnson&Johnson, Moderna, Novavax, Pfizer/BioNTechOxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VPfizer/BioNTech, Sinopharm/BeijingCanSino, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacCanSino, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VAbdala, Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Soberana02, Sputnik Light, Sputnik VOxford/AstraZeneca, Pfizer/BioNTech, SinovacCanSino, Covaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VCovaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VEpiVacCorona, Sputnik VJohnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik VPfizer/BioNTech, Sputnik VOxford/AstraZeneca, Sinopharm/Beijing, Sputnik VModerna, Pfizer/BioNTech, Sinopharm/Beijing, SinovacJohnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacJohnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik VMedigen, Moderna, Oxford/AstraZeneca, Pfizer/BioNTechModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik VJohnson&Johnson, Pfizer/BioNTech, Sinopharm/BeijingModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, SinovacPfizer/BioNTech, Sinovac, TurkovacEpiVacCorona, Oxford/AstraZeneca, QazVac, Sinopharm/Beijing, Sputnik V, ZF2001Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinopharm/Wuhan, Sputnik VModerna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik Light, Sputnik V, ZF2001Abdala, Sinopharm/Beijing, Sinovac, Soberana02, Sputnik Light, Sputnik VAbdala, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik VJohnson&Johnson, Oxford/AstraZeneca, SinovacVaccinations (daily / million vs. iso_code) grouped per country and vaccinespeople_vaccinateddaily_vaccinations_per_million

```
),
            colorbar = dict(
                title = 'Total vaccinations',
                tickprefix = '')
        )
data = [trace]
layout = go.Layout(
    title = 'Total vaccinations per country',
    geo = dict(
        showframe = True,
        showlakes = False,
        showcoastlines = True,
        projection = dict(
            type = 'natural earth'
        )
    )
)
fig = dict( data=data, layout=layout )
00.5B1B1.5B2B2.5B3BTotal vaccinationsTotal vaccinations per country
in the total vaccination of countries;
```

The country with the least vaccines is Chad.

China is the country with the most vaccines overall.

```
In [48]:
trace = go.Choropleth(
            locations = vacc_country['country'],
            locationmode='country names',
            z = vacc_country['daily_vaccinations'],
            text = vacc_country['country'],
            autocolorscale =False,
            reversescale = True,
            colorscale = 'viridis',
            marker = dict(
                line = dict(
                    color = 'rgb(0,0,0)',
                    width = 0.5)
            ),
            colorbar = dict(
                title = 'Daily vaccinations',
                tickprefix = '')
        )
data = [trace]
layout = go.Layout(
   title = 'Daily vaccinations per country',
    geo = dict(
        showframe = True,
```

in the daily vaccination of countries;

China is the country with the most daily vaccinations.

The country with the least daily vaccination is Chad.

```
color = 'rgb(0,0,0)',
                    width = 0.5)
            ),
            colorbar = dict(
                title = 'Daily vaccinations per million',
                tickprefix = '')
        )
data = [trace]
layout = go.Layout(
   title = 'Daily vaccinations per million per country',
    geo = dict(
        showframe = True,
        showlakes = False,
        showcoastlines = True,
        projection = dict(
            type = 'natural earth'
        )
    )
)
fig = dict( data=data, layout=layout )
```

Number of daily vaccines per million;

Bhudan is the country with the most daily vaccinations per million.

The country with the lowest daily vaccination per million is Chad.

```
In [50]:
trace = go.Choropleth(
            locations = vacc_country['country'],
            locationmode='country names',
            z = vacc_country['people_vaccinated'],
            text = vacc_country['country'],
            autocolorscale =False,
            reversescale = True,
            colorscale = 'viridis',
            marker = dict(
                line = dict(
                    color = 'rgb(0,0,0)',
                    width = 0.5)
            ),
            colorbar = dict(
                title = 'People vaccinated',
                tickprefix = '')
        )
data = [trace]
layout = go.Layout(
```

```
title = 'People vaccinated per country',
  geo = dict(
      showframe = True,
      showlakes = False,
      showcoastlines = True,
      projection = dict(
            type = 'natural earth'
      )
  )
)

fig = dict( data=data, layout=layout )

00.5B1B1.5B2B2.5B3BPeople vaccinatedPeople vaccinated per country
people vaccinated in countries;
```

China is the country with the most people vaccinated.

Chad is the country with the least number of people vaccinated.

```
reversescale = True,
            colorscale = 'viridis',
            marker = dict(
                line = dict(
                    color = 'rgb(0,0,0)',
                    width = 0.5)
            ),
            colorbar = dict(
                title = 'People vaccinated per hundred',
                tickprefix = '')
        )
data = [trace]
layout = go.Layout(
    title = "People vaccinated per hundred per country",
    geo = dict(
        showframe = True,
        showlakes = False,
        showcoastlines = True,
        projection = dict(
            type = 'natural earth'
        )
    )
)
```

```
fig = dict( data=data, layout=layout )
```

50100150200250300People vaccinated per hundredPeople vaccinated per hundred per country

percentage of people vaccinated in countries;

China has the highest percentage of people vaccinated.

democratic republic of congo has the lowest percentage of people vaccinated.

COVID-19 World Vaccination Progress

Basic Visualization

- 1. Vaccination by Country
 - 1.1 Total Vaccinations
 - 1.2 People Vaccinated
 - 1.3 People Fully Vaccinated
- 2. Vaccination by Country per Hundred
 - 2.1 Total Vaccinations
 - 2.2 People Vaccinated
 - 2.3 People Fully Vaccinated
- 3. Daily Vaccinations
 - 3.1 Daily Vaccinations by Country
 - 3.2 Daily Vaccinations by Country per Million

Advanced Visualization

- 1. Total Vaccination & 30-day Rolling
- 2. Daily Vaccination
 - 2.1 Day of Week
 - 2.2 Month
- 3. Total Vaccination Status Across Countries

Import packages

```
In [1]:
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
%matplotlib inline
Pandas default settings
In [2]:
# pd.set_option('display.max_rows', 500)
pd.set_option('display.max_columns', 30)
pd.set_option('display.float_format', '{:,.2f}'.format)
In [3]:
#Load Dataset
df_vaccination = pd.read_csv('../input/covid-world-vaccination-progress/country_vacci
nations.csv')
#data is from : https://gpreda/covid-world-vaccination-progress
```

Exploring the dataset

In [4]:

#Display first 5 rows

df_vaccination.head()

Out[4]:

	c u	o in ry	is o - c o d e	d a t e	total _va ccin atio ns	peo ple_ vacc inat ed	peopl e_full y_vac cinate d	daily _vacc inatio ns_ra w	dail y_v acci nati ons	total_v accinati ons_per _hundr ed	people_ vaccina ted_per _hundr ed	people_f ully_vac cinated_ per_hund red	daily_v accinati ons_pe r_milli on	vacc ines	so urc e_ na me	sourc e_we bsite
(g a is	Af gh in sst in	A F G	2 0 2 1 - 0 2 - 2 2	0.00	0.00	NaN	NaN	Na N	0.00	0.00	NaN	NaN	John son &Jo hnso n, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
1	g l a:	Af gh in sst in	A F G	2 0 2 1 - 0 2 - 2 3	Na N	Na N	NaN	NaN	1,36 7.00	NaN	NaN	NaN	34.00	John son & Johnson, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/

2	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 4	Na N	Na N	NaN	NaN	1,36 7.00	NaN	NaN	NaN	34.00	John son & Johnson, Oxford/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
3	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 5	Na N	Na N	NaN	NaN	1,36 7.00	NaN	NaN	NaN	34.00	John son &Jo hnso n, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/
4	Af gh an ist an	A F G	2 0 2 1 - 0 2 - 2 6	Na N	Na N	NaN	NaN	1,36 7.00	NaN	NaN	NaN	34.00	John son & Johnson, Oxfo rd/A straZ enec a, Pfize r/Bi	W orl d He alt h Or ga niz ati on	https: //covi d19. who.i nt/

In [5]:
df_vaccination.info()

<class 'pandas.core.frame.DataFrame'>

RangeIndex: 77709 entries, 0 to 77708

Data columns (total 15 columns):

#	Column	Non-Null Count	Dtype
0	country	77709 non-null	object
1	iso_code	77709 non-null	object
2	date	77709 non-null	object
3	total_vaccinations	40048 non-null	float64
4	people_vaccinated	37945 non-null	float64
5	people_fully_vaccinated	35465 non-null	float64
6	daily_vaccinations_raw	32591 non-null	float64
7	daily_vaccinations	77429 non-null	float64
8	total_vaccinations_per_hundred	40048 non-null	float64
9	people_vaccinated_per_hundred	37945 non-null	float64
10	people_fully_vaccinated_per_hundred	35465 non-null	float64
11	daily_vaccinations_per_million	77429 non-null	float64
12	vaccines	77709 non-null	object
13	source_name	77709 non-null	object
14	source_website	77709 non-null	object

dtypes: float64(9), object(6)

memory usage: 8.9+ MB

Content

The data (country vaccinations) contains the following information:

- Country- this is the country for which the vaccination information is provided;
- Country ISO Code ISO code for the country;
- **Date** date for the data entry; for some of the dates we have only the daily vaccinations, for others, only the (cumulative) total;
- **Total number of vaccinations** this is the absolute number of total immunizations in the country;
- **Total number of people vaccinated** a person, depending on the immunization scheme, will receive one or more (typically 2) vaccines; at a certain moment, the number of vaccination might be larger than the number of people;
- Total number of people fully vaccinated this is the number of people that received the entire set of immunization according to the immunization scheme (typically 2); at a certain moment in time, there might be a certain number of people that received one vaccine and another number (smaller) of people that received all vaccines in the scheme;
- **Daily vaccinations (raw)** for a certain data entry, the number of vaccination for that date/country;
- Daily vaccinations for a certain data entry, the number of vaccination for that date/country;
- **Total vaccinations per hundred** ratio (in percent) between vaccination number and total population up to the date in the country;
- **Total number of people vaccinated per hundred** ratio (in percent) between population immunized and total population up to the date in the country;
- Total number of people fully vaccinated per hundred ratio (in percent) between population fully immunized and total population up to the date in the country;
- **Daily vaccinations per million** ratio (in ppm) between vaccination number and total population for the current date in the country;
- Vaccines used in the country total number of vaccines used in the country (up to date);
- **Source name** source of the information (national authority, international organization, local organization etc.);
- Source website website of the source of information;

In [6]:

#Find the number or rows and columns

df_vaccination.shape

#There are 76095 rows and 15 columns

Out[6]:

(77709, 15)

In [7]:

df_vaccination.isnull().sum()

#There are no empty rows for country, iso_code or date columns.

Out[7]:

country	0
iso_code	0
date	0
total_vaccinations	37661
people_vaccinated	39764
people_fully_vaccinated	42244
daily_vaccinations_raw	45118
daily_vaccinations	280
total_vaccinations_per_hundred	37661
people_vaccinated_per_hundred	39764
people_fully_vaccinated_per_hundred	42244
daily_vaccinations_per_million	280
vaccines	0
source_name	0
source_website	0

dtype: int64

In [8]:

General Overview of the calculations in data

df_vaccination.describe()

Ou	Out[8]:												
	total_v accinat ions	people _vacci nated	people_fu lly_vacci nated	daily_vac cinations _raw	daily_v accinat ions	total_vaccin ations_per_h undred	people_vacci nated_per_h undred	people_fully_v accinated_per_ hundred	daily_vaccin ations_per_ million				
c o u nt	40,048	37,945. 00	35,465.00	32,591.0 0	77,429. 00	40,048.00	37,945.00	35,465.00	77,429.00				
m ea n	40,384 ,563.3 0	15,903, 249.33	12,278,48 6.43	276,490. 71	135,95 7.05	73.11	38.60	32.81	3,417.77				
st d	202,53 3,898. 56	63,330, 416.02	49,156,65 1.29	1,245,21 0.53	797,86 8.94	63.46	28.74	27.62	4,028.63				
m in	0.00	0.00	1.00	0.00	0.00	0.00	0.00	0.00	0.00				
2 5 %	472,14 1.25	314,53 8.00	211,664.0 0	5,240.00	972.00	13.31	9.72	5.58	704.00				
5 0 %	3,141, 270.00	1,939,9 70.00	1,442,791 .00	26,278.0 0	7,869.0 0	59.37	37.40	27.31	2,253.00				

7 5 %	15,091 ,858.2 5	8,102,7 81.00	6,399,728 .00	129,159. 00	45,644. 00	123.68	65.33	58.32	4,933.00
m a x	3,063, 391,00 0.00	1,266,4 26,000. 00	1,228,340 ,000.00	24,741,0 00.00	22,424, 286.00	333.76	123.75	121.14	117,497.00

Data Preparation

In [9]:

#drop the source_name, source_website and vaccine columns

df_vaccine_country = df_vaccination.drop(['source_name','source_website','vaccines'],
axis=1)

df_vaccine_country.head()

Out[9]:

	cou	iso _c od e	d at e	total_vacci natio ns	peopl e_vac cinate d	people_ fully_v accinate d	daily_v accinati ons_ra w	daily_ vacci nation s	total_vacc inations_p er_hundre d	people_va ccinated_p er_hundre d	people_fully _vaccinated _per_hundre d	daily_vac cinations_ per_millio n
0	Afg han ista n	A F G	2 0 2 1 - 0 2 - 2 2	0.00	0.00	NaN	NaN	NaN	0.00	0.00	NaN	NaN
1	Afg han	A F	2 0	NaN	NaN	NaN	NaN	1,367.	NaN	NaN	NaN	34.00

	ista n	G	2 1 - 0 2 - 2 3					00				
2	Afg han ista n	A F G	2 0 2 1 - 0 2 - 2 4	NaN	NaN	NaN	NaN	1,367. 00	NaN	NaN	NaN	34.00
3	Afg han ista n	A F G	2 0 2 1 - 0 2 - 2 5	NaN	NaN	NaN	NaN	1,367. 00	NaN	NaN	NaN	34.00
4	Afg han ista n	A F G	2 0 2 1 - 0 2 - 2 6	NaN	NaN	NaN	NaN	1,367. 00	NaN	NaN	NaN	34.00

In [10]:

convert Date column to date type and fill na values with 0 for calculation

```
df vaccine country["date"] = pd.to datetime(df vaccine country["date"], format = '%Y-
%m-%d')
df_vaccine_country = df_vaccine_country.replace([np.inf, -np.inf], np.nan)
df_vaccine_country = df_vaccine_country.fillna(0)
df_vaccine_country.isnull().sum()
Out[10]:
country
                                        0
iso_code
                                        0
date
                                        0
total_vaccinations
                                        0
people_vaccinated
                                        0
people_fully_vaccinated
                                        0
daily_vaccinations_raw
                                        0
daily_vaccinations
                                        0
total_vaccinations_per_hundred
                                        0
people_vaccinated_per_hundred
                                        0
people_fully_vaccinated_per_hundred
daily_vaccinations_per_million
                                        0
dtype: int64
```

In [11]:

#Function to find total, avergae, maximum and minimum of different vaccinations stat us by country

```
def vaccination_country(col_name,func_name):
    . . .
    Function that requires vaccination column name, and sum/mean/max/min function nam
e as string arguments.
    . . .
    if func_name == 'sum':
        return (df_vaccine_country[['country',col_name]].groupby(by='country')
                                  .sum()
                                  .sort_values(by=col_name,ascending= False)
                                 .reset_index()
                            )
    elif func_name == 'mean':
        return (df_vaccine_country[['country',col_name]].groupby(by='country')
                                  .mean()
                                  .sort_values(by=col_name,ascending= False)
                                  .reset_index()
                                )
    elif func_name == 'max':
        return (df_vaccine_country[['country',col_name]].groupby(by='country')
```

```
.max()
                                 .sort_values(by=col_name,ascending= False)
                                 .reset_index()
                                )
   elif func_name == 'min':
        return (df_vaccine_country[['country',col_name]].groupby(by='country')
                                 .min()
                                 .sort_values(by=col_name,ascending= False)
                                 .reset_index()
                                )
In [12]:
# Calculating different vaccinations for visualizations
max_total_vaccinations = vaccination_country('total_vaccinations','max')
sum_people_vaccinated = vaccination_country('people_vaccinated','sum')
sum_people_fully_vaccinated = vaccination_country('people_fully_vaccinated','sum')
avg_total_vaccinations = vaccination_country('total_vaccinations_per_hundred', 'mean')
avg_people_vaccinated = vaccination_country('people_vaccinated_per_hundred', 'mean')
avg_people_fully_vaccinated = vaccination_country('people_fully_vaccinated_per_hundre
d','mean')
avg_daily_vaccinations = vaccination_country('daily_vaccinations_per_million','mean')
In [13]:
```

```
#Function for Country with maximum and minimum daily vaccinations
def daily vaccination country(col name, func name):
    . . .
   A function that requires daily_vaccination column and max/min function name as st
ring arguments.
    . . .
    daily_vaccination = (df_vaccine_country
                                 .pivot_table(index='country',columns='date',values=c
ol_name)
                                    )
    if func_name == 'max':
        daily_vaccination['Highest Daily Vaccination'] = daily_vaccination.max(axis=1)
        daily_vaccination['Date - Highest Daily Vaccination'] = daily_vaccination.idx
max(axis=1)
        daily_vaccination.sort_values(by='Highest Daily Vaccination',ascending=False,
inplace=True)
        daily vaccination.rename axis('',axis=1,inplace=True)
        return daily vaccination[['Highest Daily Vaccination','Date - Highest Daily V
accination']].reset_index()
    elif func_name == 'min':
```

```
daily_vaccination.replace(0.00,np.nan,inplace=True)
        daily vaccination['Lowest Daily Vaccination'] = daily vaccination.min(axis=1)
        daily_vaccination['Date - Lowest Daily Vaccination'] = daily_vaccination.idxm
in(axis=1)
        daily vaccination.sort values(by='Lowest Daily Vaccination',ascending=False,i
nplace=True)
        daily_vaccination.rename_axis('',axis=1,inplace=True)
        return daily_vaccination[['Lowest Daily Vaccination','Date - Lowest Daily Vac
cination']].reset index()
In [14]:
#Calculating highest and lowest daily vaccination and the respective dates.
highest_daily_vaccination = daily_vaccination_country('daily_vaccinations','max')
lowest daily vaccination = daily vaccination country('daily vaccinations','min')
Data Visualization
1.1 Top & Bottom 5 Countries in terms of Total Vaccination
In [15]:
#Set sns theme and default figsize for all the sns visualizations.
sns.set_theme(style='whitegrid')
sns.set(rc={'figure.figsize' : (12,5)})
fig, axes = plt.subplots(2,1)
sns.barplot(x='country',y='total_vaccinations',data=max_total_vaccinations.head(),ax=
axes[0])
```

```
axes[0].set(xlabel = '', ylabel = 'Total Vaccinations', title ='Top 5 Countries in te
rms of total vaccinations!')
sns.barplot(x='country',y='total_vaccinations',data=max_total_vaccinations.tail(),ax=
axes[1])
axes[1].set(xlabel = '', ylabel = 'Total Vaccinations', title ='Bottom 5 Countries in
terms of total vaccinations!')
fig.tight_layout()
plt.show()
1.2 Top & Bottom 5 Countries in terms of People Vaccinated
In [16]:
fig, axes = plt.subplots(2,1)
sns.barplot(x='country',y='people_vaccinated',data=sum_people_vaccinated.head(),ax=ax
es[0])
axes[0].set(xlabel = '', ylabel = 'People Vaccinated', title ='Top 5 Countries in ter
ms of people vaccinated!')
sns.barplot(x='country', y='people_vaccinated',data=sum_people_vaccinated.tail(),ax=a
xes[1])
axes[1].set(xlabel = '', ylabel = 'People Vaccinated', title ='Bottom 5 Countries in
terms of people vaccinated!')
```

```
fig.tight_layout()
plt.show()
1.3 Top & Bottom 5 Countries in terms of People Fully Vaccinated
In [17]:
fig, axes = plt.subplots(2,1)
sns.barplot(x='country',y='people_fully_vaccinated',data=sum_people_fully_vaccinated.
head(),ax=axes[0])
axes[0].set(xlabel = '', ylabel = 'People Fully Vaccinated', title ='Top 5 Countries
in terms of people fully vaccinated!')
sns.barplot(x='country',y='people_fully_vaccinated',data=sum_people_fully_vaccinated.
tail(),ax=axes[1])
axes[1].set(xlabel = '', ylabel = 'People Fully Vaccinated', title ='Bottom 5 Countri
es in terms of people fully vaccinated!')
# plt.ticklabel_format(style='plain', axis='y') #Uncomment if y label needs to displa
y accurate values
fig.tight_layout()
plt.show()
2.1 Top & Bottom 5 Countries in terms of Total Vaccinations per Hundred
```

In [18]:

```
fig, axes = plt.subplots(2,1)
sns.barplot(x='country', y='total_vaccinations_per_hundred',data=avg_total_vaccinatio
ns.head(),ax=axes[0])
axes[0].set(xlabel='', ylabel='Average Vaccinations per 100', title='Top 5 Countries
in terms of average vaccinations per hundred!')
sns.barplot(x='country', y='total_vaccinations_per_hundred',data=avg_total_vaccinatio
ns.tail(),ax=axes[1])
axes[1].set(xlabel='', ylabel='Average Vaccinations per 100', title='Bottom 5 Countri
es in terms of average vaccinations per hundred!')
fig.tight_layout(h_pad=3)
plt.show()
2.2 Top & Bottom 5 Countries in terms of People Vaccinated per Hundred
In [19]:
fig, axes = plt.subplots(2,1)
sns.barplot(x='country', y='people_vaccinated_per_hundred',data=avg_people_vaccinated.
head(),ax=axes[0])
axes[0].set(xlabel='', ylabel='People Vaccinated per 100', title='Top 5 Countries in
terms of average people vaccinated per hundred!')
sns.barplot(x='country', y='people_vaccinated_per_hundred',data=avg_people_vaccinated.
tail(),ax=axes[1])
axes[1].set(xlabel='', ylabel='People Vaccinated per 100', title='Bottom 5 Countries
in terms of average people vaccinated per hundred!')
```

```
fig.tight layout()
plt.show()
2.3 Top & Bottom 5 Countries in terms of People Fully Vaccinated per Hundred
In [20]:
fig, axes = plt.subplots(2,1)
sns.barplot(x='country', y='people_fully_vaccinated_per_hundred',data=avg_people_full
y_vaccinated.head(),ax=axes[0])
axes[0].set(xlabel='', ylabel='People Fully Vaccinated per 100', title='Top 5 Countri
es in terms of average people fully vaccinated per hundred!')
sns.barplot(x='country', y='people_fully_vaccinated_per_hundred',data=avg_people_full
y_vaccinated.tail(),ax=axes[1])
axes[1].set(xlabel='', ylabel='People Fully Vaccinated per 100', title='Bottom 5 Coun
tries in terms of average people fully vaccinated per hundred!')
fig.tight_layout(h_pad=3)
plt.show()
3.1 Highest & Lowest 5 Daily Vaccination by Country
unfold moreshow hidden code
In [22]:
fig, axes = plt.subplots(1,2)
```

```
sns.barplot(data=daily_top5_highest,x="country", y="Highest Daily Vaccination",ax=axe
s[0],hue='Date - Highest Daily Vaccination')
axes[0].set(xlabel='',ylabel='Daily Vaccination',title='Highest Daily Vaccination by
Country')
sns.barplot(data=daily_top5_lowest,x="country", y="Lowest Daily Vaccination",ax=axes
[1],hue='Date - Lowest Daily Vaccination')
axes[1].set(xlabel='',ylabel='Daily Vaccination',title='Lowest Daily Vaccination by C
ountry')
# plt.ticklabel_format(style='plain', axis='y')
fig.tight_layout()
plt.show()
3.2 Top & Bottom 5 Daily Vaccination by Country per Million
In [23]:
fig, axes = plt.subplots(2,1)
sns.barplot(x='country', y='daily_vaccinations_per_million',data=avg_daily_vaccinatio
ns.head(),ax=axes[0])
axes[0].set(xlabel='', ylabel='Daily Vaccinations per Million', title='Top 5 Countrie
s in terms of daily vaccinations per million!')
sns.barplot(x='country', y='daily_vaccinations_per_million',data=avg_daily_vaccinatio
ns.tail(),ax=axes[1])
```

```
axes[1].set(xlabel='', ylabel='Daily Vaccinations per Million', title='Bottom 5 Count
ries in terms of daily vaccinations per million!')
fig.tight_layout(h_pad=3)
plt.show()
Advanced Data Visualization
Import Plotly Library
In [24]:
from plotly.offline import init notebook mode
import plotly.express as px
init_notebook_mode(connected=True)
1. Total Vaccination & 30-day Rolling by Top 5 Country
In [25]:
#Top 5 country with highest total vaccinations
list(max_total_vaccinations['country'].head())
Out[25]:
['China', 'India', 'United States', 'Brazil', 'Indonesia']
In [26]:
# Filter the top 5 countries and find their 30 day rolling average of total_vaccinati
ons
top5_country_total = ['China', 'India', 'United States', 'Brazil', 'Indonesia']
top5_country_total_day = df_vaccine_country[df_vaccine_country['country'].isin(top5_c
```

ountry total)].copy()

```
top5_country_total_day['30 - Day Rolling'] = top5_country_total_day['total_vaccinatio
ns'].rolling(window=30).mean()
In [27]:
fig = px.line(top5_country_total_day,x="date",y="total_vaccinations",color='country',
                labels={"country" : 'Top 5 Country', 'date' : 'Date', 'total_vaccina
tions' : "Total Vaccinations"},
                title="Total Vaccination Progress - Top 5 Country",template='plotly_d
ark')
for country in top5_country_total_day['country'].unique():
   fig.add_scatter(x=top5_country_total_day[top5_country_total_day['country'] == cou
ntry]['date']
                    ,y=top5_country_total_day[top5_country_total_day['country'] == co
untry]['30 - Day Rolling']
                    ,mode="lines",name='30 Day Rolling Vaccination ' + country)
2.1 Daily Vaccination by Day of Week by Top 5 Country
In [28]:
# Get the name of the day of vaccinations
top5_country_total_day['Day of Week'] = top5_country_total_day['date'].apply(lambda x:
x.day_name())
top5_country_total_day['Day of Week'].unique()
Out[28]:
array(['Sunday', 'Monday', 'Tuesday', 'Wednesday', 'Thursday', 'Friday',
       'Saturday'], dtype=object)
In [29]:
```

```
fig = px.box(top5_country_total_day,x='Day of Week',y='daily_vaccinations',color='cou
ntry',
             labels={"country" : 'Top 5 Country', 'daily_vaccinations' : "Daily Vacci
nation"},
                title="Dailly Vaccination by Day of Week - Top 5 Country", template='p
lotly_dark')
2.2 Daily Vaccination by Month by Top 5 Country
In [30]:
# Get the name of the month of vaccinations
top5_country_total_day['Month'] = top5_country_total_day['date'].apply(lambda x:x.mon
th_name())
top5_country_total_day['Month'].unique()
Out[30]:
array(['January', 'February', 'March', 'April', 'May', 'June', 'July',
       'August', 'September', 'October', 'November', 'December'],
      dtype=object)
In [31]:
linkcode
fig = px.bar(top5_country_total_day,x='Month',y='daily_vaccinations',color='country',
             labels={"country" : 'Top 5 Country', 'daily vaccinations' : "Daily Vacci
nation"},
                title="Dailly Vaccination by Month - Top 5 Country", template='plotly_
dark')
3. Total Vaccination Status Across Countries
In [32]:
fig = px.choropleth(max_total_vaccinations,locations='country',locationmode='country')
names',
```

```
color='total_vaccinations',hover_name="country", template='plotly_
dark',

title= 'Total Vaccination Status Across Countries',projection='nat
ural earth',

labels={'country' : 'Country','total_vaccinations' : 'Total Vaccin
ations'})
```