

# EDA on COVID-19 Clinical Trials

## 1. Introduction

The COVID-19 pandemic initiated an unprecedented surge in clinical research activities around the world. Thousands of clinical trials were launched to discover treatments, vaccines, and diagnostic methods.

This project performs **Exploratory Data Analysis (EDA)** on a dataset of **COVID-19 related clinical trials** to understand trends, patterns, and important insights.

## 2. Objectives

- To explore the **status** of clinical trials.
- To analyze **study types** (Interventional, Observational).
- To identify the most common **medical conditions** studied.
- To find the major **sponsors** behind the studies.
- To study the **geographic distribution** of trials.
- To investigate the **phases** of Interventional studies.

## 3. Dataset Overview

The data is loaded from a CSV file using Python's pandas library.

The dataset contains columns such as:

- **Status:** Current state of the trial (e.g., Completed, Recruiting).
- **Conditions:** Disease(s) being studied.
- **Sponsor:** Organization leading or funding the study.
- **Study Type:** Type of research (Interventional, Observational).
- **Phase:** Phase of the trial (Phase 1–4).
- **Locations:** Where the trial is being conducted

## 4. Code, Explanation

### 4.1 Import Visualization Libraries

```
[126] import pandas as pd
import numpy as np
from matplotlib import pyplot as plt
%matplotlib inline
import matplotlib
import seaborn as sns
from plotly.subplots import make_subplots
import plotly.graph_objects as go
```

- **pandas:** For data handling.
- **numpy:** For numerical computations.
- **matplotlib.pyplot and seaborn:** For creating visualizations.

## 4.2 Reading the Dataset

importing the csv file as a dataframe


```
[127] df = pd.read_csv('/content/COVID clinical trials.csv')
```

## 4.3 Checking Basic Information

 df.shape

 (5783, 27)

```
[172] df.columns
```

 Index(['Rank', 'NCT Number', 'Title', 'Acronym', 'Status', 'Study Results', 'Conditions', 'Interventions', 'Outcome Measures', 'Sponsor/Collaborators', 'Gender', 'Age', 'Phases', 'Enrollment', 'Funded Bys', 'Study Type', 'Study Designs', 'Other IDs', 'Start Date', 'Primary Completion Date', 'Completion Date', 'First Posted', 'Results First Posted', 'Last Update Posted', 'Locations', 'Study Documents', 'URL'], dtype='object')

## 4.4 Data Cleaning

df.isnull().sum()

|                         |      |
|-------------------------|------|
|                         | 0    |
| Rank                    | 0    |
| NCT Number              | 0    |
| Title                   | 0    |
| Acronym                 | 3303 |
| Status                  | 0    |
| Study Results           | 0    |
| Conditions              | 0    |
| Interventions           | 886  |
| Outcome Measures        | 35   |
| Sponsor/Collaborators   | 0    |
| Gender                  | 10   |
| Age                     | 0    |
| Phases                  | 2461 |
| Enrollment              | 34   |
| Funded Bys              | 0    |
| Study Type              | 0    |
| Study Designs           | 35   |
| Other IDs               | 1    |
| Start Date              | 34   |
| Primary Completion Date | 36   |
| Completion Date         | 36   |
| First Posted            | 0    |
| Results First Posted    | 5747 |
| Last Update Posted      | 0    |
| Locations               | 585  |
| Study Documents         | 5601 |
| URL                     | 0    |

dtype: int64

## 4.5 Explore the Gender distribution in the studies

```
df['Gender'].unique()
```

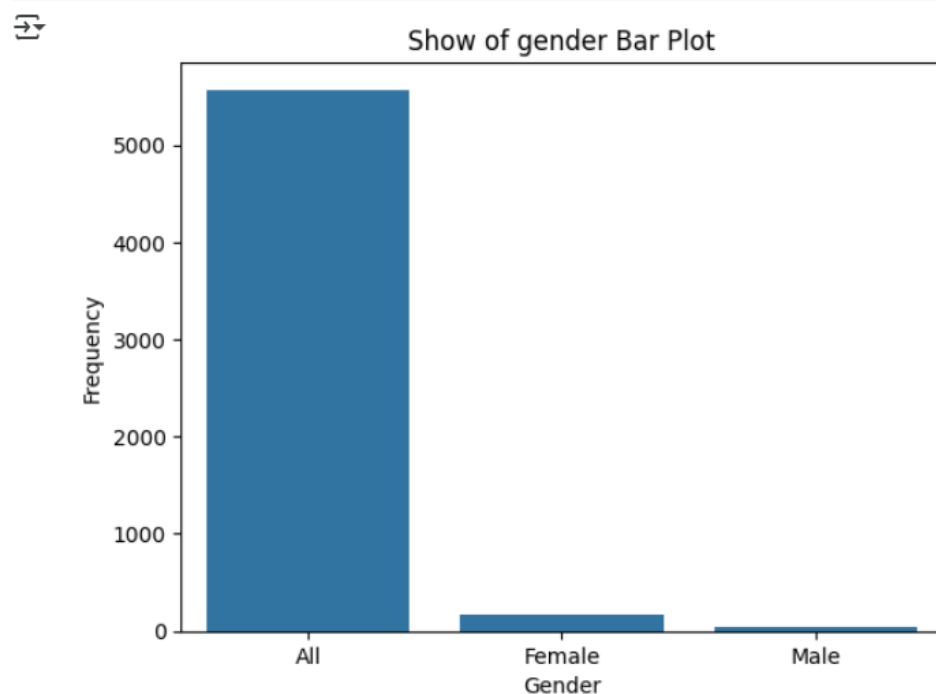
```
array(['All', 'Female', 'Male', nan], dtype=object)
```

```
[219] df['Gender'].value_counts()
```

|        | count |
|--------|-------|
| Gender |       |
| All    | 5567  |
| Female | 162   |
| Male   | 44    |

dtype: int64

```
sns.barplot(x=df['Gender'].value_counts().index,  
            y=df['Gender'].value_counts().values)  
plt.xlabel('Gender')  
plt.ylabel('Frequency')  
plt.title('Show of gender Bar Plot')  
plt.show()
```



**Observation:** The majority of studies are categorized as "All" genders, with few specifically labeled as "Female" or "Male."

## 4.6 Exploring Study Status Distribution

```
df.Status.unique()
```

```
array(['Active, not recruiting', 'Not yet recruiting', 'Recruiting',  
      'Enrolling by invitation', 'Suspended', 'Completed', 'Withdrawn',  
      'Terminated', 'No longer available', 'Available',  
      'Approved for marketing', 'Temporarily not available'],  
      dtype=object)
```

```
df['Status'].value_counts()
```

|                         | count |
|-------------------------|-------|
| Recruiting              | 2805  |
| Completed               | 1025  |
| Not yet recruiting      | 1004  |
| Active, not recruiting  | 526   |
| Enrolling by invitation | 181   |
| Withdrawn               | 107   |
| Terminated              | 74    |
| Suspended               | 27    |
| Available               | 19    |
| No longer available     | 12    |
| Approved for marketing  | 2     |

```
sns.countplot(y="Status", data=df, color="red")
```

<Axes: xlabel='count', ylabel='Status'>



**Observation:** The study status distribution reveals that most studies are still in the process of recruiting participants, indicating that the research is ongoing.

## 4.7 Cleaning Age Column

### Cleaninig Age Column

```
df.Age.unique()
```

```
array(['18 Years and older \xa0 (Adult, Older Adult)',  
      'Child, Adult, Older Adult', '18 Years to 48 Years \xa0 (Adult)',  
      '18 Years to 75 Years \xa0 (Adult, Older Adult)',  
      '18 Years to 45 Years \xa0 (Adult)',  
      '18 Years to 99 Years \xa0 (Adult, Older Adult)',  
      '18 Years to 55 Years \xa0 (Adult)',  
      '15 Years and older \xa0 (Child, Adult, Older Adult)',  
      '18 Years to 80 Years \xa0 (Adult, Older Adult)',  
      '45 Years and older \xa0 (Adult, Older Adult)',  
      '20 Years to 100 Years \xa0 (Adult, Older Adult)',  
      '8 Years to 88 Years \xa0 (Child, Adult, Older Adult)',  
      '5 Years to 65 Years \xa0 (Child, Adult, Older Adult)',  
      'up to 99 Years \xa0 (Child, Adult, Older Adult)',  
      '18 Years to 85 Years \xa0 (Adult, Older Adult)',  
      '18 Years to 65 Years \xa0 (Adult, Older Adult)',  
      'up to 29 Days \xa0 (Child)',  
      '18 Years to 70 Years \xa0 (Adult, Older Adult)',  
      '18 Years to 59 Years \xa0 (Adult)',
```

```
from string import digits  
  
def remove_digits(text):  
    return text.translate(str.maketrans('', '', digits))  
  
df["Age"] = df["Age"].apply(lambda text: remove_digits(text))  
df[['Age']].head()
```

- **from string import digits:** Imports the digits constant, which contains all digits ('0123456789').
- **remove\_digits(text):** A function that removes all digit characters from a given string text by using str.translate() to replace digits (from digits) with nothing (i.e., removes them).
- **df["Age"].apply(lambda text: remove\_digits(text)):** For each value in the "Age" column of the df DataFrame, the remove\_digits function is applied to remove any digits.
- **df[['Age']].head():** Displays the first 5 rows of the "Age" column after removing digits.



|   | Age                                  |
|---|--------------------------------------|
| 0 | Years and older (Adult, Older Adult) |
| 1 | Years and older (Adult, Older Adult) |
| 2 | Years and older (Adult, Older Adult) |
| 3 | Child, Adult, Older Adult            |
| 4 | Years to Years (Adult)               |

```
from nltk.corpus import stopwords
stopwords = stopwords.words('english')
def remove_stopwords(text):
    ... return " ".join([word for word in str(text).split() if word not in stopwords])

df["Age"] = df["Age"].apply(lambda text: remove_stopwords(text))
df[['Age']].head()
```

- **from nltk.corpus import stopwords:** Imports common English stopwords (like "the", "is", "and") from NLTK.
- **remove\_stopwords(text):** Defines a function that removes these stopwords from a given text by keeping only the words not in the stopwords list.
- **df["Age"].apply(lambda text: remove\_stopwords(text)):** Applies the remove\_stopwords function to each entry in the "Age" column.
- **df[['Age']].head():** Displays the first 5 rows of the cleaned "Age" column.

|   | Age                              |
|---|----------------------------------|
| 0 | Years older (Adult, Older Adult) |
| 1 | Years older (Adult, Older Adult) |
| 2 | Years older (Adult, Older Adult) |
| 3 | Child, Adult, Older Adult        |
| 4 | Years Years (Adult)              |

```
df.Age.unique()
```

```
array(['Years older (Adult, Older Adult)', 'Child, Adult, Older Adult',  
      'Years Years (Adult)', 'Years Years (Adult, Older Adult)',  
      'Years older (Child, Adult, Older Adult)',  
      'Years Years (Child, Adult, Older Adult)',  
      'Years (Child, Adult, Older Adult)', 'Days (Child)',  
      'Years (Child, Adult)', 'Years older (Older Adult)',  
      'Years Years (Child, Adult)', 'Years (Child)',  
      'Months older (Child, Adult, Older Adult)',  
      'Year Years (Child, Adult, Older Adult)', 'Years Years (Child)',  
      'Months Years (Child, Adult, Older Adult)', 'Minutes (Child)',  
      'Weeks Weeks (Child)', 'Year older (Child, Adult, Older Adult)',  
      'Month Years (Child, Adult, Older Adult)', 'Year Years (Child)',  
      'Year Years (Child, Adult)', 'Month Years (Child, Adult)',  
      'Month Years (Child)', 'Hours (Child)', 'Months (Child)',  
      'Months Years (Child, Adult)', 'Years Years (Older Adult)',  
      'Months older (Adult, Older Adult)', 'Months Years (Child)',  
      'Days Years (Child, Adult)', 'Month (Child)',  
      'Month older (Child, Adult, Older Adult)',  
      'Weeks Years (Child, Adult)', 'Months Months (Child)',  
      'Days older (Child, Adult, Older Adult)', 'Year (Child)'],  
      dtype=object)
```

```
df["Age"]=df["Age"].apply(lambda x:str(x).replace('Years','')if 'Years' in str(x) else str(x))  
df["Age"]=df["Age"].apply(lambda x:str(x).replace('Year','')if 'Year' in str(x) else str(x))
```

- This code removes the words **"Years"** and **"Year"** from each value in the **"Age"** column.
- It checks if "Years" or "Year" exists in the text, and if yes, replaces them with an empty string.

```
df.Age.unique()
```

```
array([' older (Adult, Older Adult)', 'Child, Adult, Older Adult',  
      ' (Adult)', ' (Adult, Older Adult)',  
      ' older (Child, Adult, Older Adult)',  
      ' (Child, Adult, Older Adult)', ' (Child, Adult, Older Adult)',  
      'Days (Child)', ' (Child, Adult)', ' older (Older Adult)',  
      ' (Child, Adult)', ' (Child)',  
      'Months older (Child, Adult, Older Adult)', ' (Child)',  
      'Months (Child, Adult, Older Adult)', 'Minutes (Child)',  
      'Weeks Weeks (Child)', 'Month (Child, Adult, Older Adult)',  
      'Month (Child, Adult)', 'Month (Child)', 'Hours (Child)',  
      'Months (Child)', 'Months (Child, Adult)', ' (Older Adult)',  
      'Months older (Adult, Older Adult)', 'Months (Child)',  
      'Days (Child, Adult)', 'Month (Child)',  
      'Month older (Child, Adult, Older Adult)', 'Weeks (Child, Adult)',  
      'Months Months (Child)', 'Days older (Child, Adult, Older Adult)'],  
      dtype=object)
```



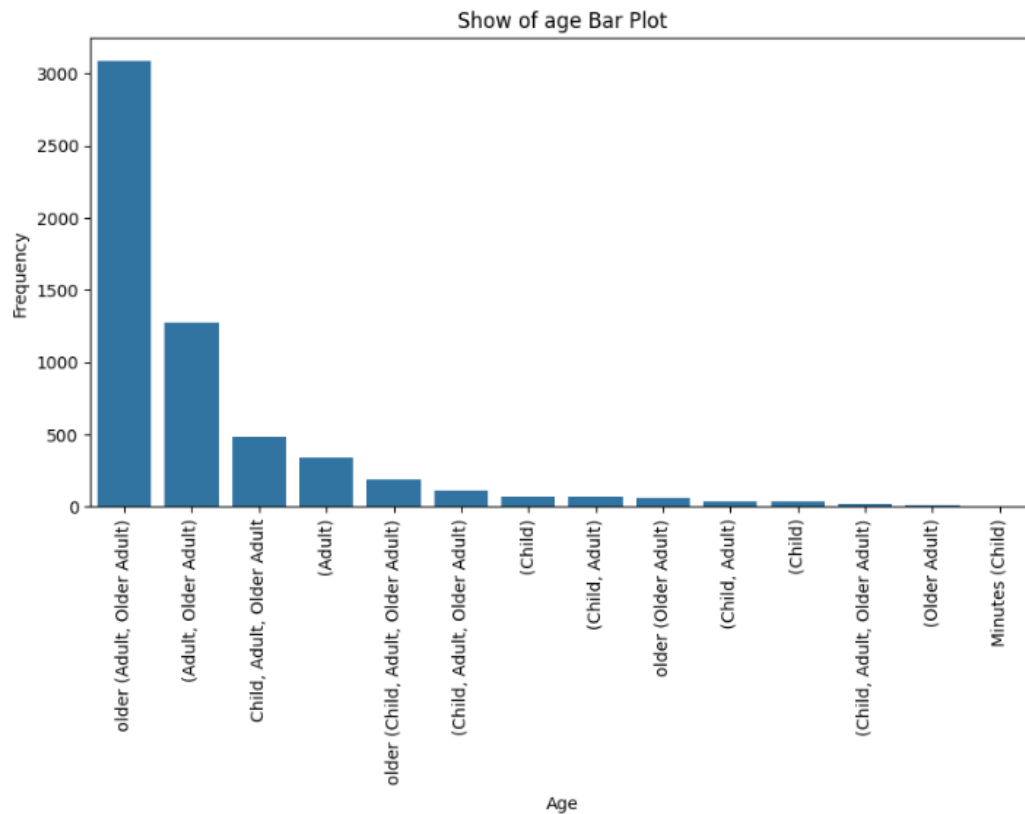
```
df["Age"]=df["Age"].apply(lambda x:str(x).replace('Months','')if 'Months' in str(x) else str(x))
df["Age"]=df["Age"].apply(lambda x:str(x).replace('Month','')if 'Month' in str(x) else str(x))
df["Age"]=df["Age"].apply(lambda x:str(x).replace('Days','')if 'Days' in str(x) else str(x))
df["Age"]=df["Age"].apply(lambda x:str(x).replace('Weeks','')if 'Weeks' in str(x) else str(x))
df["Age"]=df["Age"].apply(lambda x:str(x).replace('Hours','')if 'Hours' in str(x) else str(x))
```

- This code **removes** the words "**Months**", "**Month**", "**Days**", "**Weeks**", and "**Hours**" from each value in the "**Age**" column.
- For each word, it checks if it exists in the text and replaces it with an empty string.

```
df.Age.unique()
```

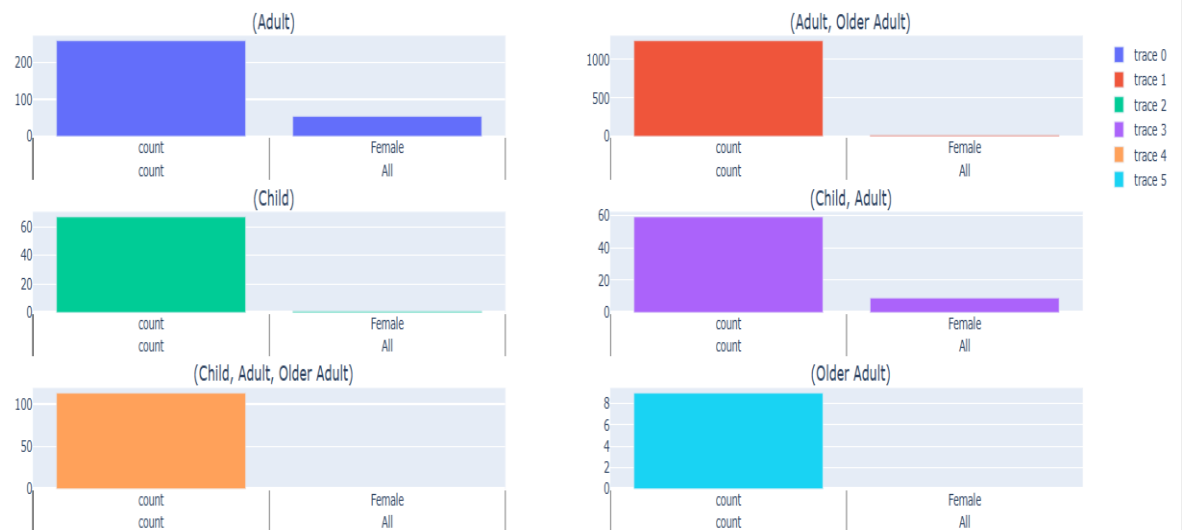
```
array([' older (Adult, Older Adult)', 'Child, Adult, Older Adult',
      ' (Adult)', ' (Adult, Older Adult)',
      ' older (Child, Adult, Older Adult)',
      ' (Child, Adult, Older Adult)', ' (Child, Adult, Older Adult)',
      ' (Child)', ' (Child, Adult)', ' older (Older Adult)',
      ' (Child, Adult)', ' (Child)', 'Minutes (Child)',
      ' (Older Adult)'], dtype=object)
```

```
plt.figure(figsize=(10,5))
sns.barplot(x=df['Age'].value_counts().index,
            y=df['Age'].value_counts().values)
plt.xlabel('Age')
plt.ylabel('Frequency')
plt.title('Show of age Bar Plot')
plt.xticks(rotation=90)
plt.show()
```



**Observation:** Adult, Older Adult age bracket are mostly studied --Child and older Adult age bracket has the lowest studies.

```
i = 0
fig = make_subplots(rows=3, cols=2, subplot_titles=list(pd.DataFrame(df.groupby(['Age'])['Gender'].value_counts().unstack().index)))
for row in range(1,4):
    for col in range(1,3):
        dt = pd.DataFrame(df.groupby(['Age'])['Gender'].value_counts().unstack().iloc[i])
        # Check if dt is a Series and convert it to DataFrame if necessary
        if isinstance(dt, pd.Series):
            dt = dt.to_frame(name='Gender')
            #This converts it to a DataFrame with 'Gender' as column name.
        fig.add_trace(go.Bar(x=dt.index, y=dt.Gender.values), row=row, col=col) #Use dt.index instead of dt.Gender.index
        i+=1
fig.show()
```



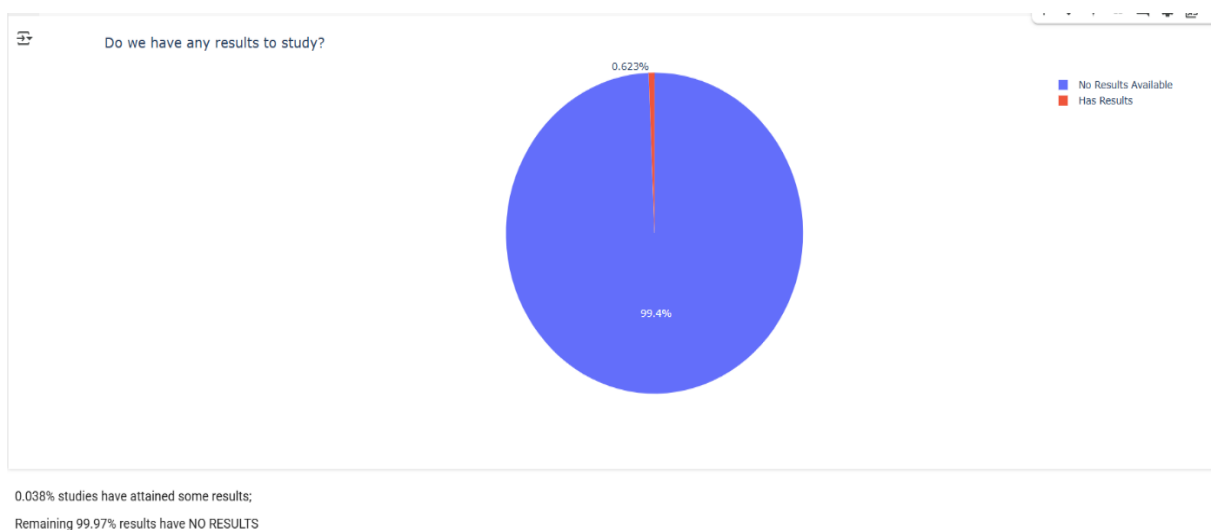
### Observations:

Most studies have taken data from All Genders;

In (Adult) and (Child, Adult) Category there is significant number of Female patients considered for the studies

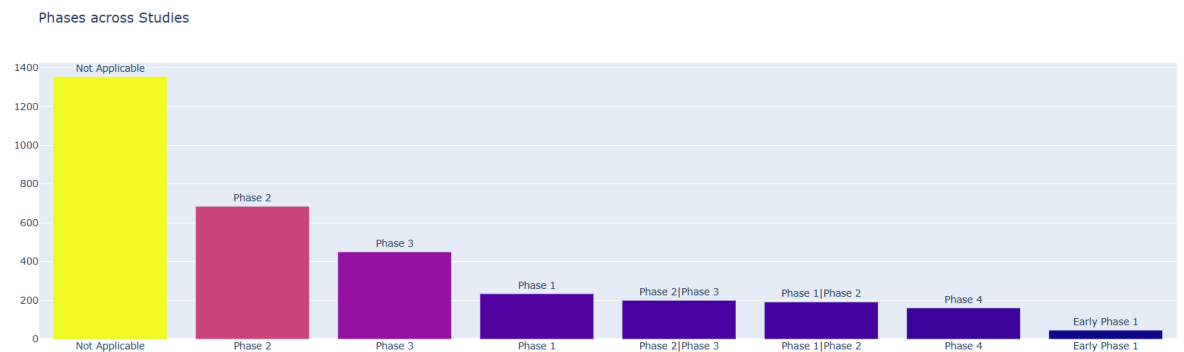
## 4.8 Exploring study results

```
import plotly.express as px
fig = px.pie(df, 'Study Results')
fig.update_layout(title='Do we have any results to study?')
fig.show()
```

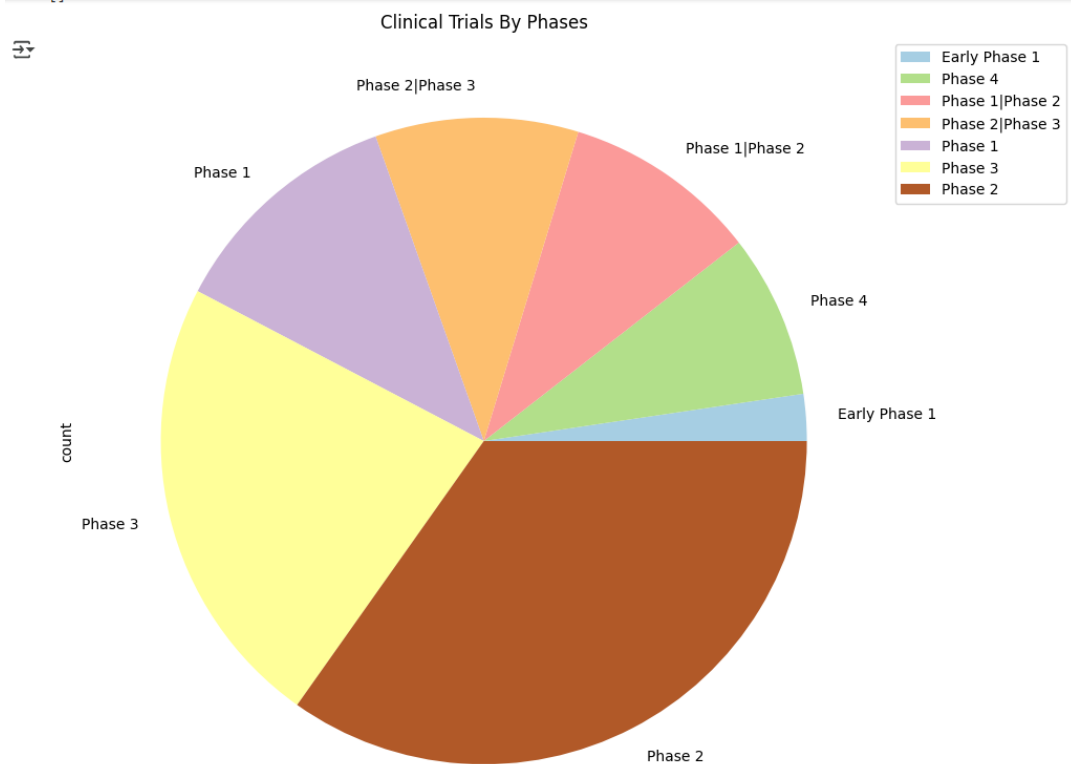


## 4.9 Exploring Study Phases

```
fig = go.Figure(go.Bar(
    x= df.groupby('Phases').agg('count')['Rank'].sort_values(ascending=False).index,
    y= df.groupby('Phases').agg('count')['Rank'].sort_values(ascending=False).values,
    text=df.groupby('Phases').agg('count')['Rank'].sort_values(ascending=False).index,
    textposition='outside',
    marker_color=df.groupby('Phases').agg('count')['Rank'].sort_values(ascending=False).values
))
fig.update_layout(title='Phases across Studies')
fig.show()
```



```
df.drop(df.index[df['Phases']=='Not Applicable'], inplace=True)
ax = df['Phases'].value_counts().sort_values().plot(kind='pie', figsize=(20,10), colormap='Paired', title='Clinical Trials By Phases')
ax.legend(bbox_to_anchor=(1.0, 1.0))
ax.plot()
```



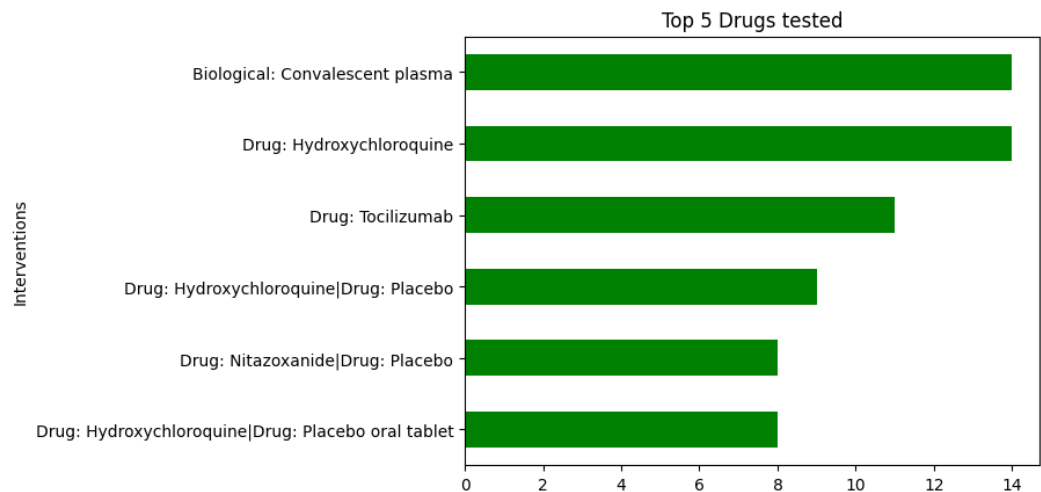
**Observation:** Most Study where applicable are in Phase 2 and Phase 3

```
df.Interventions.unique()
```

```
array(['Drug: Drug COVID19-0001-USR|Drug: normal saline',  
      'Other: Lung CT scan analysis in COVID-19 patients',  
      'Diagnostic Test: COVID 19 Diagnostic Test', ...,  
      'Biological: FluBlok|Other: Placebo',  
      'Biological: ASP2390|Biological: Placebo',  
      'Other: Antibiotic treatment|Other: No antibiotic treatment'],  
      dtype=object)
```

```
[239] interventions = df[df['Study Type']=='Interventional']  
interventions['Interventions'].value_counts().head(6).sort_values().plot(kind='barh', color='g', title='Top 5 Drugs tested')
```

```
<Axes: title={'center': 'Top 5 Drugs tested'}, ylabel='Interventions'>
```



```
df = df.ffill(axis = 1)
```

"For each row, if a cell is missing, fill it with the value from the left."

```
df.head()
```

| Rank | NCT Number | Title       | Acronym   | Status       | Study Results           | Conditions           | Interventions                                     | Outcome Measures                                  | Sponsor/Collaborators                             | ...  | Other IDs        | Start Date       | Primary Completion Date | Completion Date   | First Posted     | Results First Posted |
|------|------------|-------------|---|--------------|-------------------------|----------------------|---|---|---|--|------------------|------------------|-------------------------|-------------------|------------------|----------------------|
| 1    | 2          | NCT04595136 | Study to Evaluate the Efficacy of COVID19-0001... | COVID-19     | Not yet recruiting      | No Results Available | SARS-CoV-2 Infection                              | Drug: Drug COVID19-0001-USR Drug: normal saline   | Change on viral load results from baseline aft... | United Medical Specialties                         | COVID19-0001-USR | November 2, 2020 | December 15, 2020       | January 29, 2021  | October 20, 2020 | October 20, 2020     |
| 2    | 3          | NCT04395482 | Lung CT Scan Analysis of SARS-CoV2 Induced Lun... | TAC-COVID19  | Recruiting              | No Results Available | covid19   | Other: Lung CT scan analysis in COVID-19 patients | A qualitative analysis of parenchymal lung dam... | University of Milano Bicocca                       | TAC-COVID19      | May 7, 2020      | June 15, 2021           | June 15, 2021     | May 20, 2020     | May 20, 2020         |
| 3    | 4          | NCT04416061 | The Role of a Private Hospital in Hong Kong Am... | COVID-19     | Active, not recruiting  | No Results Available | COVID   | Diagnostic Test: COVID 19 Diagnostic Test         | Proportion of asymptomatic subjects Proportion... | Hong Kong Sanatorium & Hospital                    | RC-2020-08       | May 25, 2020     | July 31, 2020           | August 31, 2020   | June 4, 2020     | June 4, 2020         |
| 4    | 5          | NCT04395924 | Maternal-fetal Transmission of SARS-Cov-2         | TMF-COVID-19 | Recruiting              | No Results Available | Maternal Fetal Infection Transmission(COVID-19... | Diagnostic Test: Diagnosis of SARS-Cov2 by RT...  | COVID-19 by positive PCR in cord blood and / o... | Centre Hospitalier Régional d'Orléans Centre d...  | CHRO-2020-10     | May 5, 2020      | May 2021                | May 2021          | May 20, 2020     | May 20, 2020         |
| 5    | 6          | NCT04516954 | Convalescent Plasma for COVID-19 Patients         | CPCP         | Enrolling by invitation | No Results Available | COVID 19  | Biological: Convalescent COVID 19 Plasma          | Evaluate the safety Change in requirement for ... | Vinnmec Research Institute of Stem Cell and Gen... | ISC.20.11.1      | August 1, 2020   | November 30, 2020       | December 30, 2020 | August 18, 2020  | August 18, 2020      |

```
[414] df.isnull().sum()
```

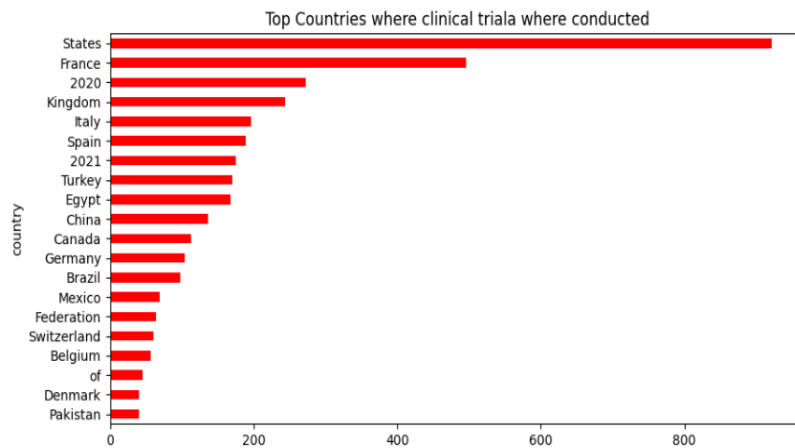


|                         |   |
|-------------------------|---|
|                         | 0 |
| Rank                    | 0 |
| NCT Number              | 0 |
| Title                   | 0 |
| Acronym                 | 0 |
| Status                  | 0 |
| Study Results           | 0 |
| Conditions              | 0 |
| Interventions           | 0 |
| Outcome Measures        | 0 |
| Sponsor/Collaborators   | 0 |
| Gender                  | 0 |
| Age                     | 0 |
| Phases                  | 0 |
| Enrollment              | 0 |
| Funded Bys              | 0 |
| Study Type              | 0 |
| Study Designs           | 0 |
| Other IDs               | 0 |
| Start Date              | 0 |
| Primary Completion Date | 0 |
| Completion Date         | 0 |
| First Posted            | 0 |
| Results First Posted    | 0 |
| Last Update Posted      | 0 |
| Locations               | 0 |
| Study Documents         | 0 |
| URL                     | 0 |

```
[415] df['country'] = [country.split()[-1] for country in df.Locations]
```

```
countries = df[df['country']!='']
countries['country'].value_counts().head(20).sort_values().plot(kind='barh', color='red', figsize=(10,5), title='Top Countries where clinical trials were conducted')
```

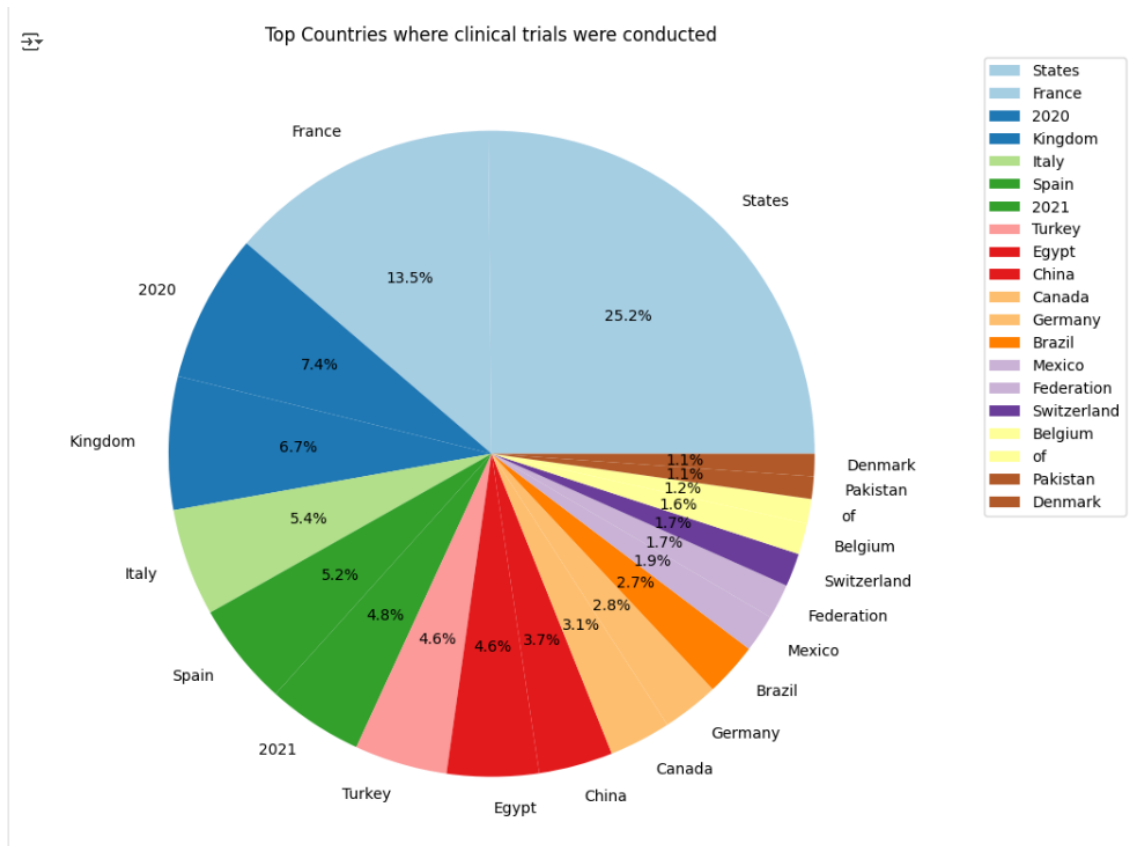
```
<Axes: title={'center': 'Top Countries where clinical trials were conducted'}, ylabel='country'>
```



```
# Filter top 20 countries by number of clinical trials
top_countries = df['country'].value_counts().head(20)

# Plot
ax = top_countries.plot(
    kind='pie',
    figsize=(10, 10),
    colormap='Paired',
    autopct='%1.1f%%', # Show percentage
    title='Top Countries where clinical trials were conducted',
    ylabel='', # Hide the y-label
    legend=False
)

# Display the plot
plt.legend(loc='upper left', bbox_to_anchor=(1.1, 1.0))
plt.tight_layout()
plt.show()
```



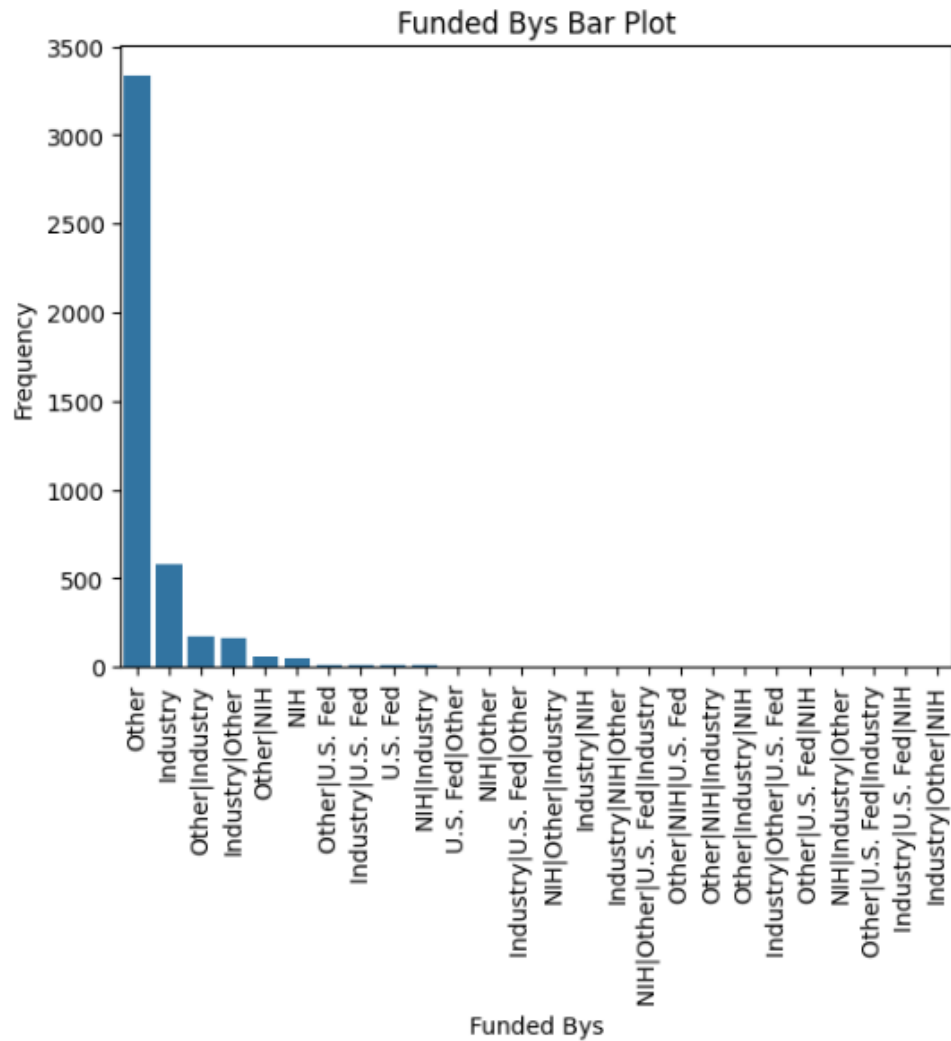
**Observation:** Most studies have taken data from State and France

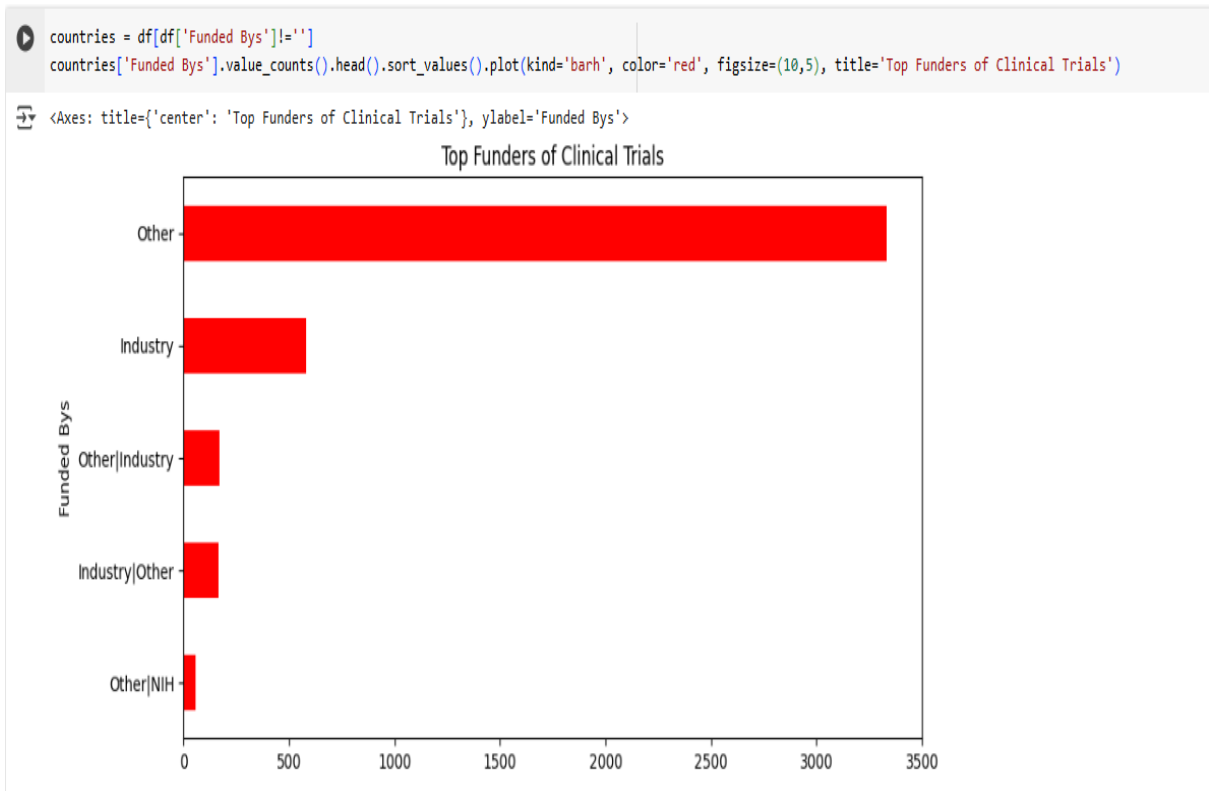


```

sns.barplot(x=df['Funded Bys'].value_counts().index,
            y=df['Funded Bys'].value_counts().values)
plt.xlabel('Funded Bys')
plt.ylabel('Frequency')
plt.title('Funded Bys Bar Plot')
plt.xticks(rotation=90)
plt.show()

```





Observation: Max Funding is by Industry



## 4.10 States Explore the Gender distribution in the studies

```
[637] dfc = df.groupby('country')
```

dfc.first()

|         | Rank | NCT Number  | Title   | Acronym      | Status             | Study Results        | Conditions                        | Interventions                                     | Outcome Measures                                  | Sponsor/Collaborators                             | ... | Other IDs              | Start Date        | Primary Completion Date | Completion Date |      |
|---------|------|-------------|---|--------------|--------------------|----------------------|-----------------------------------|---|---|---|-----|------------------------|-------------------|-------------------------|-----------------|------|
| country |      |             |   |              |                    |                      |                                   |   |   |   |     |                        |                   |                         |                 |      |
| 2020    | 36   | NCT04545498 | COVID-19 and Hereditary Metabolic Diseases        | COVID19-MH-M | Not yet recruiting | No Results Available | Covid19 Metabolism, Inborn Errors | Covid19 Metabolism, Inborn Errors                 | Frequency of MH-M imbalance triggered by COVID... | University Hospital, Lille                        | ... | 2020_52 2020-A02886-33 | January 2021      | January 2023            | January 2023    | No 2 |
| 2021    | 46   | NCT04834908 | Evaluation of Equine Antibody Treatment in Pat... | PROTECT      | Not yet recruiting | No Results Available | SARS-CoV-2 Infection              | Biological: Equine COVID-19 Antiserum Drug: ST... | Phase 1 Unexpected serious adverse events Phas... | Bharat Serums and Vaccines Limited                | ... | BSV_EQ-AB_20_08        | May 2021          | March 2022              | September 2022  |      |
| Africa  | 423  | NCT04709302 | Effects of COVID-19 on Endothelium in HIV-Posi... | ENDOCOVID    | Recruiting         | No Results Available | Covid19 HIV ART                   | Biological: COVID-19 Biological: HIV Drug: ART    | Number of patients developing Acute Respirator... | Medical University of Graz University of Oslo ... | ... | ENDOCOVID              | February 2021     | June 2022               | December 2022   | 1    |
| Albania | 73   | NCT04811391 | COVID-19 Vaccine Effectiveness in Albanian Hea... | COVEAL       | Recruiting         | No Results Available | Covid19                           | Biological: COVID-19 vaccine                      | COVID-19 vaccine effectiveness COVID-19 PCR co... | Institute of Public Health, Albania World Heal... | ... | COVE-0001              | February 19, 2021 | May 20, 2022            | May 20, 2022    | Me   |

- Shows the first row for each country group

```
dfcc=dfc.get_group('States')
```

- dfcc is a new dataframe that contains only studies from "States".

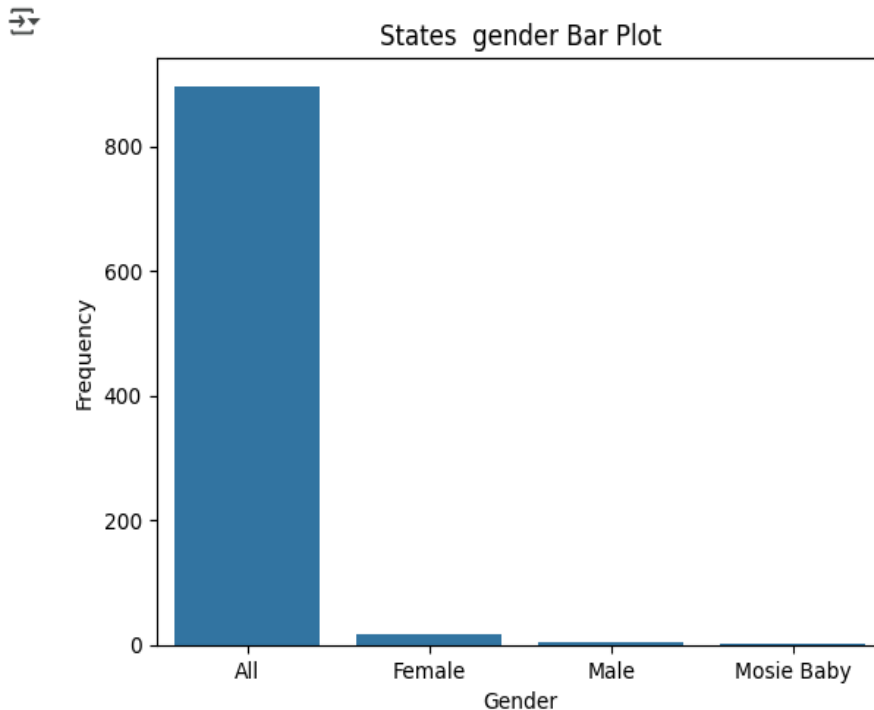
dfcc.head()

|    | Rank | NCT Number  | Title   | Acronym   | Status                 | Study Results        | Conditions | Interventions                                     | Outcome Measures                                   | Sponsor/Collaborators                             | ... | Start Date        | Primary Completion Date | Completion Date | First Posted     | Results First Posted | Last Update Posted | Locations  | Study Documents                                      |
|----|------|-------------|---|---|------------------------|----------------------|------------|---|--|---|-----|-------------------|-------------------------|-----------------|------------------|----------------------|--------------------|--|--|
| 10 | 11   | NCT04355897 | COVID-19 Plasma in Treatment of COVID-19 Patients | COVID-19 Plasma in Treatment of COVID-19 Patients | Recruiting             | No Results Available | COVID 19   | Biological: Convalescent COVID 19 Plasma          | Reduce mortality Reduce requirement for mechan...  | The Christ Hospital                               | ... | April 28, 2020    | July 2020               | August 2020     | April 21, 2020   | April 21, 2020       | May 20, 2020       | The Christ Hospital, Cincinnati, Ohio, United States | The Christ Hospital, Cincinnati, Ohio, United States |
| 12 | 13   | NCT04659759 | COVID-19 Pregnancy Related Immunological Clin...  | COVID-PRICE                                       | Recruiting             | No Results Available | Covid19    | Other: COVID-19 exposure Biological: COVID-19 ... | Maternal COVID-19 serology (IgG and IgM) Matern... | Thomas Jefferson University Nemours               | ... | November 17, 2020 | December 31, 2021       | June 30, 2022   | December 9, 2020 | December 9, 2020     | March 5, 2021      | Thomas Jefferson University Hospital, Philadel...    | Thomas Jefferson University Hospital, Philadel...    |
| 27 | 28   | NCT04424004 | MURDOCK Cabarrus County COVID-19 Prevalence an... | C3PI  | Active, not recruiting | No Results Available | COVID 19   | Other: COVID-19 PCR and serology testing          | Estimate the prevalence of COVID-19 infection ...  | Duke University North Carolina Department of H... | ... | June 9, 2020      | June 30, 2021           | June 30, 2021   | June 9, 2020     | June 9, 2020         | November 17, 2020  | Duke CTSI Translational Population Health Offi...    | Duke CTSI Translational Population Health Offi...    |
| 37 | 38   | NCT04372004 | Comparison of the Efficacy of Rapid Tests to L... | CATCH COVID-19                                    | Recruiting             | No Results Available | COVID-19   | Diagnostic Test: diagnostic tests for COVID-19... | detection of viral infection using serology an...  | Texas Cardiac Arrhythmia Research Foundation      | ... | May 8, 2020       | May 2021                | June 2021       | May 1, 2020      | May 1, 2020          | August 11, 2020    | Texas Cardiac Arrhythmia Institute, Austin, Te...    | Texas Cardiac Arrhythmia Institute, Austin, Te...    |
| 40 | 41   | NCT04412486 | COVID-19 Convalescent Plasma (CCP) Transfusion    | COVID-19 Convalescent Plasma (CCP) Transfusion    | Recruiting             | No Results Available | COVID-19   | Biological: COVID Convalescent Plasma             | Change in PaO2/FiO2 after CCP transfusion Cha...   | Galen D. Marshall, Jr., MD PhD University of M... | ... | June 1, 2020      | May 31, 2022            | May 31, 2022    | June 2, 2020     | June 2, 2020         | July 2, 2020       | University of Mississippi Medical Center, Jack...    | University of Mississippi Medical Center, Jack...    |

```

sns.barplot(x=dfcc['Gender'].value_counts().index,
            y=dfcc['Gender'].value_counts().values)
plt.xlabel('Gender')
plt.ylabel('Frequency')
plt.title('States gender Bar Plot')
plt.show()

```



## 5. Conclusion

### 1. Top Countries for Clinical Trials

- The **United States (States)** conducted the maximum number of COVID-19 clinical trials.
- **France, United Kingdom, Italy, and Spain** were also major contributors.
- This shows that developed countries were leading clinical research during the pandemic.

### 2. Funding Sources

- Most clinical trials were funded under the category "**Other**".
- The second largest funding source was "**Industry**" (pharmaceutical companies, biotech firms).
- Combination funding like **Industry|Other** also appeared but was much less frequent.

- **Conclusion:** Clinical trials were majorly **privately or independently funded** rather than solely by governments or official health bodies.

### **3. Gender Distribution in Studies (for 'States')**

- A majority of clinical trials were open to **All Genders** (both male and female participants).
- Some studies focused specifically on **Male** or **Female** participants, but these were much fewer.
- **Conclusion:** Researchers mostly designed studies to include **both genders**, aiming for a broader understanding of COVID-19's impact.

### **4. Timeline Observations (if any)**

- A lot of trials spiked around **2020-2021**, which aligns perfectly with the global emergency and vaccine development phases.