State-level variation for initial COVID-19 dynamics in the United States: The role of local government

interventions

EASTON R. WHITE^{1,2}, LAURENT HÉBERT-DUFRESNE ^{3,4}

²Department of Biology, University of Vermont, VT 05405, USA

³ Vermont Complex Systems Center, University of Vermont, VT 05405, USA

⁴Department of Computer Science, University of Vermont, VT 05405, USA

Abstract

In the development of an epidemic, metrics such as R_0 , doubling time, and case fatality

rates are important in understanding and predicting the course of an epidemic. However, if

collected over country or regional scales, these metrics hide important smaller-scale, local

dynamics. We examine how commonly used epidemiological metrics differ for each individual

state within the United States during the initial COVID-19 outbreak. We found that the

case number, and trajectory of cases, differs considerably between states. We show that early

non-pharmaceutical, government actions, were the most important determinant of epidemic

dynamics. Although individual states are clearly not independent, they can serve as mini,

natural experiments in how different demographic patterns and government responses can

impact the course of an epidemic. Thus, these results should be used to better understand,

in near real-time, what actions are working most effectively.

Keywords: SARS-CoV-2, COVID-19, spatial heterogeneity, doubling time

Daily updates to figures in this manuscript are available at: https://github.com/eastonwhite

1

/COVID19 US States

¹Easton.White@uvm.edu

Introduction

The global SARS-CoV-2 (COVID-19) pandemic began in Wuhan, China in late 2019 (WHO 2020). As of April 06th, 1,345,101 cases have been reported across 184 countries and regions. There has been several sets of efforts to track the progression of the outbreak across the world and within countries. For example, John Hopkins University Center for Systems Science and Engineering (CSSE) has compiled data from various sources, including the US Center for Disease Control and the World Health Organization, to present a global picture of COVID-19 cases and deaths (Dong et al. 2020). These efforts have allowed for international scientific research and political decision-making. Although data are collected at local scales (e.g. within hospitals), in an emerging pandemic data is typically reported at the country level. This allows for interesting comparisons between countries (Anderson et al. 2020) and for information from an earlier affected country to be used to slow the outbreak in other places. For instance, South Korea was able to flatten their outbreak curve through early and widespread testing as well as strict quarantine policies (citation). However, country-level analyses still hide more local dynamics that are important to the overall epidemic progression. Spatial heterogeneity is important for population dynamics generally (Levin 1992) and in particular for understanding the progression of infectious disease dynamics (Grenfell et al. 1995). Spatial heterogeneity can include differences in local population density, movement patterns, suitability of environmental conditions for transmission, among other factors. For instance, Keeling et al. (2001) showed how spatial distribution and size of farms affected the 2001 UK Foot and Mouth Epidemic.

Here we examine the progression of COVID-19 for state-level differences within the United States. We examine how commonly-used metrics, including doubling time, can vary state to state and compared to the US as a whole. Although not independent units, we can use state-level data to understand the progression of the outbreak across different replicates within a country. We then show that demography, education, wealth, and other hypotheses

were poor predictors of doubling time. Instead, we show that doubling time was more tightly correlated with state-level governmental actions, including restricting businesses. We also provide a link to daily updates of our findings as the exact quantitative results are likely to change over the course of the epidemic.

Results and Discussion

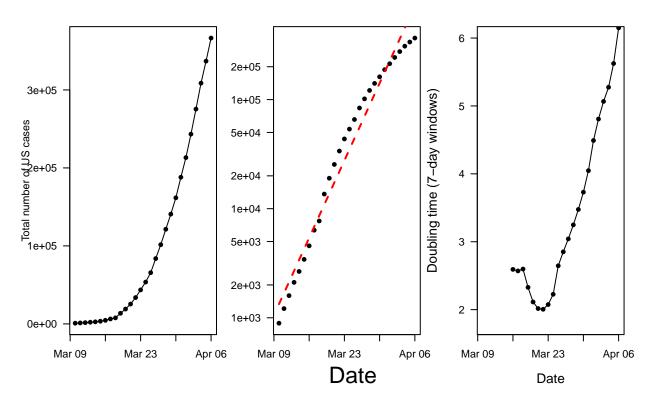


Figure 1: (Left panel) Cases versus time for the whole United States. (Right panel) Log number of cases versus time for the whole United States. The red, dashed line is the line of best fit for all the data and the blue, solid line is the line of best fit since February 29th.

We used data compiled by John Hopkins University Center for Systems Science and Engineering (Dong *et al.* 2020). The United States has seen exponential growth in the number of cases, especially since February 29th (Fig. 1). Country-level results, however, hide underlying dynamics within each state.

Therefore, we examined how the number of cases changed over time within each state. To

properly compare the progression of the epidemic across states, we looked at the log number of cases since the first day a state reported 25 cases (Fig. 2). On a log scale, a straight line of the cases over time indicates exponential growth where the slope of the line is the exponential growth parameter.

State-level variation in COVID-19 trajectories

We found considerable differences between states in how the outbreak has progressed (Fig. 2). States like New York, New Jersey, and Michigan have experienced a doubling of cases approximately every two days. Conversely, Massachusetts has experienced a doubling time closer to five days. We mapped doubling time across the US and found regional differences where the West Coast and Northeast have seen large doubling times, i.e. slower outbreak dynamics (Fig. 3).

Predictors of early (pre-intervention) state-level trajectories - NEED TO WORK ON THIS NEXT

Each US state varies considerably across a number of important axes: wealth, access to healthcare, number of international travelers, age distribution, population density, among other factors. In addition, much of the response to COVID-19 has been done at the state, as opposed to federal, government level in the US (citations). These responses have varied a lot with different laws and such... (citations).

We had two hypotheses to explain the state-level variation in COVID-19 trajectories before government intervention occurred: human demographics as well as wealth and education indicators.

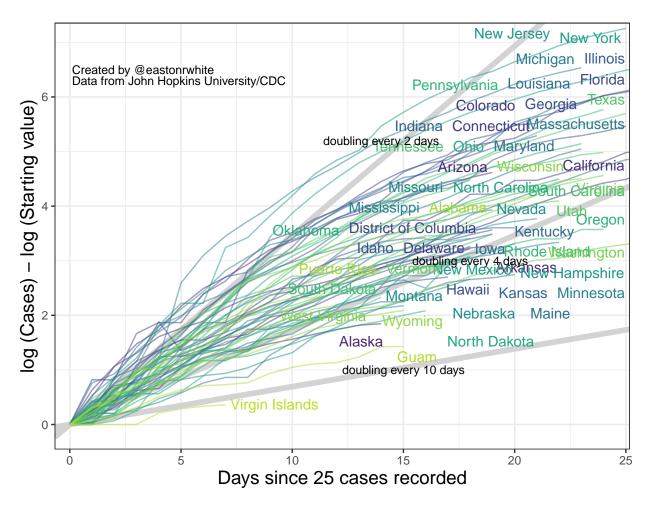
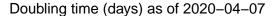
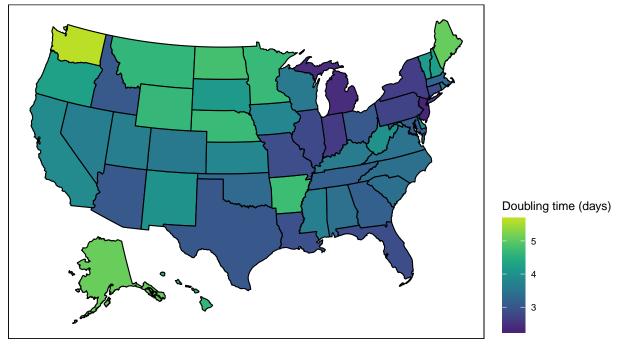


Figure 2: The log number of cases over time for each individual state that recorded more than 25 cases over at least three days. The light grey diagonal lines represent the growth trajectory for doubling times of 2, 4, and 10 days. The log number of the starting value (intial number of cases on first day when at least 25 cases were recorded) had to be subtracted on the y-axis to standarize the graph across states.





Created by: @eastonrwhite, Data from John Hopkins University/CDC

Figure 3: Doubling time (in number of days) across the US states that recorded more than 25 cases over at least three days.

Predictors of overall state-level trajectories

We had three hypotheses to explain the state-level variation in COVID-19 trajectories: human demographics, wealth and education indicators, and governmental actions.

We found that demography, education, and wealth were poor predictors of the state-level doubling times (Fig. 4). Therefore, we also examined the correlation between doubling time and state government interventions, and timing of those interventions. We specifically looked at whether or not a state had implemented a specific action by the first day they had 25 or more cases. We looked at closing of public schools, limiting large gatherings (usually of more than 10 people), restricting business, and stay at home orders. Of these factors, only restricting businesses was a significant predictor of doubling time (Fig. 5).

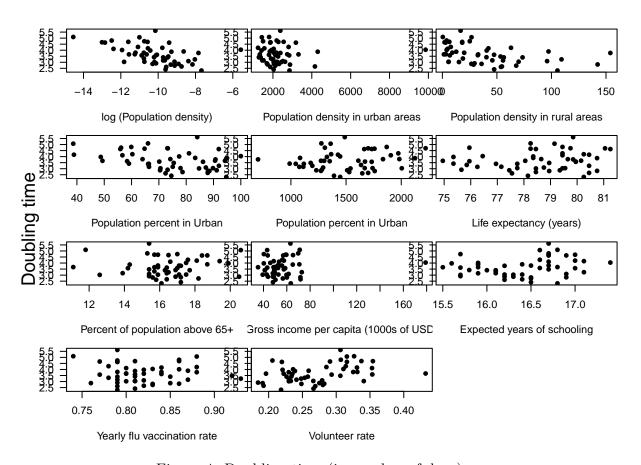


Figure 4: Doubling time (in number of days)....

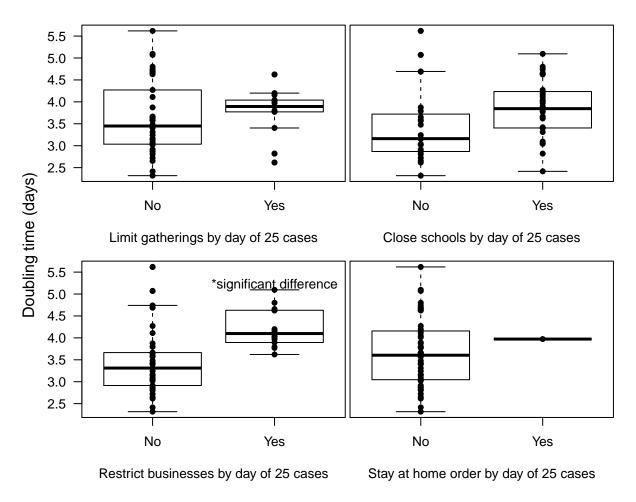


Figure 5: Doubling time (in number of days) across the US states that recorded more than 25 cases over at least three days.

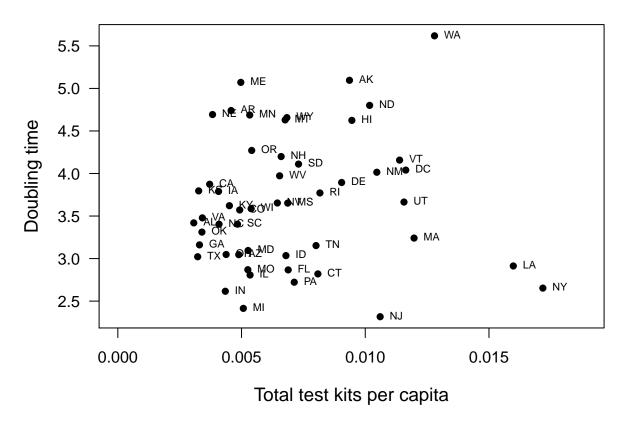


Figure 6: Doubling time (in number of days) across the US states that recorded more than 25 cases over at least three days.

Conclusions and Future Work

We found a large degree of heterogeneity in the number of cases over time across US states. After state-level government actions were implemented, doubling time was most strongly correlated to restrictions on businesses. More detailed work will be needed to understand how these dynamics differ within each state, especially as many government actions started on these more local scales.

Code availability and acknowledglements

All code and corresponding data is freely available at https://github.com/eastonwhite/COV ID19_US_States. The original raw data has been compiled by the Johns Hopkins University Center for Systems Science and Engineering at (https://github.com/CSSEGISandData/CO VID-19). The authors received no specific funding for this work.

References

- Anderson, R.M., Heesterbeek, H., Klinkenberg, D. & Hollingsworth, T.D. (2020). How will country-based mitigation measures influence the course of the COVID-19 epidemic? *The Lancet*, 395, 931–934.
- Dong, E., Du, H. & Gardner, L. (2020). An interactive web-based dashboard to track COVID-19 in real time. *The Lancet Infectious Diseases*, S1473309920301201.
- Grenfell, B.T., Bolker, B.M. & Kleczkowski, a. (1995). Seasonality and Extinction in Chaotic Metapopulations. *Proceedings of the Royal Society B: Biological Sciences*, 259, 97–103.
- Jombart, T., van Zandvoort, K., Russell, T., Jarvis, C., Gimma, A. & Abbott, S. *et al.* (2020).

 Inferring the number of COVID-19 cases from recently reported deaths. *medRxiv*.

- Keeling, M.J., Woolhouse, M.E.J., Shaw, D.J., Matthews, L., Chase-Topping, M. & Haydon, D.T. et al. (2001). Dynamics of the 2001 UK foot and mouth epidemic: Stochastic dispersal in a heterogeneous landscape. Science, 294, 813–817.
- Levin, S.A. (1992). The Problem of Pattern and Scale in Ecology. Ecology, 73, 1943–1967.
- Park, A.W. (2012). Infectious disease in animal metapopulations: The importance of environmental transmission. *Ecology and Evolution*, 2, 1398–1407.
- Schreiber, S.J. (2010). Interactive effects of temporal correlations, spatial heterogeneity and dispersal on population persistence. *Proceedings of the Royal Society B: Biological Sciences*, 277, 1907–1914.
- WHO. (2020). Report of the WHO-China Joint Mission on Coronavirus Disease 2019 (COVID-19).