

```
# Importing libraries
```

```
import pandas as pd
```

```
import seaborn as sns
```

```
import matplotlib.pyplot as plt
```

```
import urllib.request, json
```

```
In [2]:# Several methods to retrieve JSON
```

```
# df=pd.read_json('https://www....json')
```

```
# data = json.load('/home/.../json')
```

```
# with urllib.request.urlopen("https://.../json") as url:
```

```
# data = json.load(url)
```

```
# print(data)
```

```
# with open('/.../Covid19.json', 'r') as myfile:
```

```
# data=myfile.read()
```

```
# # parse file
```

```
# obj = json.loads(data)
```

```
In [3]:df=pd.read_json('/.../Covid19.json')
```

```
In [4]:#### The dataset below is as of 10-02-2023.
```

```
In [5]:# Showing information form JSON file, Here is the dataset we have to work on
```

```
df
```

```
Out[5]:
```

	sno	state_name	active	positive	cured	death	new_active	new_positive	new_cured
0	1	dolnośląskie	0	10751	10622	129	0	10751	10622
1	2	kujawsko-pomorskie	4	2339098	2324361	14733	3	2339098	2324361
2	3	lubelskie	0	66891	66595	296	0	66891	66595
3	4	łódzkie	0	746100	738065	8035	0	746100	738065
4	5	małopolskie	3	851428	839122	12303	3	851428	839122
5	6	mazowieckie	4	99373	98187	1182	6	99375	98189
6	7	opolskie	6	1177805	1163653	14146	6	1177806	1163654
7	8	podkarpackie	0	11591	11587	4	0	11591	11587
8	9	podlaskie	38	2007692	1981131	26523	62	2007718	1981157
9	10	pomorskie	32	259191	255146	4013	40	259199	255150
10	11	śląskie	233	1278016	1266736	11047	268	1278074	1266763
11	12	świętokrzyskie	49	1056797	1046034	10714	47	1056803	1046041
12	13	warmińsko-mazurskie	60	312858	308584	4214	100	312900	308584
13	14	wielkopolskie	41	479527	474701	4785	43	479533	474705
14	15	zachodniopomorskie	2	442579	437246	5331	4	442581	437248

```
In [6]:# In this file are 15 rows and 14 columns
```

```
df.shape
```

```
Out[6]:(15, 14)
```

```
In [7]:# In this file type of data present in the columns(int,object)
```

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 15 entries, 0 to 14
Data columns (total 14 columns):
#   Column                Non-Null Count  Dtype
---  -
0   sno                    15 non-null    int64
1   state_name            15 non-null    object
2   active                15 non-null    int64
3   positive              15 non-null    int64
4   cured                 15 non-null    int64
5   death                 15 non-null    int64
6   new_active            15 non-null    int64
7   new_positive          15 non-null    int64
8   new_cured             15 non-null    int64
9   new_death             15 non-null    int64
10  death_reconsille      15 non-null    object
11  total                  15 non-null    object
12  state_code            15 non-null    int64
13  actualdeath24hrs      15 non-null    int64
dtypes: int64(11), object(3)
memory usage: 1.8+ KB
```

In ... # Viewing the descriptive statistics of the data like mean, std deviation, min and max values present in the df.describe()

```
Out[8]:
```

	sno	active	positive	cured	death	new_active	new_positiv
count	15.000000	15.000000	1.500000e+01	1.500000e+01	15.000000	15.000000	1.500000e+01
mean	8.000000	31.466667	7.426465e+05	7.347847e+05	7830.333333	38.800000	7.426565e+05
std	4.472136	59.562532	7.198113e+05	7.132340e+05	7272.052763	70.127639	7.198153e+05
min	1.000000	0.000000	1.075100e+04	1.062200e+04	4.000000	0.000000	1.075100e+04
25%	4.500000	1.000000	1.792820e+05	1.766665e+05	2597.500000	1.500000	1.792870e+05
50%	8.000000	4.000000	4.795270e+05	4.747010e+05	5331.000000	6.000000	4.795330e+05
75%	11.500000	39.500000	1.117301e+06	1.104844e+06	11675.000000	45.000000	1.117304e+06
max	15.000000	233.000000	2.339098e+06	2.324361e+06	26523.000000	268.000000	2.339098e+06

In [9]:# Displaying all the column names present in data
df.columns

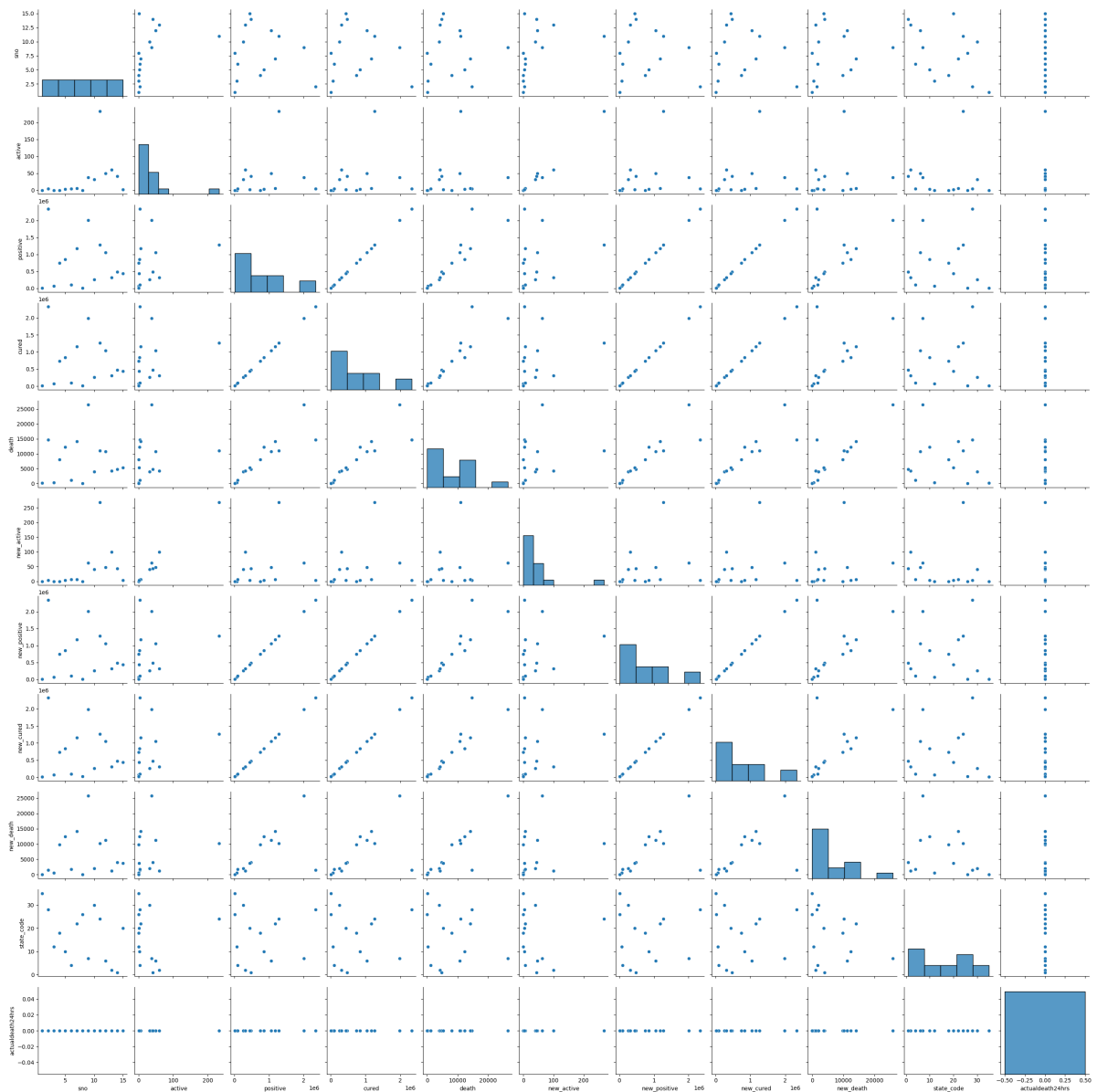
```
Out[9]:Index(['sno', 'state_name', 'active', 'positive', 'cured', 'death',
              'new_active', 'new_positive', 'new_cured', 'new_death',
              'death_reconsille', 'total', 'state_code', 'actualdeath24hrs'],
              dtype='object')
```

In [10]:# Display the last 6 rows
df.tail(6)

```
Out[10]:
```

	sno	state_name	active	positive	cured	death	new_active	new_positive	new_cui
9	10	pomorskie	32	259191	255146	4013	40	259199	2551
10	11	śląskie	233	1278016	1266736	11047	268	1278074	12667
11	12	świętokrzyskie	49	1056797	1046034	10714	47	1056803	10460
12	13	warmińsko-mazurskie	60	312858	308584	4214	100	312900	3085
13	14	wielkopolskie	41	479527	474701	4785	43	479533	4747
14	15	zachodniopomorskie	2	442579	437246	5331	4	442581	4372

```
In [11]:# The total number of active cases in Poland 472
df['active'].sum(axis = 0)
Out[11]:472
In [12]:# The total number of deaths in Poland 117.455
df['death'].sum(axis = 0)
Out[12]:117455
In [13]:# The total number of positive in Poland 11.139.697
df['positive'].sum(axis = 0)
Out[13]:11139697
In [14]:# The total number of cured in Poland 11.021.770
df['cured'].sum(axis = 0)
Out[14]:11021770
In [15]:# The total number of new_active cases in Poland 582
df['new_active'].sum(axis = 0)
Out[15]:582
In [16]:# The total number of new_deaths in Poland 98.483
df['new_death'].sum(axis = 0)
Out[16]:98483
In [17]:# The total number of new_positive in Poland 11.139.848
df['new_positive'].sum(axis = 0)
Out[17]:11139848
In [18]:# The total number of new_cured in Poland 11.027.971
df['new_cured'].sum(axis = 0)
Out[18]:11027971
In [19]:### Data Visualization
In [20]:# Plotting scatter plots of all data
sns.pairplot(data=df)
# plt.show()
Out[20]:<seaborn.axisgrid.PairGrid at 0x7fc8ad3e5900>
```



In [21]:# Storing total cases

```
total_df = df.sum()
```

```
total_df
```

Out[21]:sno

120

state_name dolnośląskie kujawsko-pomorskie lubelskie łódzkie...

active 472

positive 11139697

cured 11021770

death 117455

new_active 582

new_positive 11139848

new_cured 11027971

new_death 98483

death_reconsille

total

state_code 245

actualdeath24hrs 0

dtype: object

In [... # Chart visualization

```
my_data = [472,117455,11139697,11021770]
```

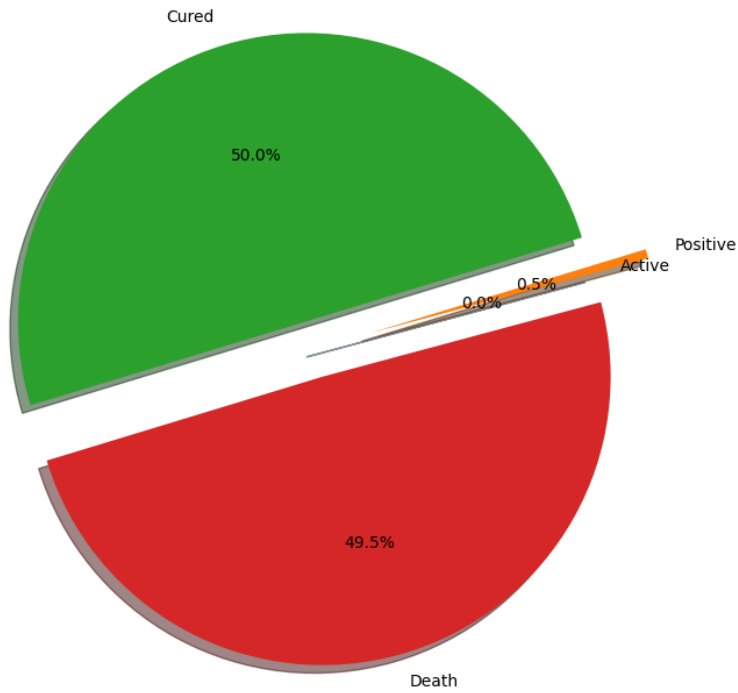
```
my_labels = 'Active','Positive','Cured','Death'
```

```
my_explode = (0,0.2,0.1,0.1)
```

```
fig1, ax1 = plt.subplots(figsize=(13, 8))
```

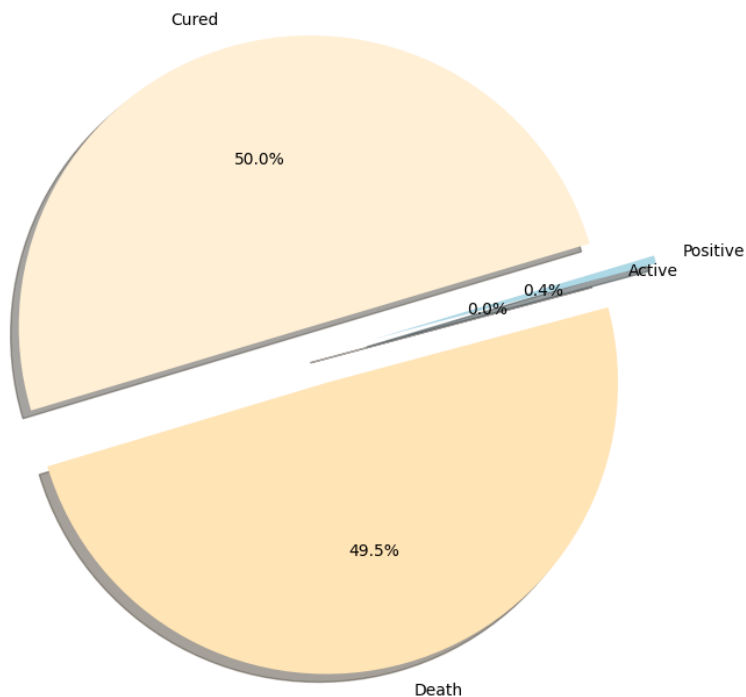
```
plt.pie(my_data, labels=my_labels, autopct='%1.1f%%', startangle=15, shadow = True, explode=my_ex
```

```
plt.axis('equal')
plt.show()
```



In [...# Chart visualization

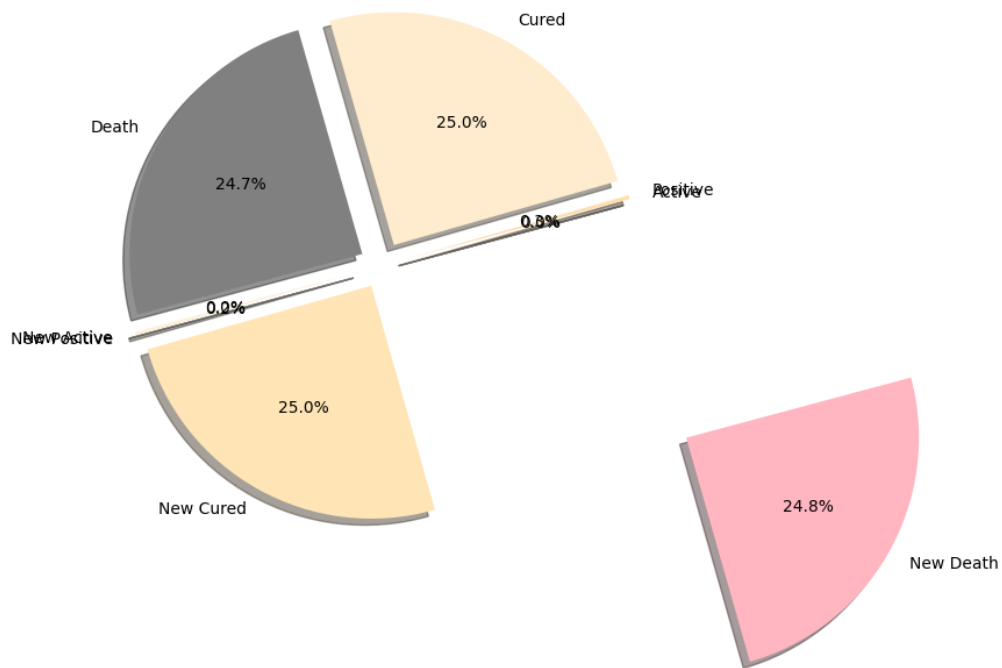
```
my_data = [582,98483,11139848,11027971]
my_labels = 'Active','Positive','Cured','Death'
my_explode = (0,0.2,0.1,0.1)
my_colors = ['blanchedalmond','lightblue','papayawhip','moccasin']
fig1, ax1 = plt.subplots(figsize=(13, 8))
plt.pie(my_data, labels=my_labels, autopct='%1.1f%%', startangle=15, shadow = True, colors=my_color:
plt.axis('equal')
plt.show()
```



In [...# Chart visualization

```
my_data = [472,117455,11139697,11021770,582,98483,11139848,11027971]
my_labels = 'Active','Positive','Cured','Death','New Active','New Positive','New Cured','New Death'
my_explode = (0.1,0.1,0.1,0.1,0.1,0.1,0.1,1.5)
my_colors = ['gray','navajowhite','blanchedalmond','grey','lightblue','papayawhip','moccasin','lightpink']
fig1, ax1 = plt.subplots(figsize=(13, 8))
```

```
plt.pie(my_data, labels=my_labels, autopct='%1.1f%%', startangle=15, shadow = True, colors=my_colors)
plt.axis('equal')
plt.show()
```

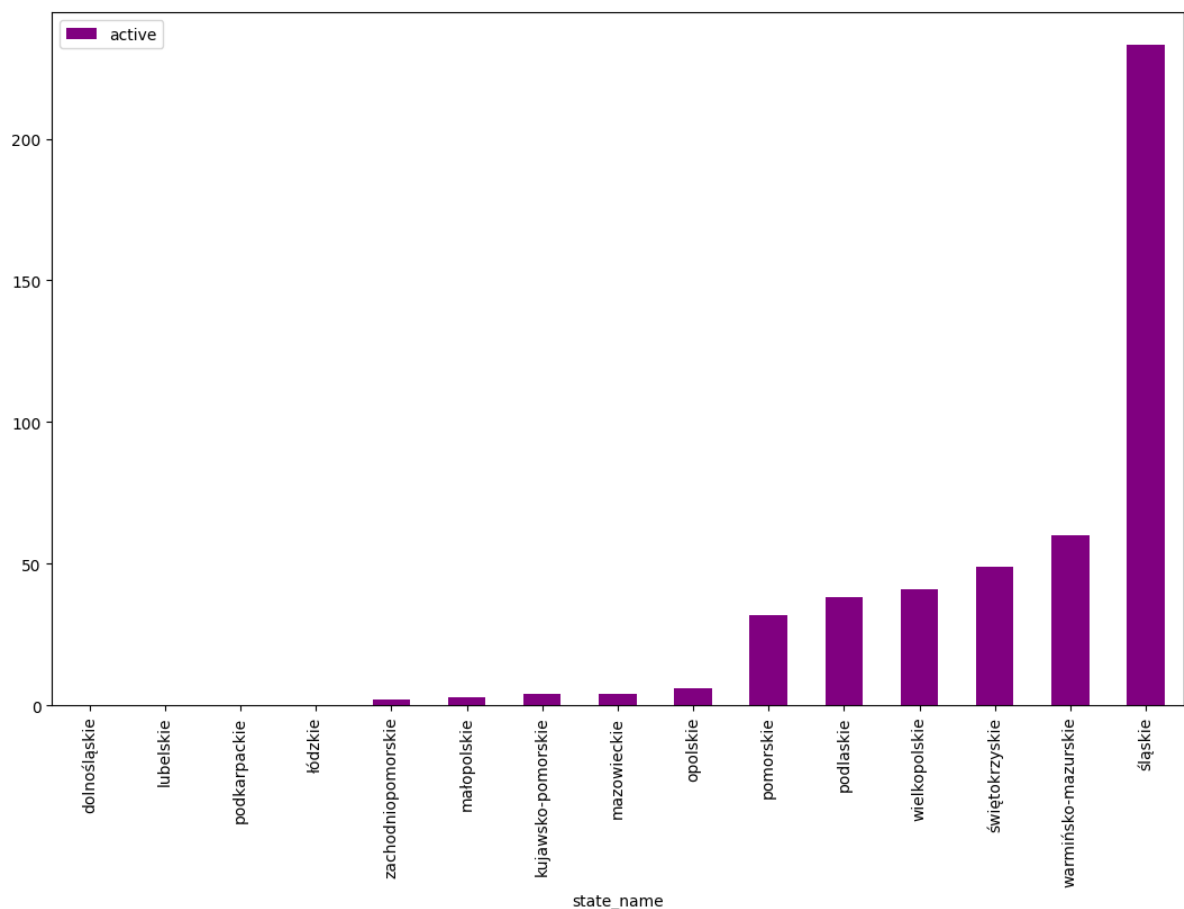


In [25]: # Figure size

```
plt.rcParams['figure.figsize']=(13,8)
```

In [2...]: # We get a picture of the states in an increasing order based on their active level of cases.

```
df[["state_name", 'active']].groupby(["state_name"]).mean().sort_values(by='active').plot.bar(color='purple')
plt.show()
```

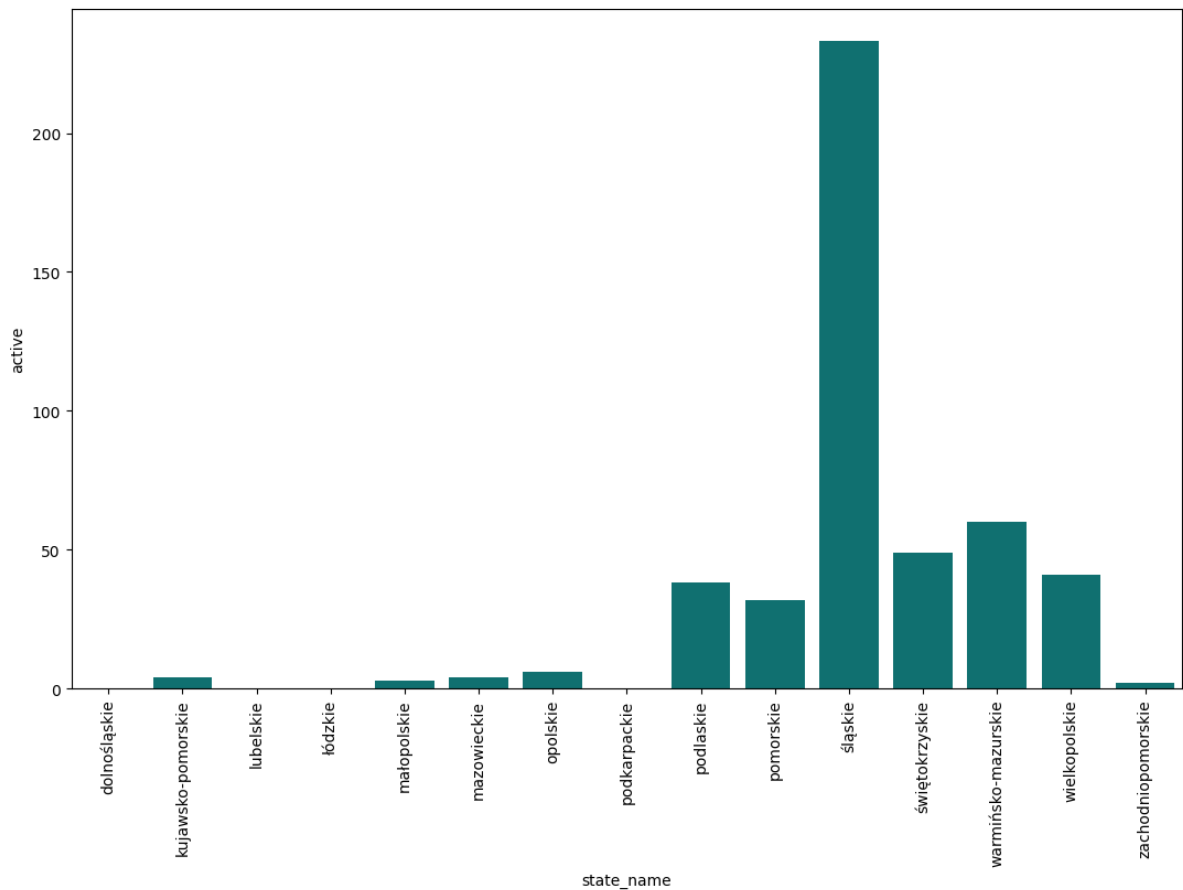


In [27]: # Number of active cases

```
plt.figure(figsize=(13, 8))
```

```
plt.xticks(rotation=90)
```

```
sns.barplot(x='state_name',y='active',color='teal',data=df);
plt.show()
```



```
In [28]:df.columns
```

```
Out[28]:Index(['sno', 'state_name', 'active', 'positive', 'cured', 'death',
               'new_active', 'new_positive', 'new_cured', 'new_death',
               'death_reconsille', 'total', 'state_code', 'actualdeath24hrs'],
              dtype='object')
```

```
In [29]:df.drop(['sno','state_code'],axis=1,inplace=True)
```

```
In [30]:df.head(5)
```

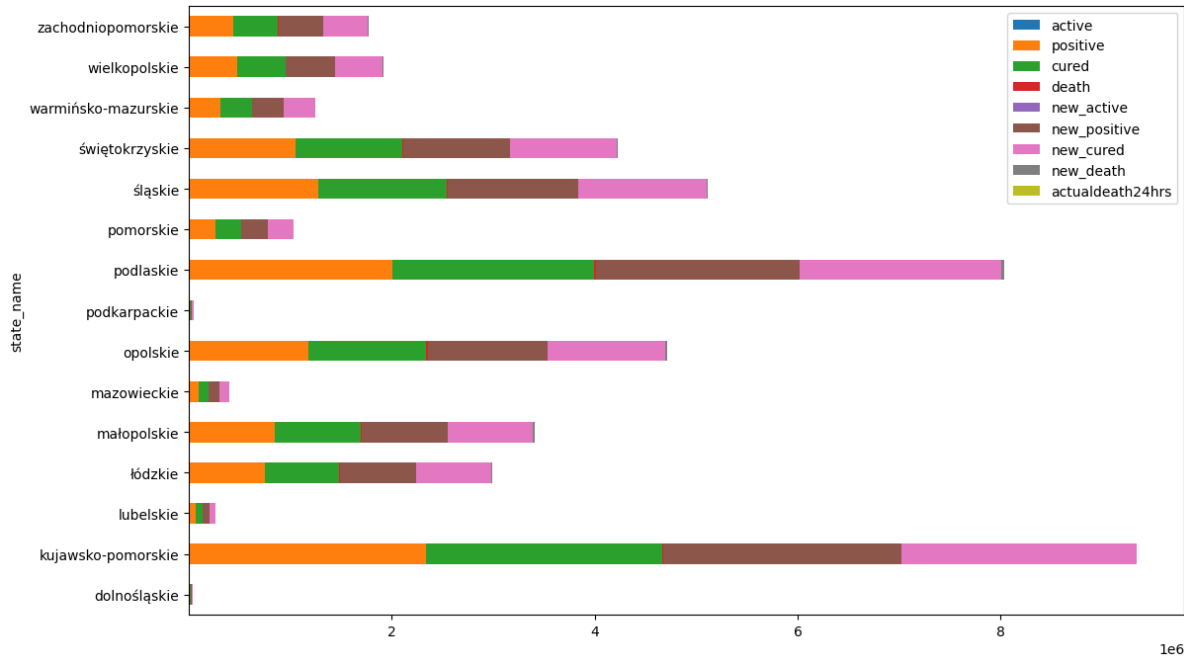
```
Out[30]:
```

	state_name	active	positive	cured	death	new_active	new_positive	new_cured	new_deat
0	dolnośląskie	0	10751	10622	129	0	10751	10622	1
1	kujawsko-pomorskie	4	2339098	2324361	14733	3	2339098	2324362	147
2	lubelskie	0	66891	66595	296	0	66891	66595	52
3	łódzkie	0	746100	738065	8035	0	746100	738065	985
4	małopolskie	3	851428	839122	12303	3	851428	839122	1244

```
In [31]:df=df.set_index('state_name')
```

```
In [37]:# Stacked bar plot
```

```
df.plot.barh(stacked=True,figsize=(13,8))
plt.show()
```



```
In [40]:# Bar plot for active and death cases
df1=df[['active', 'death']]
df1.plot.barh(color={"active": "black", "death": "orange"},figsize=(13,8))
plt.show()
```

