

final_report_notebook

June 10, 2020

1 Final Report Notebook

This is our main program for importing data, manipulating it, and exporting it to .CSV files used for creating a network graph of COVID infection. We chose to import data from a John's Hopkins dataset that we can derive infection rate of different U.S. counties from. In this project we are specifically looking at the state of New York

```
[125]: from IPython.display import Image

Image(filename="new-york-county-map.gif")
```

```
[125]: <IPython.core.display.Image object>
```

```
[123]: import pandas as pd
import matplotlib.pyplot as plt
import networkx as nx
import datetime as dt
import numpy as np
import math
from sklearn import linear_model
import joblib
import warnings
from IPython.display import display
from IPython.display import Video
from IPython.core.interactiveshell import InteractiveShell
```

The method below is used to filter data from a Pandas dataframe based on the state parameter given. This is useful when looking to pull specific state data.

```
[102]: # useful for filtering data from a certain state or province
def filter_data_by_state(data, state):
    data = data[data['Province_State'] == state]
    return data[data['Admin2'] != 'Unassigned']
```

The create_nodes_for_data method takes in a Pandas dataframe, and populates a Networkx graph with nodes created with county names.

[103]: *# makes nodes in graph for each row in data*

```
def create_nodes_for_data(data):
    G = nx.Graph()
    for index, row in data.iterrows():
        G.add_node(str(row.iloc[5]))
    return G
```

The `populate_data_into_nodes_state` method is used to pull data from a Pandas dataframe and populate each node in the initialized Networkx graph 'G' with important data like latitude, longitude, time of first infection, and so on. The method will also assign each node a relative time, based on first infection.

[104]: *# fills nodes with data and assign relative_time*

```
def populate_data_into_nodes_state(G, data, logistic_curve=False,
    ↪pop_data=None, load_curves=False,
                                max_iterations=10000):
    already_infected = set()
    logistic_curves = dict()
    time = 0
    start_date = dt.datetime(2020, 1, 22)
    current_date = dt.datetime(2020, 1, 22)
    end_date = dt.datetime(2020, 5, 27) # change as more data comes in
    while current_date <= end_date:
        day_data = data[data[current_date.strftime('%m/%d/%y')] > 0]
        was_added = False
        for index, row in day_data.iterrows():
            county = str(row.iloc[5])
            if county not in already_infected:
                G.nodes[county]['relative_time'] = time
                G.nodes[county]['real_time'] = current_date.strftime('%m/%d/
    ↪%y')

                G.nodes[county]['lat'] = row.iloc[8]
                G.nodes[county]['long'] = row.iloc[9]
                G.nodes[county]['data'] = row.iloc[12:]
                if pop_data is not None:
                    pop = pop_data[pop_data['County'] == county]
                    if not pop.empty:
                        G.nodes[county]['pop'] = pop['Population']
                    else:
                        G.nodes[county]['pop'] = 2000 # give unknown counties
    ↪2000 population
                if logistic_curve:
                    if load_curves:
                        load_logistic_curve(county, logistic_curves)
                    else:
                        logistic_curves[county] = fit_logistic_curve(G.
    ↪nodes[county], max_iterations)
```

```

        already_infected.add(county)
        was_added = True
    if was_added:
        time += 1
        current_date += dt.timedelta(days=1)
    return time, logistic_curves

```

The `create_edges_for_graph_first_infection` method is used to create edges between the nodes in a NetworkX graph, based on the relative time of infection, and the distance between the two counties.

```

[105]: # uses relative_time field to determine where to place edges, distance can be
        ↳ limited but by default is not
def create_edges_for_graph_first_infection(G, max_relative_time,
        ↳ distance_limit):
    time = 0
    prev_nodes = []
    temp = []
    while time <= max_relative_time:
        for node in G.nodes(data=True):
            if 'relative_time' in node[1] and node[1]['relative_time'] == time:
                for prev_node in prev_nodes:
                    distance = calculate_distance(prev_node, node)
                    if distance <= distance_limit:
                        G.add_edge(prev_node[0], node[0], weight=distance)
                temp.append(node)
        prev_nodes = temp.copy()
        temp.clear()
        time += 1

```

Much like the above method, the `create_edges_for_graph_threshold_distance` method checks if the number of infected in the county, with proportion to the distance to the next county, meets a certain specified threshold. If it does meet this threshold, the method will create an edge between the two counties.

```

[106]: # uses real_time and distance to determine where to place edges threshold and
        ↳ distance_limit must be provided
def create_edges_for_graph_threshold_distance(G, threshold, distance_limit):
    start_date = dt.datetime(2020, 1, 22)
    current_date = dt.datetime(2020, 1, 22)
    end_date = dt.datetime(2020, 5, 27)
    prev_nodes = []
    while current_date <= end_date:
        for node in G.nodes(data=True):
            if 'real_time' in node[1] and node[1]['real_time'] == current_date.
            ↳ strftime('%m/%d/%y'):
                for prev_node in prev_nodes:
                    distance = calculate_distance(prev_node, node)

```

```

        day = int(current_date.strftime('%#j')) - int(start_date.
→strftime('%#j'))
        weight = prev_node[1]['data'][int(day)]/distance
        if weight >= threshold and distance <= distance_limit:
            G.add_edge(prev_node[0], node[0], weight=weight)
        prev_nodes.append(node)
        current_date += dt.timedelta(days=1)

```

This method uses a logistic regression model from SciKit-Learn to predict a fitted logistic curve for each county, and then runs a simulation to create edges in the network.

This simulation is based off a radius of infection, calculated by the logistic regression model. If a county is within the infection radius distance of an already infected county, that county will also begin to become infected.

```

[107]: # uses regression to fit a logistic growth curve to the each county and then
→run a simulation to create edges
def create_edges_for_graph_logistic_simulation(G, logistic_curves, time_limit,
→radius_weight):
    time = np.zeros((1, 1), dtype=int)
    infected = []
    found_first = False
    while time[0][0] <= time_limit:
        if not found_first:
            for node in G.nodes(data=True):
                if node[0] in logistic_curves and logistic_curves[node[0]].
→predict(time)[0] > 1:
                    infected.append(node)
                    found_first = True
                    break
            time[0][0] += 1
            if found_first:
                break
        while time[0][0] <= time_limit:
            for node1 in infected:
                radius = radius_weight * logistic_curves[node1[0]].predict(time)[0]
→# / float(node1[1]['pop'])
                for node2 in G.nodes(data=True):
                    if node2 in infected:
                        continue
                    if 'lat' in node2[1]:
                        distance = calculate_distance(node1, node2)
                        if distance < radius:
                            G.add_edge(node1[0], node2[0], weight=distance)
                            infected.append(node2)

            time[0][0] += 1

```

The below method is key for creating our infection simulation. This will take in a node from the Networkx graph, and use the Scikit-Learn python library to create a logistic regression model to fit logistic curves to the infection data in that node. It will then return the logistic curve.

The logistic regression model will the node to fit a curve based on the data within that node. It will optimize the data to the logistic function:

$$f(x) = \frac{L}{1 + e^{-k(x-x_0)}}$$

where x_0 is the x value of the sigmoid midpoint, L is the curves maximum value, and k is the growth rate of the curve.

```
[108]: # uses sklearn to fit a logistic growth curve to the data in the node
def fit_logistic_curve(node, max_iterations):
    x = np.linspace(0, len(node['data'])-1, len(node['data']), dtype=int)
    y = np.zeros(len(x), dtype=int)
    for i in range(0, len(y)):
        y[i] = int(node['data'][i])
    x = x[:, np.newaxis]
    warnings.filterwarnings("ignore")
    clf = linear_model.LogisticRegression(C=1e5, max_iter=max_iterations)
    clf.fit(x, y)
    # # used to plot logistic curves
    # plt.figure(1, figsize=(4, 3))
    # # plt.clf()
    # x = np.linspace(0, 250, 251)
    # x = x[:, np.newaxis]
    # plt.plot(x, clf.predict(x))
    # plt.show()
    return clf
```

This method is used to save the produced logistic curves to a certain directory. The default directory is logistic_curves, but this can be modified to save to different locations.

```
[109]: # saves logistic_curves to a directory named logistic_curves and a modifier can
    ↪ be added for different save locations
def save_logistic_curves(logistic_curves, modifier=''):
    if modifier is not '':
        modifier = '_' + modifier
    for key in logistic_curves:
        joblib.dump(logistic_curves[key], 'logistic_curves%s/%s.pkl' %
    ↪ (modifier, key))
```

This method will load logistic curves from a certain directory. The default directory is logistic_curves, but this can be modified to load from different locations.

```
[110]: # loads logistic curves from a directory named logistic_curves and a modifier
    ↪ can be added for different load locations
def load_logistic_curve(county, logistic_curves, modifier=''):
```

```

    if modifier is not '':
        modifier = '_' + modifier
    logistic_curves[country] = joblib.load('logistic_curves%s/%s.pkl' % (
    ↪modifier, country))

```

This method will run previously defined methods to create a Networkx graph of the spread of Covid-19 based on first infection in a location and connects location which were infected next.

```

[111]: def create_graph_first_infected(data, distance_limit=30000):
        G = create_nodes_for_data(data)
        max_time = populate_data_into_nodes_state(G, data)[0]
        create_edges_for_graph_first_infection(G, max_time, distance_limit)
        return G

```

This method will run previously defined methods in order to create a networkx graph of the spread Covid-19 based on a threshold and distance_limit. This method may be better on larger sample sizes so far locations are not infecting each other

```

[112]: def create_graph_infected_distance(data, threshold=2.5, distance_limit=75):
        G = create_nodes_for_data(data)
        populate_data_into_nodes_state(G, data)
        create_edges_for_graph_threshold_distance(G, threshold=threshold,
    ↪distance_limit=distance_limit)
        return G

```

This method will run previously defined methods in order to create a networkx graph of the spread Covid-19 based on a Logistic simulation

```

[113]: def create_graph_logistic_simulation(data, time_limit=250, radius_weight=.5):
        G = create_nodes_for_data(data)
        pop_data = pd.read_csv('pop_ny_data_2020.csv')
        max_relative_time, logistic_curves = populate_data_into_nodes_state(G,
    ↪data, logistic_curve=True, pop_data=pop_data,
        ↪
    ↪max_iterations=50000)
        create_edges_for_graph_logistic_simulation(G, logistic_curves, time_limit,
    ↪radius_weight)
        save_logistic_curves(logistic_curves)
        return G

```

This method will run previously defined methods in order to create a networkx graph of the spread Covid-19 based on a logistic simulation loaded from saved logistic curves.

```

[114]: def create_graph_logistic_simulation_load_lc(data, time_limit=250,
    ↪radius_weight=.5):
        G = create_nodes_for_data(data)
        pop_data = pd.read_csv('pop_ny_data_2020.csv')

```

```

    max_relative_time, logistic_curves = populate_data_into_nodes_state(G,
↪data, logistic_curve=True, pop_data=pop_data,

↪load_curves=True)
    create_edges_for_graph_logistic_simulation(G, logistic_curves, time_limit,
↪radius_weight)
    return G

```

This method converts the networkx graph created to two csv files which can be imported into cytoscape.

```

[115]: def convert_to_csv(G):
    output_node = ''
    ↪'county,relative_time,real_time,lat,long,degree,degree_centrality,closeness_centrality\n'
    output_edge = 'source,target,distance\n'
    for node in G.nodes(data=True):
        if 'relative_time' in node[1]:
            output_node += '%s,%s,%s,%s,%s,%s,%s,%s\n' % (node[0],
↪node[1]['relative_time'], node[1]['real_time'],
                                                                    node[1]['lat'],
↪node[1]['long'], node[1]['degree'],
                                                                    node[1]['degree_centrality'],
↪node[1]['closeness_centrality'])
        for edge in G.edges(data=True):
            output_edge += '%s,%s,%s\n' % (edge[0], edge[1], edge[2]['weight'])

    return output_node, output_edge

```

The convert_to_cyto_layout method is used to convert the cytoscape data and network csv files into csv files usable for the cytoscape coordinateLayout plugin.

```

[116]: def convert_to_cyto_layout(node_file_in, edge_file_in, node_file_out,
↪edge_file_out, scale=25):
    nodes_out_text = ''
    edges_out_text = ''
    nodes_in = pd.read_csv(node_file_in)
    cities = dict()
    i = 1
    for index, row in nodes_in.iterrows():
        cities[row.iloc[0]] = i
        nodes_out_text += '%s %s %s %s %s\n' % (i, str(row.iloc[0]).replace('↪
↪', '_'), '1', int(row.iloc[3])/scale,
                                                                    int(row.iloc[4])/scale)

        i += 1
    edges_in = pd.read_csv(edge_file_in)
    for index, row in edges_in.iterrows():

```

```

edges_out_text += '%s %s %s\n' % (cities[row.iloc[0]], cities[row.
↪iloc[1]], row.iloc[2])

with open(node_file_out, 'w') as f:
    f.write(nodes_out_text)
    f.close()

with open(edge_file_out, 'w') as f:
    f.write(edges_out_text)
    f.close()

```

These two methods are used to calculate the distance between two nodes. This calculation uses the latitude and longitude of the counties that the nodes represent.

```

[117]: # calculates the distance between two nodes
def calculate_distance(node1, node2):
    lat1 = node1[1]['lat']
    lat2 = node2[1]['lat']
    lon1 = node1[1]['long']
    lon2 = node2[1]['long']
    R = 3958.5 # radius of earth in miles
    dlat = deg_to_rad(lat2 - lat1)
    dlon = deg_to_rad(lon2 - lon1)
    a = (math.sin(dlat/2))**2 + math.cos(deg_to_rad(lat1)) * math.
↪cos(deg_to_rad(lat2)) * (math.sin(dlon/2))**2
    c = 2 * math.atan2(math.sqrt(a), math.sqrt(1-a))
    return math.floor(R * c)

# helper function for calculate distance
def deg_to_rad(deg):
    return deg * (math.pi / 180)

```

This method is used to calculate useful statistics of nodes in the graph such as, centrality and degree.

```

[118]: def calculate_node_stats(G):
    for item in nx.degree_centrality(G).items():
        G.nodes[item[0]]['degree_centrality'] = item[1]
    for item in nx.closeness_centrality(G).items():
        G.nodes[item[0]]['closeness_centrality'] = item[1]
    for item in G.degree():
        G.nodes[item[0]]['degree'] = item[1]

```

This method is used to write data from the Networkx graph to a csv format, and will then write it to a local directory file.


```
[119]: def write_csv(filename, G):
        csv = convert_to_csv(G)
        with open('%s_network.csv' % filename, 'w') as f:
            f.write(csv[1])
            f.close()

        with open('%s_data.csv' % filename, 'w') as f:
            f.write(csv[0])
            f.close()
```

The main method of our program. This will read in the John's Hopkins csv data into a Pandas dataframe, will filter that dataframe to the state of New York, and will use the methods above to create various visualizations.

```
[120]: def main():
        data = pd.read_csv('time_series_covid19_confirmed_US.csv')

        new_york = filter_data_by_state(data, 'New York')
        display(new_york)

        # visualizes data in graph vs time
        dates = list(range(0, len(data.columns[12:].values)))
        plt.figure()
        for index, row in new_york.iterrows():
            plt.plot(dates, row.iloc[12:], label=row.iloc[5])
        plt.title(str.title('NY Counties cases vs. time'))
        plt.legend(framealpha=2, frameon=True, ncol=3, loc='upper left')
        plt.show()

        # creates graph of first infection
        G = create_graph_first_infected(new_york)
        calculate_node_stats(G)
        write_csv('first_infection', G)

        # creates graph of threshold distance
        H = create_graph_infected_distance(new_york)
        calculate_node_stats(H)
        write_csv('threshold_distance', H)

        # creates graph of logistic simulation
        J = create_graph_logistic_simulation(new_york)
        # J = create_graph_logistic_simulation_load_lc(new_york, radius_weight=1)
        calculate_node_stats(J)
        write_csv('logistic_simulation', J)
```

```

    # this will make files for the coordinateLayout plugin for cytoscape for
    ↪ each graph
    convert_to_cyto_layout('first_infection_data.csv', 'first_infection_network.
    ↪ csv',
                           'first_infection_data_cyto_layout.csv',
    ↪ 'first_infection_network_cyto_layout.csv')
    convert_to_cyto_layout('threshold_distance_data.csv',
    ↪ 'threshold_distance_network.csv',
                           'threshold_distance_data_cyto_layout.csv',
    ↪ 'threshold_distance_network_cyto_layout.csv')
    convert_to_cyto_layout('logistic_simulation_data.csv',
    ↪ 'logistic_simulation_network.csv',
                           'logistic_simulation_data_cyto_layout.csv',
    ↪ 'logistic_simulation_network_cyto_layout.csv')

if __name__ == '__main__':
    main()

```

	UID	iso2	iso3	code3	FIPS	Admin2	Province_State \
1833	84036001	US	USA	840	36001.0	Albany	New York
1834	84036003	US	USA	840	36003.0	Allegany	New York
1835	84036005	US	USA	840	36005.0	Bronx	New York
1836	84036007	US	USA	840	36007.0	Broome	New York
1837	84036009	US	USA	840	36009.0	Cattaraugus	New York
1838	84036011	US	USA	840	36011.0	Cayuga	New York
1839	84036013	US	USA	840	36013.0	Chautauqua	New York
1840	84036015	US	USA	840	36015.0	Chemung	New York
1841	84036017	US	USA	840	36017.0	Chenango	New York
1842	84036019	US	USA	840	36019.0	Clinton	New York
1843	84036021	US	USA	840	36021.0	Columbia	New York
1844	84036023	US	USA	840	36023.0	Cortland	New York
1845	84036025	US	USA	840	36025.0	Delaware	New York
1846	84036027	US	USA	840	36027.0	Dutchess	New York
1847	84036029	US	USA	840	36029.0	Erie	New York
1848	84036031	US	USA	840	36031.0	Essex	New York
1849	84036033	US	USA	840	36033.0	Franklin	New York
1850	84036035	US	USA	840	36035.0	Fulton	New York
1851	84036037	US	USA	840	36037.0	Genesee	New York
1852	84036039	US	USA	840	36039.0	Greene	New York
1853	84036041	US	USA	840	36041.0	Hamilton	New York
1854	84036043	US	USA	840	36043.0	Herkimer	New York
1855	84036045	US	USA	840	36045.0	Jefferson	New York
1856	84036047	US	USA	840	36047.0	Kings	New York
1857	84036049	US	USA	840	36049.0	Lewis	New York
1858	84036051	US	USA	840	36051.0	Livingston	New York
1859	84036053	US	USA	840	36053.0	Madison	New York

1860	84036055	US	USA	840	36055.0	Monroe	New York
1861	84036057	US	USA	840	36057.0	Montgomery	New York
1862	84036059	US	USA	840	36059.0	Nassau	New York
...
1866	84036067	US	USA	840	36067.0	Onondaga	New York
1867	84036069	US	USA	840	36069.0	Ontario	New York
1868	84036071	US	USA	840	36071.0	Orange	New York
1869	84036073	US	USA	840	36073.0	Orleans	New York
1870	84036075	US	USA	840	36075.0	Oswego	New York
1871	84036077	US	USA	840	36077.0	Otsego	New York
1872	84036079	US	USA	840	36079.0	Putnam	New York
1873	84036081	US	USA	840	36081.0	Queens	New York
1874	84036083	US	USA	840	36083.0	Rensselaer	New York
1875	84036085	US	USA	840	36085.0	Richmond	New York
1876	84036087	US	USA	840	36087.0	Rockland	New York
1877	84036089	US	USA	840	36089.0	St. Lawrence	New York
1878	84036091	US	USA	840	36091.0	Saratoga	New York
1879	84036093	US	USA	840	36093.0	Schenectady	New York
1880	84036095	US	USA	840	36095.0	Schoharie	New York
1881	84036097	US	USA	840	36097.0	Schuyler	New York
1882	84036099	US	USA	840	36099.0	Seneca	New York
1883	84036101	US	USA	840	36101.0	Steuben	New York
1884	84036103	US	USA	840	36103.0	Suffolk	New York
1885	84036105	US	USA	840	36105.0	Sullivan	New York
1886	84036107	US	USA	840	36107.0	Tioga	New York
1887	84036109	US	USA	840	36109.0	Tompkins	New York
1888	84036111	US	USA	840	36111.0	Ulster	New York
1889	84036113	US	USA	840	36113.0	Warren	New York
1890	84036115	US	USA	840	36115.0	Washington	New York
1891	84036117	US	USA	840	36117.0	Wayne	New York
1892	84036119	US	USA	840	36119.0	Westchester	New York
1893	84036121	US	USA	840	36121.0	Wyoming	New York
1894	84036123	US	USA	840	36123.0	Yates	New York
3179	84080036	US	USA	840	80036.0	Out of NY	New York

	Country_Region	Lat	Long_	...	5/29/20	5/30/20	5/31/20	\
1833	US	42.600603	-73.977239	...	1834	1843	1860	
1834	US	42.257484	-78.027505	...	45	45	45	
1835	US	40.852093	-73.862828	...	0	0	0	
1836	US	42.159032	-75.813261	...	557	561	566	
1837	US	42.247782	-78.679231	...	86	87	88	
1838	US	42.912617	-76.557316	...	89	90	91	
1839	US	42.227692	-79.366918	...	82	83	84	
1840	US	42.138911	-76.763880	...	137	137	137	
1841	US	42.494300	-75.608876	...	130	132	133	
1842	US	44.745309	-73.678754	...	95	95	95	
1843	US	42.248193	-73.630891	...	382	383	387	
1844	US	42.595092	-76.070489	...	39	41	41	

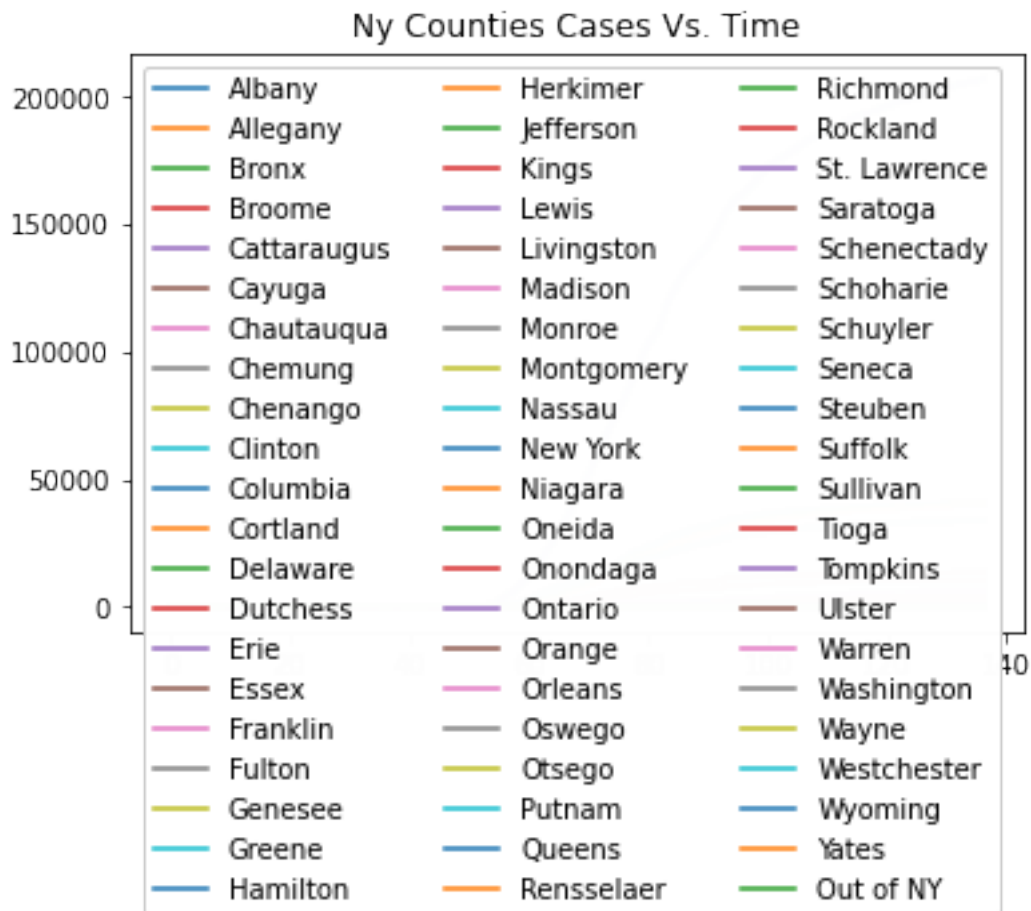
1845	US	42.198376	-74.967093	...	79	79	79
1846	US	41.764861	-73.743567	...	3887	3899	3909
1847	US	42.762490	-78.730637	...	5935	6014	6070
1848	US	44.116308	-73.772978	...	36	36	37
1849	US	44.590409	-74.299260	...	21	22	22
1850	US	43.113639	-74.417988	...	206	207	208
1851	US	43.002260	-78.191352	...	199	200	202
1852	US	42.275797	-74.123849	...	237	238	238
1853	US	43.661466	-74.497220	...	5	5	5
1854	US	43.420342	-74.961453	...	107	107	108
1855	US	44.042010	-75.946535	...	74	74	74
1856	US	40.636183	-73.949356	...	0	0	0
1857	US	43.784416	-75.449040	...	20	20	20
1858	US	42.725963	-77.779662	...	118	118	119
1859	US	42.916539	-75.672666	...	310	311	316
1860	US	43.146389	-77.693229	...	2860	2909	2942
1861	US	42.901235	-74.440116	...	84	89	91
1862	US	40.740665	-73.589419	...	40226	40307	40396
...
1866	US	43.004919	-76.199712	...	2092	2133	2170
1867	US	42.851457	-77.308744	...	205	206	207
1868	US	41.403375	-74.302408	...	10361	10389	10406
1869	US	43.251698	-78.232007	...	216	228	229
1870	US	43.427789	-76.146352	...	108	108	109
1871	US	42.634926	-75.031514	...	71	71	72
1872	US	41.426301	-73.749655	...	1241	1248	1252
1873	US	40.710881	-73.816847	...	0	0	0
1874	US	42.713481	-73.510899	...	478	486	491
1875	US	40.585822	-74.148086	...	0	0	0
1876	US	41.150279	-74.025605	...	13100	13128	13151
1877	US	44.497618	-75.065500	...	202	202	204
1878	US	43.109042	-73.866539	...	473	477	480
1879	US	42.816688	-74.052783	...	680	686	689
1880	US	42.588317	-74.443390	...	49	50	50
1881	US	42.391840	-76.877330	...	11	11	11
1882	US	42.780810	-76.824971	...	57	58	58
1883	US	42.268914	-77.382992	...	241	241	241
1884	US	40.883201	-72.801217	...	39445	39532	39643
1885	US	41.715795	-74.763946	...	1364	1375	1387
1886	US	42.168528	-76.308358	...	126	128	130
1887	US	42.449458	-76.472298	...	157	161	164
1888	US	41.890279	-74.262521	...	1663	1678	1685
1889	US	43.561730	-73.843370	...	254	255	255
1890	US	43.311538	-73.430434	...	232	235	235
1891	US	43.154944	-77.029765	...	113	113	115
1892	US	41.162784	-73.757417	...	33349	33429	33481
1893	US	42.701451	-78.221996	...	82	84	86
1894	US	42.635055	-77.103699	...	39	39	39

3179	US	0.000000	0.000000	...	0	0	0
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	6/1/20	6/2/20	6/3/20	6/4/20	6/5/20	6/6/20	6/7/20
1833	1882	1900	1920	1930	1941	1953	1961
1834	48	48	49	49	51	51	51
1835	0	0	0	0	0	0	0
1836	574	578	589	593	605	613	620
1837	89	89	90	91	92	92	94
1838	92	93	96	96	98	100	101
1839	85	86	89	95	97	99	101
1840	137	137	137	137	137	137	137
1841	133	133	133	133	133	133	134
1842	96	97	97	97	97	97	97
1843	389	391	399	400	411	412	414
1844	41	41	41	41	41	41	41
1845	81	82	82	82	84	85	85
1846	3924	3936	3951	3962	3984	3995	4000
1847	6123	6173	6234	6308	6359	6429	6486
1848	37	38	38	38	38	38	38
1849	23	23	23	23	23	23	23
1850	210	212	213	215	219	222	224
1851	202	202	202	205	205	206	208
1852	239	241	241	241	242	246	246
1853	5	5	5	5	5	5	5
1854	109	111	113	113	115	122	123
1855	74	74	74	74	75	75	77
1856	0	0	0	0	0	0	0
1857	20	20	20	20	20	20	20
1858	119	119	120	120	121	121	121
1859	316	317	319	319	323	325	327
1860	2964	2989	3048	3081	3117	3167	3190
1861	91	93	94	96	96	98	100
1862	40479	40572	40644	40713	40797	40853	40904
...
1866	2197	2228	2256	2295	2329	2375	2392
1867	208	209	215	218	219	220	221
1868	10422	10449	10460	10471	10484	10508	10514
1869	236	241	246	247	247	254	255
1870	110	110	112	112	112	114	115
1871	72	73	73	73	73	74	74
1872	1257	1262	1264	1268	1270	1274	1277
1873	0	0	0	0	0	0	0
1874	492	493	495	495	497	499	502
1875	0	0	0	0	0	0	0
1876	13185	13223	13259	13280	13297	13315	13325
1877	205	205	206	207	207	209	209
1878	483	484	489	493	496	501	502
1879	693	696	701	701	706	710	711

1880	50	51	51	51	51	54	54
1881	11	12	12	12	12	12	12
1882	59	59	60	60	61	61	61
1883	241	242	243	245	246	251	251
1884	39705	39980	40062	40153	40239	40278	40329
1885	1389	1392	1393	1405	1409	1411	1415
1886	130	131	133	133	134	134	134
1887	164	164	165	165	167	171	171
1888	1691	1696	1701	1704	1711	1714	1718
1889	255	255	256	256	256	257	257
1890	235	237	238	238	240	240	240
1891	116	116	120	121	122	123	124
1892	33552	33633	33691	33767	33854	33924	33954
1893	86	86	87	87	88	89	89
1894	39	39	39	39	39	39	39
3179	0	0	0	0	0	0	0

[63 rows x 149 columns]



Here is a video of our final simulation visualization in cytoscape. It shows of COVID theoretically spread from county to county in the state of New York.

```
[126]: Video('NewGraphDemo.mp4')
```

```
[126]: <IPython.core.display.Video object>
```

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[ ]:
```