Project Milestone 5

May 25, 2020

[2]:
```
# Project Milestone 5
# Lenin Kamma 05/25/2020
```

[3]:
```
# Import libraries
import numpy as np
import pandas as pd
import sqlite3
import matplotlib.pyplot as plt
import squarify
```

[4]:
```
# Import finalized saved csv files into a dataframe
```

[5]:
```
county_population = pd.read_csv("C:/Lenin Data Science/DSC540/Project/
 ↪FinalCountyPopulationData.csv")
case_count = pd.read_csv("C:/Lenin Data Science/DSC540/Project/
 ↪FinalCaseCountData.csv")
surface_area = pd.read_csv("C:/Lenin Data Science/DSC540/Project/
 ↪FinalSurfaceArea.csv")
hospital_data = pd.read_csv("C:/Lenin Data Science/DSC540/Project/
 ↪FinalHospitalData.csv")
```

[6]:
```
# View each dataframe
county_population.head(10)
```

[6]:

	Unnamed: 0	state	STATE	COUNTY	estimate_2019	county	Countycode
0	1	Alabama	1	1	55869	Autauga	1001
1	2	Alabama	1	3	223234	Baldwin	1003
2	3	Alabama	1	5	24686	Barbour	1005
3	4	Alabama	1	7	22394	Bibb	1007
4	5	Alabama	1	9	57826	Blount	1009
5	6	Alabama	1	11	10101	Bullock	1011
6	7	Alabama	1	13	19448	Butler	1013
7	8	Alabama	1	15	113605	Calhoun	1015
8	9	Alabama	1	17	33254	Chambers	1017
9	10	Alabama	1	19	26196	Cherokee	1019

[7]:
```
# Update the county column  to County and DROP unwanted columns
county_population['County'] = county_population['county'].rename()
county_population.drop(columns=['STATE','COUNTY', 'county','Unnamed: 0' ],␣
↪inplace=True)
county_population
```

[7]:

	state	estimate_2019	Countycode	County
0	Alabama	55869	1001	Autauga
1	Alabama	223234	1003	Baldwin
2	Alabama	24686	1005	Barbour
3	Alabama	22394	1007	Bibb
4	Alabama	57826	1009	Blount
…	…	…	…	…
3137	Wyoming	42343	56037	Sweetwater
3138	Wyoming	23464	56039	Teton
3139	Wyoming	20226	56041	Uinta
3140	Wyoming	7805	56043	Washakie
3141	Wyoming	6927	56045	Weston

[3142 rows x 4 columns]

[8]:
```
# View case count dataset
case_count.head(10)
```

[8]:

	FIPS_code	Date	US_county	US_state	No_of_cases	No_of_deaths
0	10001	2020-05-21	Kent	Delaware	1281	55
1	10003	2020-05-21	New Castle	Delaware	3053	144
2	10005	2020-05-21	Sussex	Delaware	4006	117
3	1001	2020-05-21	Autauga	Alabama	147	3
4	1003	2020-05-21	Baldwin	Alabama	270	8
5	1005	2020-05-21	Barbour	Alabama	100	1
6	1007	2020-05-21	Bibb	Alabama	52	1
7	1009	2020-05-21	Blount	Alabama	48	1
8	1011	2020-05-21	Bullock	Alabama	71	1
9	1013	2020-05-21	Butler	Alabama	321	11

[130]:
```
surface_area.head(10)
```

[130]:

	Unnamed: 0	STCOU	LND110210D	Statecode
0	2	1001	594.44	AL
1	3	1003	1589.78	AL
2	4	1005	884.88	AL
3	5	1007	622.58	AL
4	6	1009	644.78	AL
5	7	1011	622.81	AL
6	8	1013	776.83	AL
7	9	1015	605.87	AL

```
8          10   1017      596.53        AL
9          11   1019      553.70        AL
```

[131]:
```
hospital_data.head(10)
```

[131]:
```
   COUNTYFIPS  BEDS
0        1001    85
1        1003   398
2        1005    74
3        1007    35
4        1009    40
5        1011    61
6        1013    94
7        1015   590
8        1017   115
9        1019    60
```

[132]:
```
pd.to_numeric(hospital_data["COUNTYFIPS"])
```

[132]:
```
0        1001
1        1003
2        1005
3        1007
4        1009
        ...
2502    72113
2503    72125
2504    72127
2505    72145
2506    72153
Name: COUNTYFIPS, Length: 2507, dtype: int64
```

[133]:
```
# Create a dataframe with State, County, County Code (FIPS code), Surface Area␣
 ↪and Population
# Drop unwanted columns from the join
df_pop_density = pd.merge(county_population,surface_area, how='left', left_on =␣
 ↪'Countycode', right_on = 'STCOU').drop(columns=
                                  ['STCOU','Unnamed: 0'])
```

[134]:
```
df_pop_density
```

[134]:
```
        state  estimate_2019  Countycode      County  LND110210D Statecode
0     Alabama          55869        1001     Autauga      594.44        AL
1     Alabama         223234        1003     Baldwin     1589.78        AL
2     Alabama          24686        1005     Barbour      884.88        AL
3     Alabama          22394        1007        Bibb      622.58        AL
4     Alabama          57826        1009      Blount      644.78        AL
```

```
…       …              …          …          …           …          …
3137  Wyoming          42343      56037  Sweetwater    10426.65        WY
3138  Wyoming          23464      56039       Teton     3995.38        WY
3139  Wyoming          20226      56041       Uinta     2081.26        WY
3140  Wyoming           7805      56043    Washakie     2238.55        WY
3141  Wyoming           6927      56045      Weston     2398.09        WY

[3142 rows x 6 columns]
```

[135]:
```
# Calculate population density (population per square mile))
df_pop_density['Popdensity'] = df_pop_density['estimate_2019'] /
 ↪df_pop_density['LND110210D']
```

[136]:
```
# Remove unwanted columns from pop density dataframe
df_pop_density.drop(columns=['state', 'estimate_2019','County','LND110210D'],␣
 ↪inplace=True)
```

[137]:
```
# View data
df_pop_density.head(10)
```

[137]:
```
   Countycode Statecode  Popdensity
0        1001         AL   93.985936
1        1003         AL  140.418171
2        1005         AL   27.897568
3        1007         AL   35.969675
4        1009         AL   89.683303
5        1011         AL   16.218429
6        1013         AL   25.035078
7        1015         AL  187.507221
8        1017         AL   55.745729
9        1019         AL   47.310818
```

[138]:
```
df_pop_density['Countycode']
```

[138]:
```
0        1001
1        1003
2        1005
3        1007
4        1009
        …
3137    56037
3138    56039
3139    56041
3140    56043
3141    56045
Name: Countycode, Length: 3142, dtype: int64
```

[139]:
```
# Merge df_pop_density, hospital_data on county code
df_work = pd.
 ↪merge(df_pop_density,hospital_data,left_on='Countycode',right_on='COUNTYFIPS')
```

[140]:
```
# Merge df_work with case count dataset on FIPS code(county code)
# drop duplicate columns
df_final_work = pd.
 ↪merge(df_work,case_count,left_on='Countycode',right_on='FIPS_code').
 ↪drop(columns=
                                  ['COUNTYFIPS','Countycode'])
```

[141]:
```
# This is the final Dataset for Visualizations
df_final_work
```

[141]:

	Statecode	Popdensity	BEDS	FIPS_code	Date	US_county	US_state	\
0	AL	93.985936	85	1001	2020-05-21	Autauga	Alabama	
1	AL	140.418171	398	1003	2020-05-21	Baldwin	Alabama	
2	AL	27.897568	74	1005	2020-05-21	Barbour	Alabama	
3	AL	35.969675	35	1007	2020-05-21	Bibb	Alabama	
4	AL	89.683303	40	1009	2020-05-21	Blount	Alabama	
…	…	…	…	…	…	…	…	
2364	WY	12.078099	88	56033	2020-05-21	Sheridan	Wyoming	
2365	WY	4.061036	115	56037	2020-05-21	Sweetwater	Wyoming	
2366	WY	5.872783	48	56039	2020-05-21	Teton	Wyoming	
2367	WY	9.718152	225	56041	2020-05-21	Uinta	Wyoming	
2368	WY	3.486632	18	56043	2020-05-21	Washakie	Wyoming	

	No_of_cases	No_of_deaths
0	147	3
1	270	8
2	100	1
3	52	1
4	48	1
…	…	…
2364	16	0
2365	25	0
2366	100	1
2367	13	0
2368	19	1

[2369 rows x 9 columns]

[142]:
```
df_final_work.describe()
```

[142]:

	Popdensity	BEDS	FIPS_code	No_of_cases	No_of_deaths
count	2368.000000	2369.000000	2369.000000	2369.000000	2369.000000
mean	260.998342	389.847193	30144.165049	567.322077	30.664415

```
std       866.936598   1107.779474   15090.079728   2676.871008    156.209071
min         0.214238      4.000000    1001.000000      1.000000      0.000000
25%        24.013802     25.000000   19003.000000     13.000000      0.000000
50%        58.887062     85.000000   29077.000000     51.000000      1.000000
75%       164.338456    270.000000   42121.000000    221.000000      9.000000
max     18808.384894   25613.000000   56043.000000   67551.000000   3114.000000
```

[143]:
```
# Maximum number of deaths in a county is 3114
# Maximum population density is 18808 persons per square mile
# Mean no of cases is 567
```

[144]:
```
# Create a new database covid
conn = sqlite3.connect('covid.db')
c = conn.cursor()
```

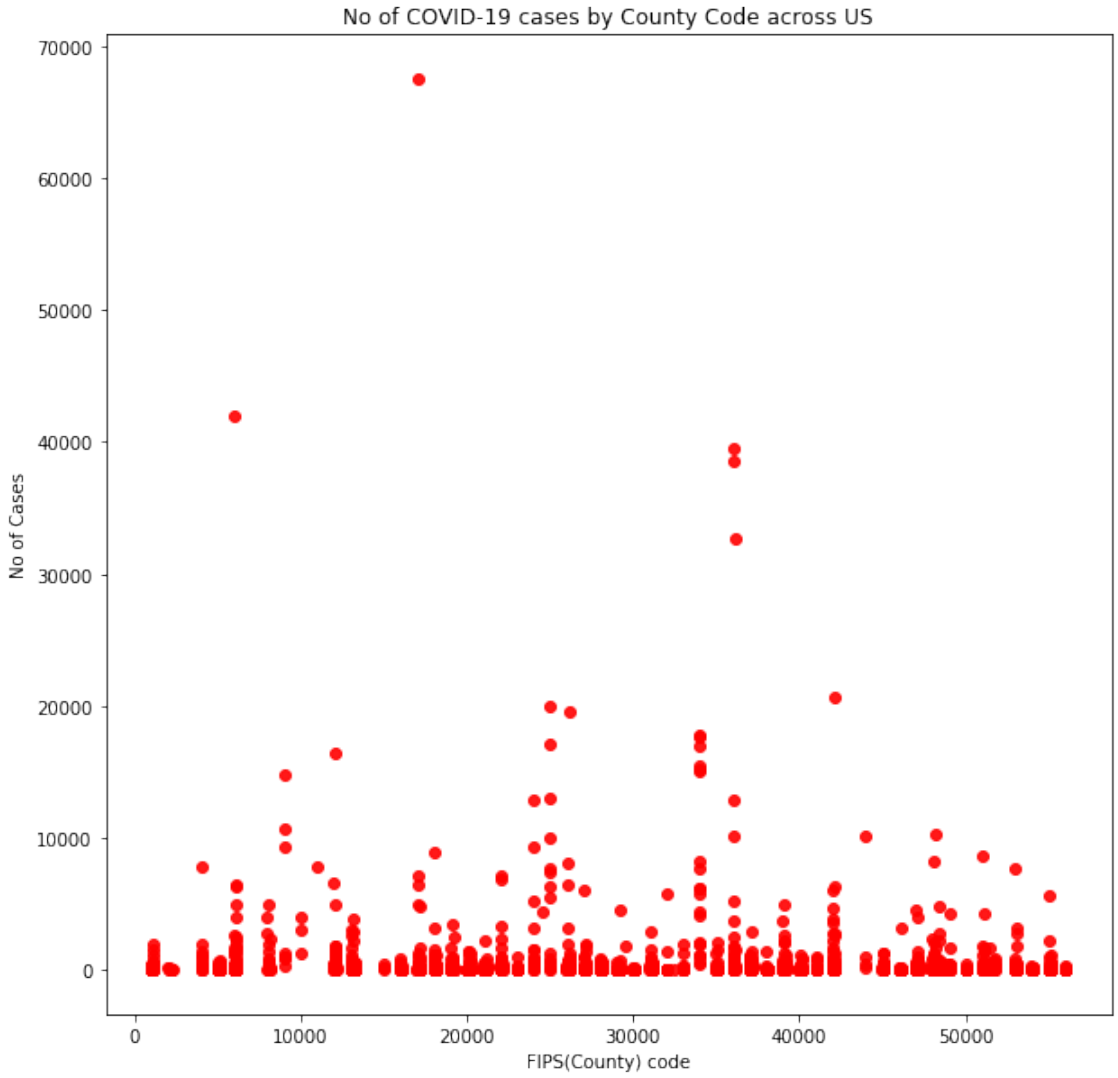
[145]:
```
# Create sql using the dataframe
df_final_work.to_sql("covid_data", conn, if_exists="replace")
```

[146]:
```
# Create a new table in the database
# First drop the table if already exists
conn.execute(
    """
    drop table covid_table
    """)
conn.execute(
    """
    create table covid_table as
    select * from covid_data
    """)
conn.commit()
```

[147]:
```
# Retrive data from database
with sqlite3.connect("covid.db") as conn:
    cursor = conn.cursor()
    rows = cursor.execute("Select * from covid_table")
```

[172]:
```
# Draw scatter plot with number of cases on y-axis and county code on x-axi
rng = np.random.RandomState(0)
colors = rng.rand(2369)
sizes = 10000 * rng.rand(2369)
plt.figure(figsize=(10,10))
plt.plot(df_final_work["FIPS_code"], df_final_work["No_of_cases"],'o',color=␣
 ↪'red',alpha=0.9)
plt.xlabel('FIPS(County) code')
plt.ylabel('No of Cases');
plt.title("No of COVID-19 cases by County Code across US")
plt.savefig('C:/Lenin Data Science/DSC540/scatter1.pdf', dpi=1200)
```

```
plt.show()
```

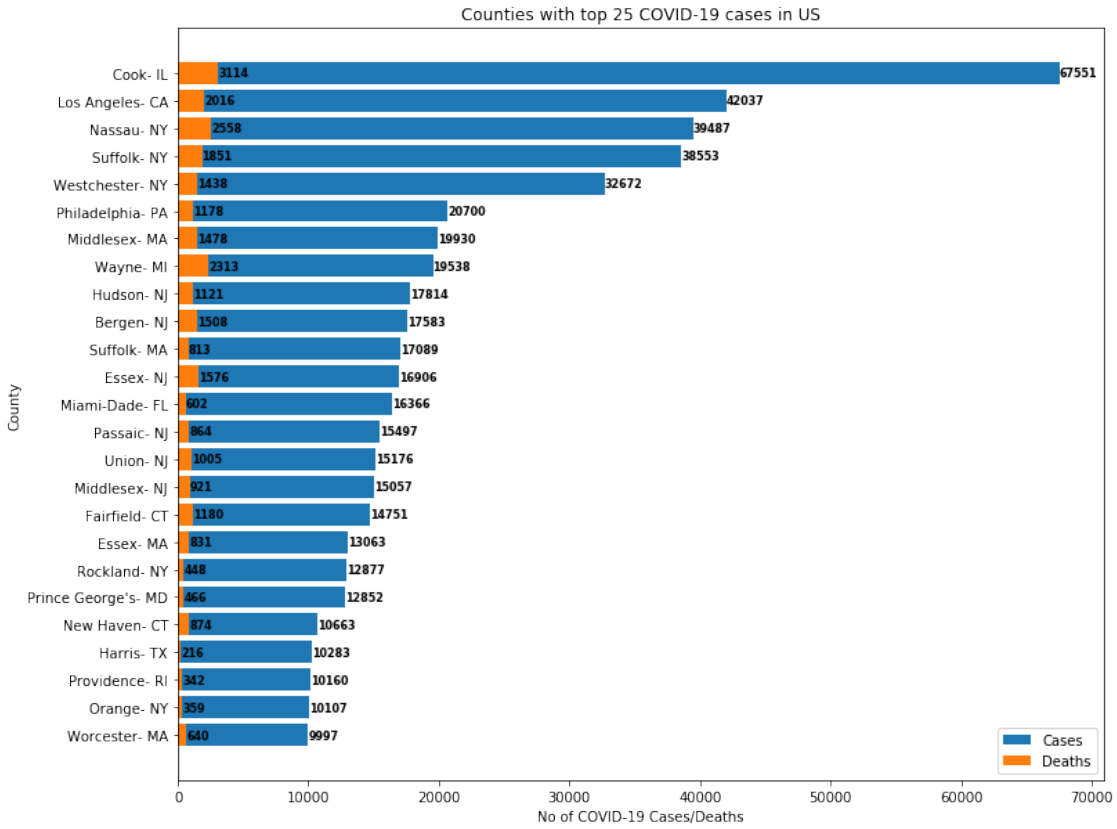


[149]:
```
# There is one outlier with number of cases close to 70000
# Most of the counties have covid cases less than 5000
```

[150]:
```
# Let's take top 25 counties from DB
list1 = []
columns = ['County', 'Count', 'Deaths']
with sqlite3.connect("covid.db") as conn:
    cursor = conn.cursor()
    rows = cursor.execute("SELECT US_county||'-'||Statecode,␣
 ↪No_of_cases,No_of_deaths from covid_table ORDER BY No_of_cases desc LIMIT␣
 ↪25")
    for row in rows:
```

```
        list1.append(row)
top25_df = pd.DataFrame(list1, columns=columns)
top25_df = top25_df.sort_values(by='Count')
```

[169]:
```
# Draw bar plot with number of cases/Number of deaths on y-axis and county code␣
 ↪on x-axis
fig, ax = plt.subplots(figsize=(12,10))
ax.barh(top25_df['County'], top25_df['Count'], align='center')
ax.barh(top25_df['County'], top25_df['Deaths'], align='center')
legend_val = ['Cases', 'Deaths']
plt.xlabel('No of COVID-19 Cases/Deaths')
plt.ylabel('County')
plt.title("Counties with top 25 COVID-19 cases in US")
for i, v in enumerate(top25_df['Count']):
    ax.text(v + 10, i, str(v), color='black', fontweight='bold', fontsize=8,␣
 ↪ha='left', va='center')
for i, v in enumerate(top25_df['Deaths']):
    ax.text(v + 10, i, str(v), color='black', fontweight='bold', fontsize=8,␣
 ↪ha='left', va='center')
ax.legend(legend_val,loc='best')
plt.savefig('C:/Lenin Data Science/DSC540/bar1.pdf', dpi=1200)
plt.show()
```



[152]:
```
# No. of cases are highest in Cook (IL) county
# There are 24 counties with more than 10000 cases (as of 05/23/2020)
```

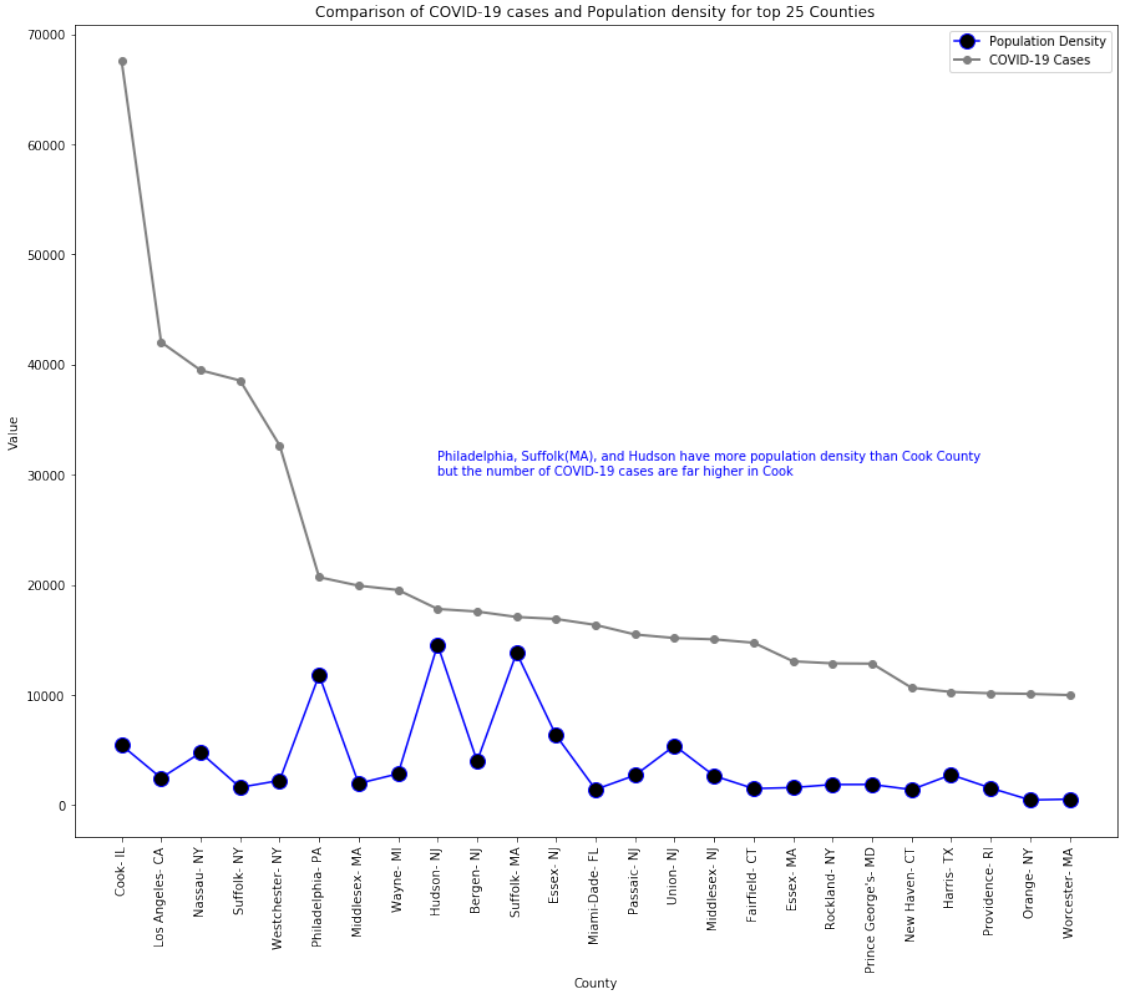
[153]:
```
# Let's take top 25 counties from DB
list1 = []
columns = ['County', 'Popdensity', 'Casecount']
with sqlite3.connect("covid.db") as conn:
    cursor = conn.cursor()
    rows = cursor.execute("SELECT US_county||'-'||Statecode, Popdensity,␣
 ↪No_of_cases from covid_table ORDER BY No_of_cases desc LIMIT 25")
    for row in rows:
        list1.append(row)
top25_den_df = pd.DataFrame(list1, columns=columns)
top25_den_df
```

[153]:

	County	Popdensity	Casecount
0	Cook- IL	5448.079507	67551
1	Los Angeles- CA	2473.978284	42037
2	Nassau- NY	4765.819050	39487
3	Suffolk- NY	1618.991283	38553
4	Westchester- NY	2247.400697	32672
5	Philadelphia- PA	11812.557793	20700
6	Middlesex- MA	1970.725832	19930
7	Wayne- MI	2858.029996	19538
8	Hudson- NJ	14557.068630	17814
9	Bergen- NJ	4000.695249	17583
10	Suffolk- MA	13824.711952	17089
11	Essex- NJ	6330.520561	16906
12	Miami-Dade- FL	1431.686445	16366
13	Passaic- NJ	2718.597974	15497
14	Union- NJ	5408.720591	15176
15	Middlesex- NJ	2670.881487	15057
16	Fairfield- CT	1509.596889	14751
17	Essex- MA	1601.904337	13063
18	Rockland- NY	1877.205416	12877
19	Prince George's- MD	1883.873708	12852
20	New Haven- CT	1413.966684	10663
21	Harris- TX	2766.880151	10283
22	Providence- RI	1560.271062	10160
23	Orange- NY	474.245094	10107
24	Worcester- MA	549.800433	9997

[168]:
```
# Compare number of cases with population density
fig, ax = plt.subplots(figsize=(15,12))
```

```
plt.plot(top25_den_df['County'], top25_den_df['Popdensity'], marker='o',␣
 ↪markerfacecolor='black', markersize=12, color='blue', label="Population␣
 ↪Density")
plt.plot(top25_death_df['County'], top25_death_df['Casecount'], marker='o',␣
 ↪color='gray', linewidth=2, label="COVID-19 Cases")
plt.xlabel('County')
plt.xticks(rotation=90)
plt.ylabel('Value');
plt.title("Comparison of COVID-19 cases and Population density for top 25␣
 ↪Counties")
plt.text(8,30000,'Philadelphia, Suffolk(MA), and Hudson have more population␣
 ↪density than Cook County\nbut the number of COVID-19 cases are far higher in␣
 ↪Cook'
        ,color='blue')
plt.legend()
plt.savefig('C:/Lenin Data Science/DSC540/line1.pdf', dpi=1200)
plt.show()
```



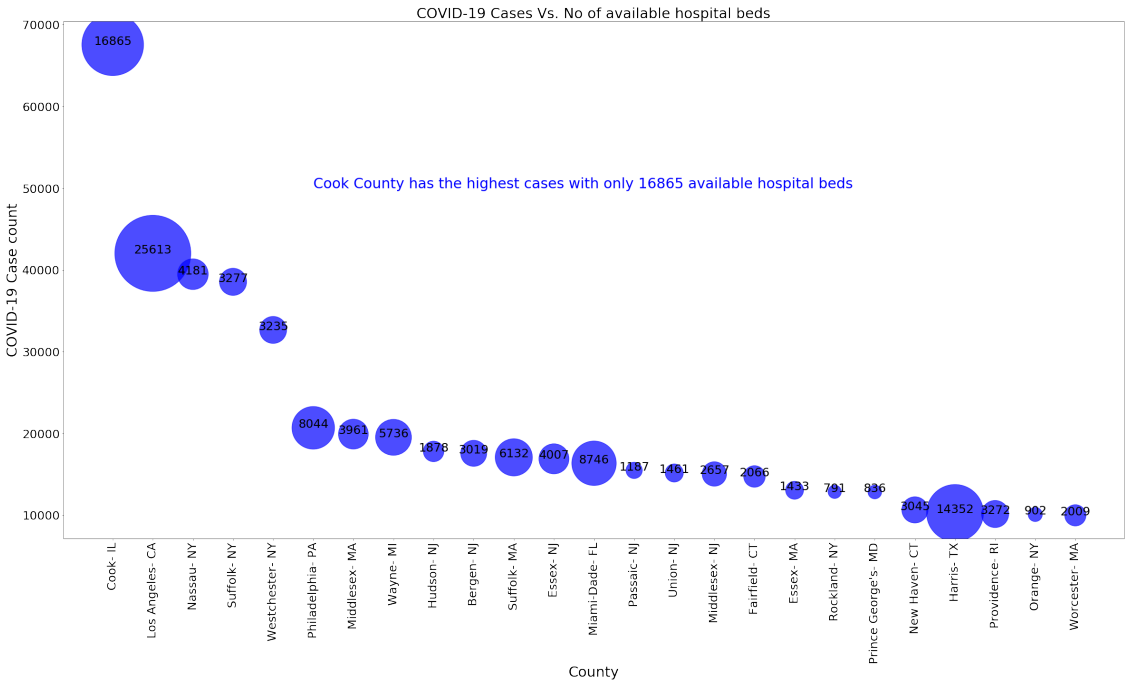
[155]:
```
# Top 25 counties with highest cases
list3 = []
columns = ['County','Casecount', 'Beds']
with sqlite3.connect("covid.db") as conn:
    cursor = conn.cursor()
    rows = cursor.execute("SELECT US_county||'-'||Statecode, No_of_cases, BEDS␣
 ↪from covid_table ORDER BY No_of_cases desc LIMIT 25")
    for row in rows:
        list3.append(row)
top25_beds_df = pd.DataFrame(list3, columns=columns)
top25_beds_df = top25_beds_df.dropna()
top25_beds_df
```

[155]:

	County	Casecount	Beds
0	Cook- IL	67551	16865
1	Los Angeles- CA	42037	25613
2	Nassau- NY	39487	4181
3	Suffolk- NY	38553	3277
4	Westchester- NY	32672	3235
5	Philadelphia- PA	20700	8044
6	Middlesex- MA	19930	3961
7	Wayne- MI	19538	5736
8	Hudson- NJ	17814	1878
9	Bergen- NJ	17583	3019
10	Suffolk- MA	17089	6132
11	Essex- NJ	16906	4007
12	Miami-Dade- FL	16366	8746
13	Passaic- NJ	15497	1187
14	Union- NJ	15176	1461
15	Middlesex- NJ	15057	2657
16	Fairfield- CT	14751	2066
17	Essex- MA	13063	1433
18	Rockland- NY	12877	791
19	Prince George's- MD	12852	836
20	New Haven- CT	10663	3045
21	Harris- TX	10283	14352
22	Providence- RI	10160	3272
23	Orange- NY	10107	902
24	Worcester- MA	9997	2009

[171]:
```
# Create a bubble plot with number of Cases and Beds
fig = plt.figure(figsize=(40,20))
area1 = (top25_beds_df['Beds'])

# Choose random colors
colors1=np.random.rand(15)
```

```
# Scatter plot and labels
plt.scatter(top25_beds_df['County'],␣
 ↪top25_beds_df['Casecount'],s=area1,color='blue', alpha=0.7)
plt.title('COVID-19 Cases Vs. No of available hospital beds', fontsize=30 )
plt.xlabel('County', fontsize=30)
plt.ylabel('COVID-19 Case count', fontsize=30)

x , y = top25_beds_df['County'], top25_beds_df['Casecount']
for i, txt in enumerate(top25_beds_df['Beds']):
    plt.annotate(txt, (x[i], y[i]), fontsize=25, horizontalalignment='center')


# Rotate xticks to show vertical
plt.xticks(rotation=90, fontsize=25)
plt.yticks(fontsize=25)

plt.text(5,50000,'Cook County has the highest cases with only 16865 available␣
 ↪hospital beds', color='blue', fontsize=30)
plt.savefig('C:/Lenin Data Science/DSC540/bubble1.pdf', dpi=2400)
plt.show()
```



[160]:
```
# Top 25 counties with highest cases
list4 = []
columns = ['County','Casecount', 'Popdensity']
```

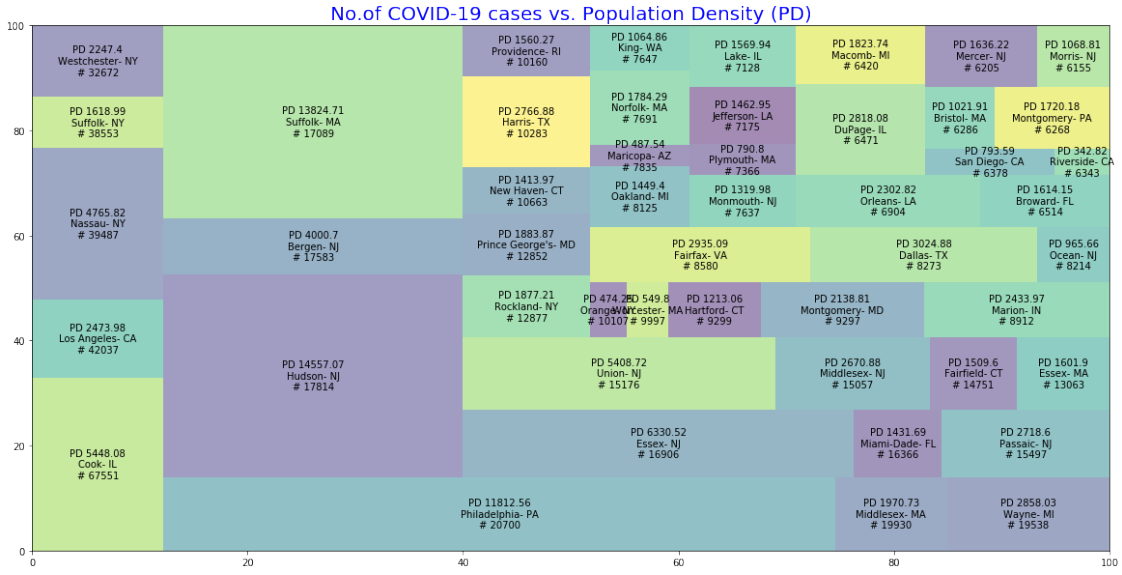
```
with sqlite3.connect("covid.db") as conn:
    cursor = conn.cursor()
    rows = cursor.execute("SELECT US_county||'-'||Statecode, No_of_cases,␣
 ↪popdensity from covid_table ORDER BY No_of_cases DESC LIMIT 50")
    for row in rows:
        list4.append(row)
top50_pop_df = pd.DataFrame(list4, columns=columns)
top50_pop_df = top50_pop_df.dropna()
top50_pop_df
```

[160]:

	County	Casecount	Popdensity
0	Cook- IL	67551	5448.079507
1	Los Angeles- CA	42037	2473.978284
2	Nassau- NY	39487	4765.819050
3	Suffolk- NY	38553	1618.991283
4	Westchester- NY	32672	2247.400697
5	Philadelphia- PA	20700	11812.557793
6	Middlesex- MA	19930	1970.725832
7	Wayne- MI	19538	2858.029996
8	Hudson- NJ	17814	14557.068630
9	Bergen- NJ	17583	4000.695249
10	Suffolk- MA	17089	13824.711952
11	Essex- NJ	16906	6330.520561
12	Miami-Dade- FL	16366	1431.686445
13	Passaic- NJ	15497	2718.597974
14	Union- NJ	15176	5408.720591
15	Middlesex- NJ	15057	2670.881487
16	Fairfield- CT	14751	1509.596889
17	Essex- MA	13063	1601.904337
18	Rockland- NY	12877	1877.205416
19	Prince George's- MD	12852	1883.873708
20	New Haven- CT	10663	1413.966684
21	Harris- TX	10283	2766.880151
22	Providence- RI	10160	1560.271062
23	Orange- NY	10107	474.245094
24	Worcester- MA	9997	549.800433
25	Hartford- CT	9299	1213.059448
26	Montgomery- MD	9297	2138.805089
27	Marion- IN	8912	2433.969215
28	Fairfax- VA	8580	2935.089649
29	Dallas- TX	8273	3024.878340
30	Ocean- NJ	8214	965.657305
31	Oakland- MI	8125	1449.397229
32	Maricopa- AZ	7835	487.537581
34	Norfolk- MA	7691	1784.289718
35	King- WA	7647	1064.858171
36	Monmouth- NJ	7637	1319.983361

37	Plymouth- MA	7366	790.802331
38	Jefferson- LA	7175	1462.953692
39	Lake- IL	7128	1569.939369
40	Orleans- LA	6904	2302.821391
41	Broward- FL	6514	1614.146257
42	DuPage- IL	6471	2818.079389
43	Macomb- MI	6420	1823.738575
44	San Diego- CA	6378	793.587741
45	Riverside- CA	6343	342.822848
46	Bristol- MA	6286	1021.907431
47	Montgomery- PA	6268	1720.178453
48	Mercer- NJ	6205	1636.221945
49	Morris- NJ	6155	1068.810031

[167]:
```
# Create a label to show on tree map
top50_pop_df['label1'] = "PD " + round(top50_pop_df["Popdensity"],2).
 ↪astype(str) +'\n'+ top50_pop_df["County"].astype(str) +'\n' + "# " +␣
 ↪top50_pop_df["Casecount"].astype(str)

# Create tree map
plt.figure(figsize=(20,10))
squarify.plot(sizes=top50_pop_df['Popdensity'],label=top50_pop_df["label1"],␣
 ↪alpha=0.5 )
plt.title("No.of COVID-19 cases vs. Population Density (PD)", fontsize=20,␣
 ↪color="blue")
plt.savefig('C:/Lenin Data Science/DSC540/tree1.pdf', dpi=1200)
plt.show()
```



[162]:
```
# Save final data file to the local drive
#Save final dataset to a csv file
df_final_work.to_csv(r'C:/Lenin Data Science/DSC540/Project/FinalDataFile.csv')
```