

# CA-COVID

April 24, 2021

```
[1]: #!/usr/bin/env python

'''
This iPython Notebook Visualized COVID-19 data from the data.cdc.gov APIs
Dataset Name: COVID-19 Case Surveillance Public Use Data with Geography
Dataser URL: https://data.cdc.gov/Case-Surveillance/
→COVID-19-Case-Surveillance-Public-Use-Data-with-Ge/n8mc-b4w4
'''

# import all the required libraries to use Panda and Plotly for data_
→visualization
import pandas as pd
import os
from sodapy import Socrata
import numpy as np
import matplotlib
import cufflinks as cf
import plotly
import plotly.offline as py
import plotly.graph_objs as go
pd.options.plotting.backend = "plotly"
cf.go_offline()

# Unauthenticated client only works with public data sets. Note 'None'
# in place of application token, and no username or password:
# client = Socrata# ("data.cdc.gov", None)

# Example authenticated client (needed for non-public datasets):
client = Socrata("data.cdc.gov", os.environ["SOCRATA_APP_TOKEN"])

# Get records where Residential State is California, returned as JSON from API /
→ converted to Python list of
# dictionaries by sodapy.
results = client.get("n8mc-b4w4", res_state="CA", limit="1000000")

# Convert result to pandas DataFrame
results_df=pd.DataFrame.from_records(results)
```

```
# Print result dataframe from the obtained data for COVID-19 cases in California
results_df
```

```
[1]:
```

	case_month	res_state	state_fips_code	res_county	county_fips_code	\
0	2020-12	CA	06	NA	NA	
1	2020-12	CA	06	NA	NA	
2	2020-12	CA	06	NA	NA	
3	2020-03	CA	06	BUTTE	06007	
4	2020-03	CA	06	BUTTE	06007	
...	...	...	...	...	...	
999995	2021-01	CA	06	ORANGE	06059	
999996	2021-01	CA	06	ORANGE	06059	
999997	2021-01	CA	06	ORANGE	06059	
999998	2021-01	CA	06	ORANGE	06059	
999999	2021-01	CA	06	ORANGE	06059	

	age_group	sex	race	ethnicity	case_onset_interval	\
0	Missing	NA	NA	NA	0.0	
1	Missing	NA	NA	NA	NaN	
2	NA	NA	NA	NA	0.0	
3	NA	NA	NA	NA	0.0	
4	NA	NA	NA	NA	0.0	
...	...	...	...	...	...	
999995	18 to 49 years	Female	Unknown	Unknown	NaN	
999996	18 to 49 years	Female	Unknown	Unknown	NaN	
999997	18 to 49 years	Female	Unknown	Unknown	NaN	
999998	18 to 49 years	Female	Unknown	Unknown	0.0	
999999	18 to 49 years	Female	Unknown	Unknown	NaN	

	process	exposure_yn	current_status	symptom_status	\
0	Missing	Missing	Laboratory-confirmed case	Symptomatic	
1	Missing	Missing	Laboratory-confirmed case	Unknown	
2	Missing	Missing	Laboratory-confirmed case	Symptomatic	
3	Missing	Missing	Laboratory-confirmed case	Symptomatic	
4	Missing	Missing	Laboratory-confirmed case	Symptomatic	
...	...	...	...	...	
999995	Missing	Missing	Laboratory-confirmed case	Unknown	
999996	Missing	Missing	Laboratory-confirmed case	Unknown	
999997	Missing	Missing	Laboratory-confirmed case	Unknown	
999998	Missing	Missing	Laboratory-confirmed case	Symptomatic	
999999	Missing	Missing	Laboratory-confirmed case	Unknown	

	hosp_yn	icu_yn	death_yn	case_positive_specimen	\
0	Yes	Missing	NA	NaN	
1	No	Missing	NA	NaN	
2	Yes	Unknown	NA	NaN	
3	No	Missing	No	NaN	

4	Yes	No	No		NaN
...	...	...	...	...	
999995	Missing	Missing	No		NaN
999996	Missing	Missing	No		NaN
999997	Missing	Missing	No		NaN
999998	No	Missing	No		NaN
999999	Missing	Missing	No		NaN

	underlying_conditions_yn
0	NaN
1	NaN
2	NaN
3	NaN
4	NaN
...	...
999995	NaN
999996	NaN
999997	NaN
999998	NaN
999999	NaN

[1000000 rows x 19 columns]

```
[2]: # select county column from the result dataframe
cases_by_county = pd.concat([results_df.res_county])
cases_by_county
```

```
[2]: 0          NA
1          NA
2          NA
3      BUTTE
4      BUTTE
...
999995  ORANGE
999996  ORANGE
999997  ORANGE
999998  ORANGE
999999  ORANGE
Name: res_county, Length: 1000000, dtype: object
```

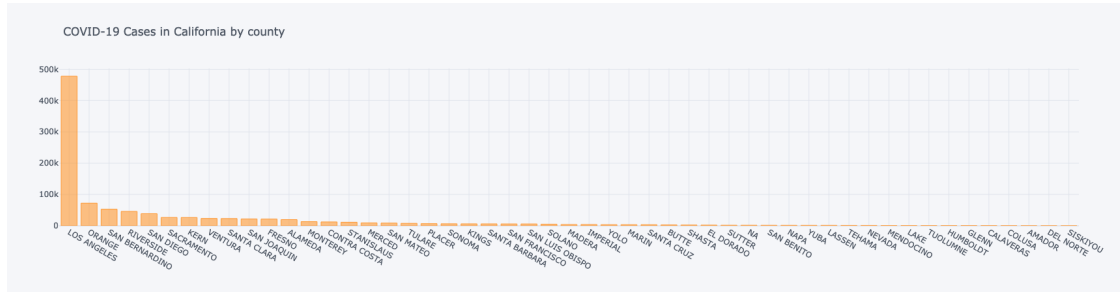
```
[3]: # count the number of cases by county
cases_by_county.value_counts()
```

```
[3]: LOS ANGELES      477821
ORANGE                72427
SAN BERNARDINO        52974
RIVERSIDE             45910
```

SAN DIEGO	38638
SACRAMENTO	26601
KERN	26458
VENTURA	23460
SANTA CLARA	23119
SAN JOAQUIN	21794
FRESNO	21429
ALAMEDA	19781
MONTEREY	13333
CONTRA COSTA	12523
STANISLAUS	11417
MERCED	9079
SAN MATEO	8615
TULARE	7856
PLACER	6859
SONOMA	6454
KINGS	6272
SANTA BARBARA	6174
SAN FRANCISCO	6039
SAN LUIS OBISPO	5939
SOLANO	4956
MADERA	4047
IMPERIAL	4036
YOLO	3570
MARIN	3551
SANTA CRUZ	3487
BUTTE	2747
SHASTA	2462
EL DORADO	2203
SUTTER	1849
NA	1813
SAN BENITO	1649
NAPA	1539
YUBA	1484
LASSEN	1483
TEHAMA	1240
NEVADA	1185
MENDOCINO	841
LAKE	805
TUOLUMNE	682
HUMBOLDT	681
GLENN	665
CALAVERAS	627
COLUSA	510
AMADOR	351
DEL NORTE	318
SISKIYOU	247

Name: res\_county, dtype: int64

```
[4]: # graph the number of cases by county
cases_by_county.value_counts().plot(kind="bar",title="COVID-19 Cases in_
↳California by county")
```



```
[5]: # We would create a Panda DataFrame that would allow us to use the county name_
↳as an index.
tmp = pd.DataFrame({'cases_by_county':cases_by_county.value_counts()})
tmp
```

```
[5]:
```

	cases_by_county
LOS ANGELES	477821
ORANGE	72427
SAN BERNARDINO	52974
RIVERSIDE	45910
SAN DIEGO	38638
SACRAMENTO	26601
KERN	26458
VENTURA	23460
SANTA CLARA	23119
SAN JOAQUIN	21794
FRESNO	21429
ALAMEDA	19781
MONTEREY	13333
CONTRA COSTA	12523
STANISLAUS	11417
MERCED	9079
SAN MATEO	8615
TULARE	7856
PLACER	6859
SONOMA	6454
KINGS	6272
SANTA BARBARA	6174
SAN FRANCISCO	6039
SAN LUIS OBISPO	5939

SOLANO	4956
MADERA	4047
IMPERIAL	4036
YOLO	3570
MARIN	3551
SANTA CRUZ	3487
BUTTE	2747
SHASTA	2462
EL DORADO	2203
SUTTER	1849
NA	1813
SAN BENITO	1649
NAPA	1539
YUBA	1484
LASSEN	1483
TEHAMA	1240
NEVADA	1185
MENDOCINO	841
LAKE	805
TUOLUMNE	682
HUMBOLDT	681
GLENN	665
CALAVERAS	627
COLUSA	510
AMADOR	351
DEL NORTE	318
SISKIYOU	247

```
[6]: # Remove the cases where we don't have the county information
# Select data where the index != NA
cases_by_county_df = tmp[tmp.index != "NA"]
# sort the results in ascending order by the number of cases
cases_by_county_df = cases_by_county_df.sort_values(by='cases_by_county',
↪ascending=True)
cases_by_county_df
```

```
[6]: cases_by_county
SISKIYOU      247
DEL NORTE     318
AMADOR        351
COLUSA        510
CALAVERAS     627
GLENN         665
HUMBOLDT      681
TUOLUMNE     682
LAKE          805
MENDOCINO     841
```

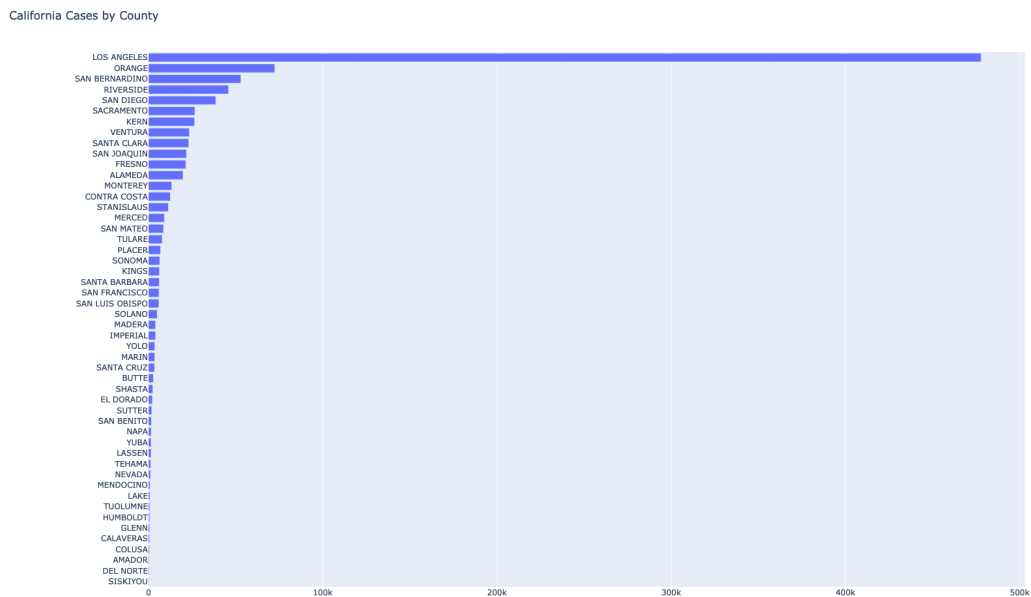
NEVADA	1185
TEHAMA	1240
LASSEN	1483
YUBA	1484
NAPA	1539
SAN BENITO	1649
SUTTER	1849
EL DORADO	2203
SHASTA	2462
BUTTE	2747
SANTA CRUZ	3487
MARIN	3551
YOLO	3570
IMPERIAL	4036
MADERA	4047
SOLANO	4956
SAN LUIS OBISPO	5939
SAN FRANCISCO	6039
SANTA BARBARA	6174
KINGS	6272
SONOMA	6454
PLACER	6859
TULARE	7856
SAN MATEO	8615
MERCED	9079
STANISLAUS	11417
CONTRA COSTA	12523
MONTEREY	13333
ALAMEDA	19781
FRESNO	21429
SAN JOAQUIN	21794
SANTA CLARA	23119
VENTURA	23460
KERN	26458
SACRAMENTO	26601
SAN DIEGO	38638
RIVERSIDE	45910
SAN BERNARDINO	52974
ORANGE	72427
LOS ANGELES	477821

```
[7]: # Create a Plotly Graph Object of type Bar and use the index on the Y axis and
      ↪ the value on the x axis
data = [go.Bar(
            y = cases_by_county_df.index,
            x = cases_by_county_df.cases_by_county,
            orientation='h'
```

```

    )]
# create the Plotly layout object
layout = go.Layout(
    height = 1000,
    margin=go.layout.Margin(l=300),
    title = "California Cases by County"
)
# create a Plotly figure object
fig = go.Figure(data=data,layout=layout)
# call the Plotly iplot() and pass the Figure object
py.iplot(fig)

```



[8]: *# In the next steps lets find the number of reported cases by county and race*

```

[9]: """
Below is the information related to the 'race' field of the dataset
From https://data.cdc.gov/Case-Surveillance/
↳ COVID-19-Case-Surveillance-Public-Use-Data-with-Ge/n8mc-b4w4
Race [American Indian/Alaska Native; Asian; Black; Multiple/Other; Native
↳ Hawaiian/Other Pacific Islander; White; Unknown; Missing; NA, if value
↳ suppressed for privacy protection.]
"""
df = pd.DataFrame({'county':results_df.res_county,'race':results_df.race})
df

```



```
[9]:      county      race
0         NA        NA
1         NA        NA
2         NA        NA
3        BUTTE        NA
4        BUTTE        NA
...
999995  ORANGE  Unknown
999996  ORANGE  Unknown
999997  ORANGE  Unknown
999998  ORANGE  Unknown
999999  ORANGE  Unknown

[1000000 rows x 2 columns]
```

```
[10]: # see all the possible race values
race_values = list()
'''
iterate over the series returned by value_counts() and convert the results to a
↳ list
'''
for key,value in pd.concat([df.race]).value_counts().items():
    race_values.append(key)
race_values
```

```
[10]: ['NA',
        'Unknown',
        'White',
        'Multiple/Other',
        'Missing',
        'Asian',
        'Black',
        'Native Hawaiian/Other Pacific Islander',
        'American Indian/Alaska Native']
```

```
[11]: # collect the cases by race and by county.
# Lets use the race as the key in a dictionary to store the results
race_df_dic = dict()
ignore_values = ['NA', 'Unknown', 'Missing'] # list of race values to ignore
race_values.sort()
for race in race_values:
    if race not in ignore_values:
        ''' below an example on how to get all the cases where the race was
        ↳ White:
            df[df["race"] == "White"]
        '''
        race_df_dic[race] = df[df['race'] == race]
```

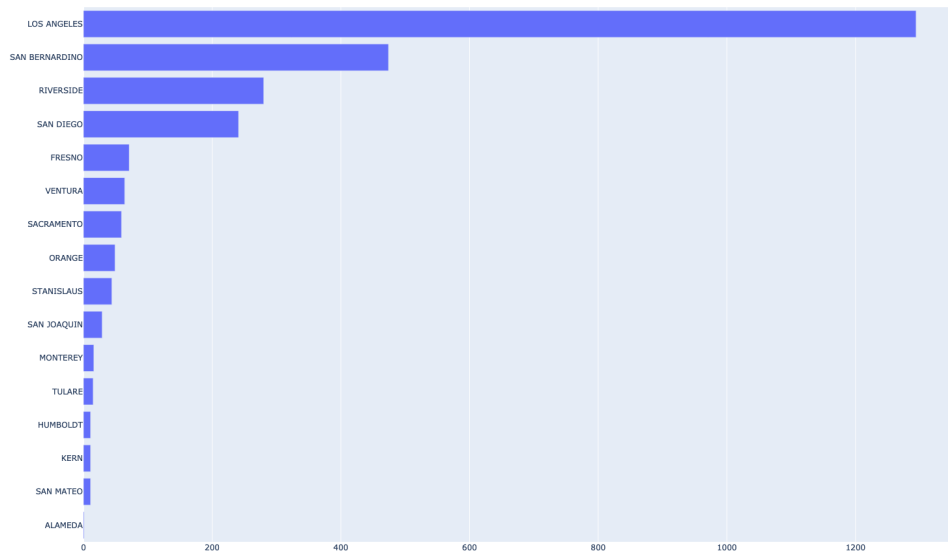
```

# get the number of cases by race by county
race_bycounty_dic = dict()

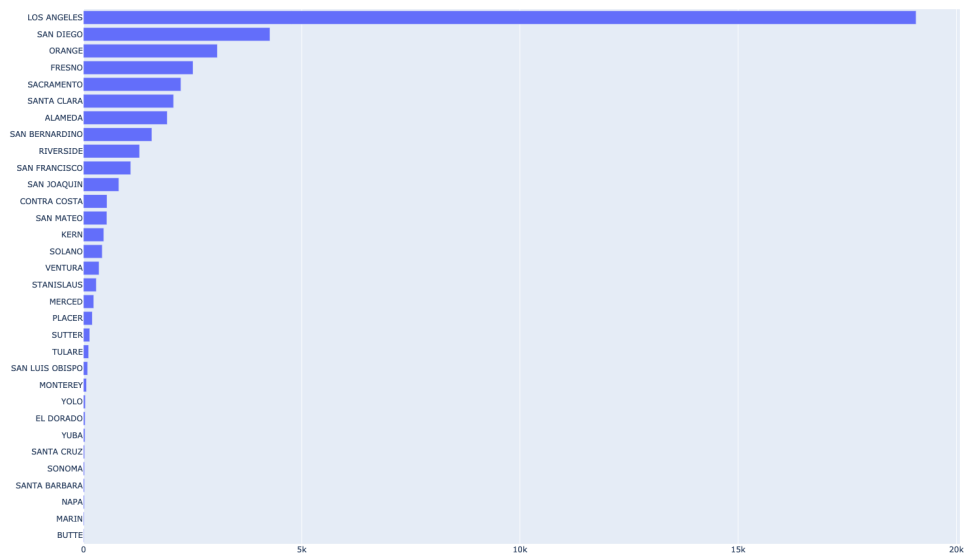
for race in race_df_dic.keys():
    # count the number of cases for each race and create a dataframe with the
    ↪ results we can graph
    race_bycounty_dic[race] = pd.DataFrame({'race': pd.
    ↪ concat([race_df_dic[race].county]).value_counts()})
    # remove any 'NA' county
    race_bycounty_dic[race] = race_bycounty_dic[race][race_bycounty_dic[race].
    ↪ index != "NA"]
    # sort the results in ascending order by the number of cases
    race_bycounty_dic[race] = race_bycounty_dic[race].
    ↪ sort_values(by='race', ascending=True)
    # create the data for the graph
    data = [go.Bar(
        y = race_bycounty_dic[race].index,
        x = race_bycounty_dic[race].race,
        orientation='h'
    )]
    # now create our figure layout
    layout = go.Layout(
        height = 1000,
        margin=go.layout.Margin(l=300),
        title = "%s race reported COVID-19 Cases in California by county" %
    ↪ (race)
    )
    # create a figure object
    fig = go.Figure(data=data, layout=layout)
    py.iplot(fig)

```

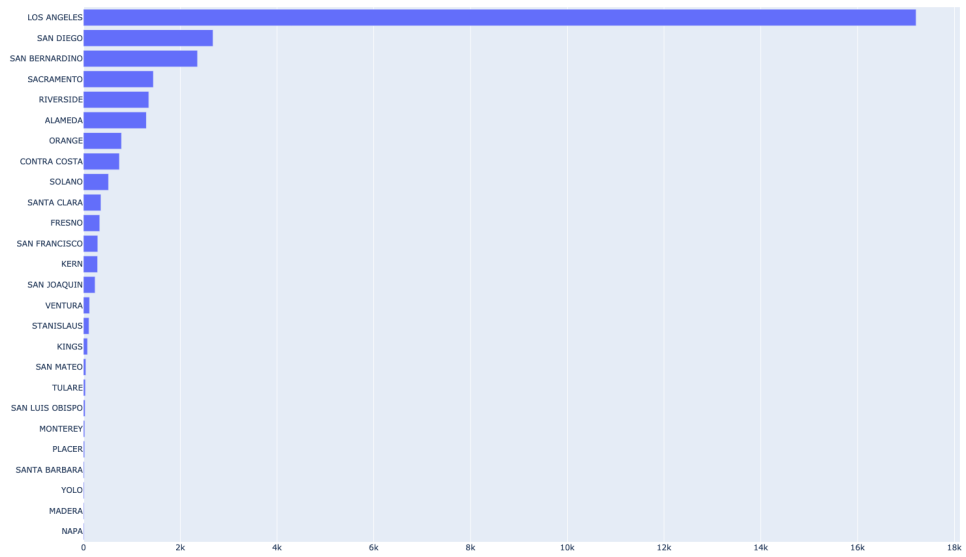
American Indian/Alaska Native race reported COVID-19 Cases in California by county



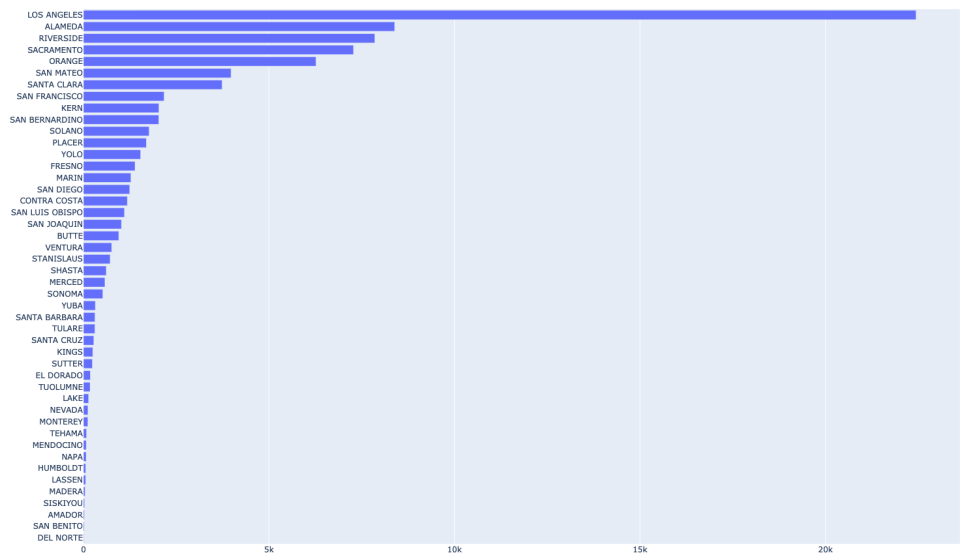
Asian race reported COVID-19 Cases in California by county



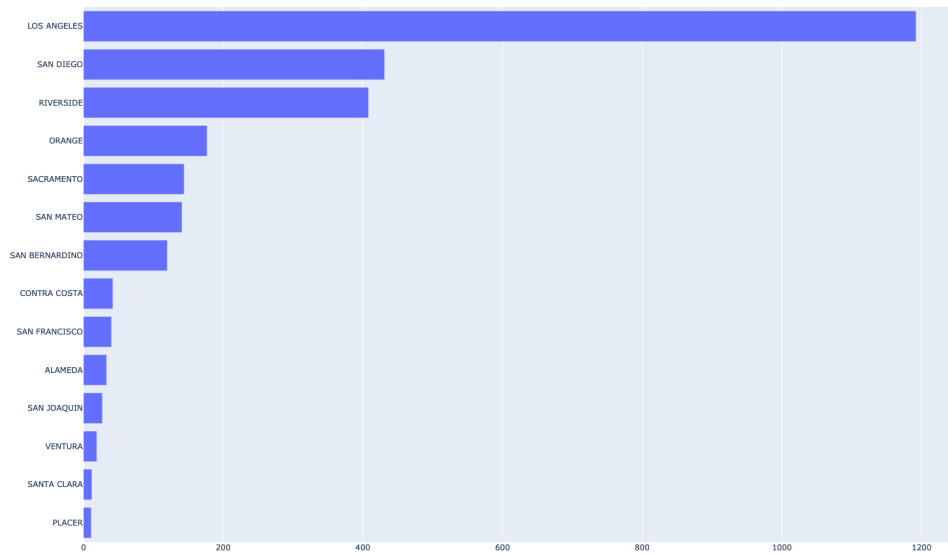
Black race reported COVID-19 Cases in California by county



Multiple/Other race reported COVID-19 Cases in California by county



Native Hawaiian/Other Pacific Islander race reported COVID-19 Cases in California by county



White race reported COVID-19 Cases in California by county

