Covid vaccine analysis

Phase 3: Coding part

Project Title: Covid vaccine analysis with python programming

Items in the dataset:

- Countries
- Dates
- Vaccines
- Total Vaccinations

Desired data to find:

- Most commonly used vaccines in countries
- Average daily vaccination count in countries -Number of countries where vaccines are used
- Choropleth map of the most used vaccine

INPUT:

```
data=pd.DataFrame(columns=['Country', 'Vaccine',
'Total_vaccine'])
    for country in df["location"].unique():
    for vaccine in df["vaccine"].unique():
    filtered_data = df[(df['location'] == country) & (df['vaccine']
== vaccine)]
    total_count = filtered_data['total_vaccinations'].max()
    data = pd.concat([data, pd.DataFrame({'Country': [country],
'Vaccine': [vaccine], 'Total_vaccine': [total_count]})],
ignore_index=True)
SUB-INPUT:
    data.head(10)
```

OUTPUT:

| | Country | Vaccine | Total_vaccine |
|---|-----------|--------------------|---------------|
| 0 | Argentina | Moderna | 6507561 |
| 1 | Argentina | Oxford/AstraZeneca | 25977231 |
| 2 | Argentina | Sinopharm/Beijing | 28322602 |
| 3 | Argentina | Sputnik V | 20405678 |
| 4 | Argentina | CanSino | 610540 |
| 5 | Argentina | Pfizer/BioNTech | 14681054 |
| 6 | Argentina | Johnson&Johnson | NaN |
| 7 | Argentina | Novavax | NaN |
| 8 | Argentina | Sinovac | NaN |
| 9 | Argentina | Covaxin | NaN |

Most commonly used vaccines

INPUT:

```
data_2=pd.DataFrame(columns=['Country', 'Vaccine'])
data["Total_vaccine"] = pd.to_numeric(data["Total_vaccine"],
errors="coerce")
for country in data["Country"].unique():
new_data = data[data["Country"] == country]
max_vaccine = new_data.loc[new_data["Total_vaccine"].idxm
ax(), "Vaccine"]
data_2 = pd.concat([data_2, pd.DataFrame({'Country': [country], 'Vaccine': [max_vaccine]})], ignore_index=True)
```

SUB-INPUT:

data_2.head()

OUTPUT:

| | Country | Vaccine |
|---|-----------|-------------------|
| 0 | Argentina | Sinopharm/Beijing |
| 1 | Austria | Pfizer/BioNTech |
| 2 | Belgium | Pfizer/BioNTech |
| 3 | Bulgaria | Pfizer/BioNTech |
| 4 | Chile | Sinovac |

INPUT:

data_2["Vaccine"].value_counts().plot(kind="bar", color=["Red","Gray","Gray","Gray"])

OUTPUT:

<Axes: >

