Dummy Title By Dummies 27 JUN 2020

Abstract

On April 24th, 2020, a researcher at MIT released a working paper finding that "The Subways Seeded the Massive Coronavirus Epidemic in New York City". While the analysis in the paper has been called into question, it remains true that the role of public transportation in the spread of COVID-19 is still unknown. In this paper, we introduce an agent-based model of the New York City subway and analyze how well it can predict the spread of COVID-19 through the boroughs of New York City.

Our findings that [insert findings here] should interest public health officials looking to make policy decisions about public transportation.

[Writer's Note: Of course, this is the ideal final result. We will focus on the early infection period and I give it a 50/50 that we even get to taking into account countermeasures and ridership losses. We will make a preliminary model, improve it, and see how far we can get.]

Competencies Briefing

Purpose: Brief others on competencies so we can discuss things

Transportation Networks

specifically subways, the NYC subway. mapping, passenger flow

python, networkx, mesa

we're working with these techs.

Writer's Note: Obviously to be removed before submission

This report was last pushed to OverLeaf on 30.05.2020. You will find our latest work at the link below: https://github.com/cheung-ho-lum/NS_Epidemics_ABM_Approach/blob/

master/Report/ABM_NYC_Subway.pdf

Background

Epidemics and COVID-19

Coronavirus disease 2019 (COVID-19) is a disease caused by the SARS-CoV-2 coronavirus. Since being identified in December 2019, it has been labelled by the WHO as a pandemic, and spread around the world. Epidemics such as the 'coronavirus' have been a subject of research for centuries, and is of special interest to those working in public health. Recent waves of new research came in 2002 (SARS), 2009 (H1N1), and 2014 (Ebola). However, in these prior epidemics, researchers did not have access to as much data as we have currently. In recent years, the state of data science research tools have also greatly improved, allowing researchers to answer questions in novel ways, but following the scientific method. In this paper, we will try to respect the expertise of public health

researchers, medical professionals, and the general public by being conservative when making conclusions.

COVID-19 in NYC

In addition to general background knowledge about COVID-19, it would behoove the reader to know about the early spread of the disease in New York. Analysis of viral RNA in patients at the Mount Sinai Health System[citation] has lead researchers to conclude that the virus first came into the community through "multiple, independent but isolated introductions" from Europe and elsewhere in the USA. Below we have also given an approximate timeline of some of the most relevant events as of June 1st, 2020. We would also like to note that as forensic researchers begin examining the data, some significant new dates or corrections may arise:

- February 25 First positive test in NYC from a 39 year-old female healthcare worker flying back from Iran.
- March 3 First confirmed P2P spread in NYC.
- March 9 Mayor holds press conference and notes that there have been 16 confirmed cases.
- March 12 Mayor declares a local state of emergency.
- March 15 Schools officially close.
- March 22 State order (PAUSE) to shelter in place comes into effect.

COVID-19 and NYC Public Transportation

Meanwhile, here is a graphic of the metro ridership from scheider.com...

Epidemic Modeling

There are two basic types of modeling for epidemics. Statistical and Mechanistic. While statistical modeling has typically given more accurate forecasts for a well-known situation [citation needed], mechanistic models such as the SIR and SEIR compartmental models help to explain why phenomena such as epidemics spread the way they do, and what impact various policy decisions will have on the result.

SEIR Model

As we will be using the SEIR model in our research, we will cover it briefly here. The SEIR compartmental model is a mathematical modeling of infectious diseases where a closed population of people move successively from compartments suspected to exposed to infected and to removed. We briefly provide an explanation of each compartment and leave the equations below.

- Susceptible These people are susceptible to getting the disease from someone infected.
- Exposed These people are no longer susceptible to the disease, and do not infect others. After a latent period, they become infectious.
- Infected These people will spread the disease to susceptible people. After a period of time they are removed by recovery, hospitalization, death.
- Removed Sometimes known as resistant or recovered. We will do our modeling with the term 'removed'. These people are no longer spreading the disease.

Writer's note: just take the equations from someone's overleaf

$$dS/dt$$

$$ds/dt = -beta * s(t) * i(t)$$

$$de/dt = beta * s(t) * i(t) - alpha * e(t)$$

$$di/dt = alpha * e(t) - gamma * i(t)$$

$$dr/dt = gamma * i(t)$$

$$S + E + I + R = 1$$

Newer Compartmental Models

While we have done some research into more advanced models and are interested in cases such as super-spreaders, we believe the SEIR model to be sufficient for the modeling done in this paper. Basic SIR is insufficient because public health officials often make policy decisions based on positive case numbers. For example, an official may decide to impose strict isolation only after 100 positive cases. But by the time there are 100 cases of 'infected' people, there may be 1000 exposed people who will meaningfully impact epidemic statistics.

Epidemics on Transportation Networks

Writer's Note: epidemics on social networks blah blah linear threshold epidemics on transportation networks a little different, passenger flow is one of the most important numbers

Agent Based Models

It's just a tool. like any model. Simulating agents in an environment. Key parameters are agent, model, and environment parameters. For example, we have the following list thus far:

```
(Agent)STATUS_SUSCEPTIBLE = 'Susceptible'
(Agent)STATUS_EXPOSED = 'Exposed'
(Agent)STATUS_INFECTED = 'Infected'
(Agent)STATUS_INFECTED_ASYMPTOMATIC = 'Asymptomatic'
```

```
(Agent)STATUS_RECOVERED = 'Recovered'
(Agent)TIME_TO_RECOVER = 10
(Agent)TIME_TO_INFECTION = 3
(Model)RUN_SPAN = 60
(Model)TOTAL_POPULATION = 50000
```

The environment is new york subway mapped as a network. All of this is still being discussed.

Transportation Networks and Epidemics

Airlines

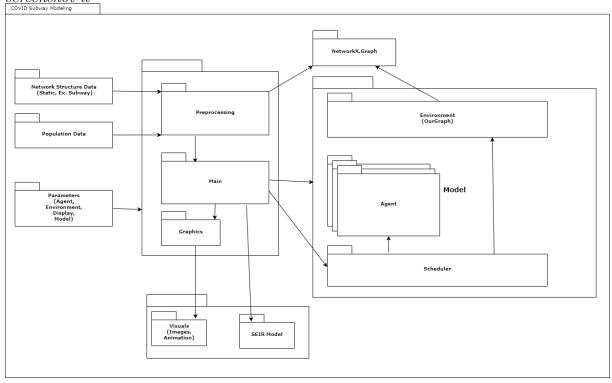
Subways and Subway Nomenclature

This guy is indispensable for figuring out how to parse some of this data: https://en.wikipedia.org/wiki/New_York_City_Subway_nomenclature The key is lines and routes

Methodology

MESA(Or our ABM)

Writer's note: I'm a bit too lazy to learn tikz-uml. I'll just make a regular uml and... screenshot it_



NetworkX

MTA Station Data

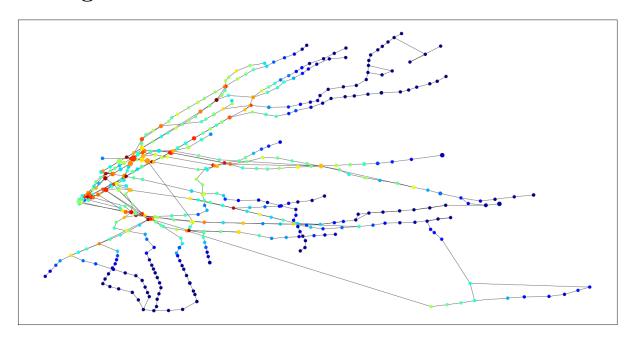
Station, Complex, Line, Route

station 167... doubled making edges. GTFS Data... no we do it ourselves

MTA Turnstile Data

discuss format. every 4 hours, every machine. aggregation.

Fitting the data



Conclusion

Writer's Note: All Models are Wrong, Some are Useful

References

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- [4] Philip Cooley, Shawn Brown, James Cajka, Bernadette Chasteen, Laxminarayana Ganapathi, John Grefenstette, Craig R. Hollingsworth, Bruce Y. Lee, Burton Levine, William D. Wheaton, and et al. The role of subway travel in an influenza epidemic: A new york city simulation. *Journal of Urban Health*, 88(5):982–995, Sep 2011.
- [5] Ana S Gonzalez-Reiche, Matthew M Hernandez, Mitchell Sullivan, Brianne Ciferri, Hala Alshammary, Ajay Obla, Shelcie Fabre, Giulio Kleiner, Jose Polanco, Zenab Khan, and et al. Introductions and early spread of sars-cov-2 in the new york city area. Nov 2020.
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- [7] M. Laskowski, B. C. P. Demianyk, J. Witt, S. N. Mukhi, M. R. Friesen, and R. D. Mcleod. Agent-based modeling of the spread of influenza-like illness in an emergency department: A simulation study. *IEEE Transactions on Information Technology in Biomedicine*, 15(6):877–889, 2011.
- [8] Faïçal Ndaïrou, Iván Area, Juan J Nieto, and Delfim F M Torres. Mathematical modeling of covid-19 transmission dynamics with a case study of wuhan, Apr 2020.