

# Documenting Social, Geographic, and Economic Inequities in COVID-19 Mortality at the County Level in the U.S. Using Generalized Additive Models

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## Abstract

We present three types of applications of generalized additive models (GAMs) to COVID-19 mortality rates in the US for the purpose of advancing methods to document inequities with respect to which communities suffered disproportionate COVID-19 mortality rates. First, GAMs can be used to describe the changing relationship between COVID-19 mortality and county-level covariates (sociodemographic, economic, and political metrics) over time. Second, GAMs can be used to perform spatiotemporal smoothing that pools information over time and space so that county level mortality rates that tend to be noisy due to small population counts or stochasticity can be summarized by a smooth, dynamic latent surface describing the mortality risk associated with geographic locations over time. Third, estimation of the associations between county-level covariates and COVID-19 mortality after controlling for spatiotemporal effects allow for the distinguishing of what patterns in COVID-19 mortality were more plausibly due to geography than variation in county-level covariates. Each of these approaches provides a valuable approach and perspective to documenting inequities in COVID-19 mortality by addressing the question of which populations have suffered the worst burden of COVID-19 mortality taking into account the nonlinear spatial, temporal, and social patterning of disease.

Abbreviations used: United States (US), Coronavirus Disease 2019 (COVID-19), Generalized Additive Model (GAM), Centers for Disease Control and Prevention (CDC)

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## Introduction

As we enter the third winter with the novel coronavirus disease COVID-19 in the United States, evidence documenting the intense disparities in COVID-19 mortality rates comparing socially advantaged and disadvantaged populations continues to mount. Eliminating inequities in health outcomes has been stated as a major policy goal of the Biden administration (The White House, 2021, 2022), representing a revitalized focus on health justice and underlining the importance of adequate data reporting systems that measure the prevalent health inequities. To this end, we use generalized additive models (GAMs) in a flexible regression framework to illustrate the evolving roles and relationships sociodemographic, geographic, and economic conditions have with respect to trends in COVID-19 mortality. The code, data, and documentation necessary to reproduce the analyses contained in this paper are online and free to access at <https://github.com/catesta01/covid.gradient.estimation>.

## Background

Having passed over 1 million COVID-19 deaths in the United States earlier this year (Donovan, 2022), and facing uncertain prospects for the third COVID-19 winter looming even as new iterations on the COVID-19 vaccines become available, it remains critical that inequities in COVID-19 outcomes are documented and

analyzed to reckon with the unjust and unfair burden of preventable illness. Even though the first vaccines were granted emergency use authorization by the U.S. Food and Drug Administration in 2020 (Mayo Clinic, 2022), with the first shots going in arms in December 2020, COVID-19 is still continuing to cause hundreds of deaths a day in the U.S. in the fall of 2022 (“United States COVID - Coronavirus Statistics - Worldometer,” 2022). The new bivalent vaccines released at the end of August 2022 contain mRNA sequences from both the original strain as well as the recently emergent BA.4 and BA.5 lineages in an effort to make the nation’s immunity more up-to-date and robust against the myriad of phylogenetic directions the COVID-19 virus is evolving to explore (Office of the Commissioner, 2022). Despite the updated bivalent boosters representing a significant step forward in prevention strategy, less than 4% of eligible Americans had taken the booster in the first month after it became available (Bendix, 2022; Lambert, 2022). As such, and with an enduring history of inequities in health care availability (Bailey et al., 2021; Blendon et al., 2002; Carpenter, 2021; Chrisler et al., 2016; Feldman et al., 2021; Okonkwo et al., 2021; Ortega & Roby, 2021; Rapp et al., 2022; Whitehead et al., 2016), it is clear that without further intervention not all communities will be equally able to benefit from the new vaccines available and inequities in COVID-19 illness and mortality may persist despite the technological innovations in vaccine technology.

## The Role of Geography in COVID-19

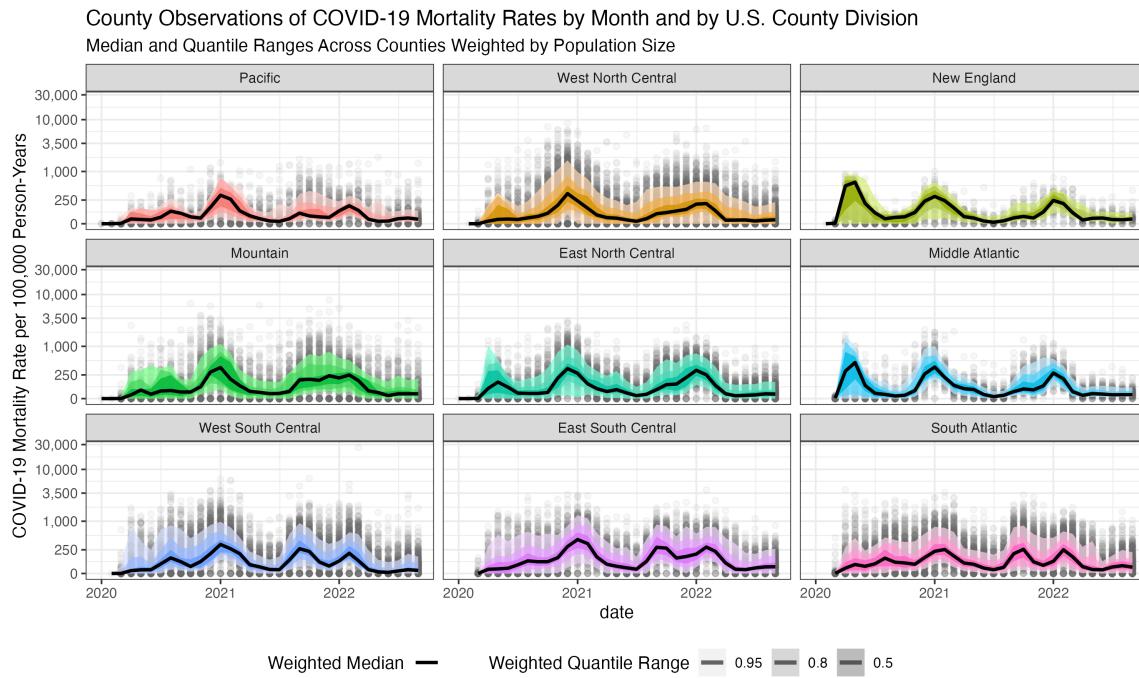


Figure 1: Estimates of monthly COVID-19 mortality rates per 100,000 person-years by county organized by Census Division. For each division, the median trendline and quantile ranges are shown weighted by county population size.

Prior literature has demonstrated that geography has played a significant role in the spread and impact of COVID-19. Methods employed to highlight the role geography plays have included quantile regression (Sigler et al., 2021), Besag-York-Mollie’s mixed models (Whittle & Diaz-Artiles, 2020), spatial cluster analysis (Sugg et al., 2021), geographically weighted regression (Mollalo et al., 2020; Park et al., 2021), and others.

In the U.S. context, one of the key aspects to the geographic story of COVID-19’s spread and diffusion was the early surge of cases and epicenter in New York City during March 2020 (Thompson, 2020) followed by subsequent waves of cases in the South and Midwest (Glenza, 2020; Scott, 2020; Shumaker & Wu, 2020). As Park et al. stated in their article *COVID-19 Deaths in the United States: Shifts in Hot Spots over the Three Phases of the Pandemic and the Spatiotemporally Varying Impact of Pandemic Vulnerability* where

they summarized US trends from March 2020 to May 2021, “hot spots have shifted from densely populated cities and the states with a high percentage of socially vulnerable individuals to the states with relatively relaxed social distancing requirements, and then to the states with low vaccination rates” (Park et al., 2021).

When considering the drivers of the COVID-19 pandemic, it’s necessary to note that geography and social conditions are inextricably linked. In July 2021, the CDC reported that “the COVID-19 cumulative death rate in non-metropolitan areas has exceeded that of metropolitan areas since December 2020,” noting that of the approximately 1/5th of Americans who live in rural areas, many “are considered highly vulnerable according to CDC’s Social Vulnerability Index (SVI), which includes factors such as housing, transportation, socioeconomic status, race, and ethnicity” (CDC, 2021). Moreover, rural communities often have lower health insurance rates, higher disability rates, older populations, and limited access to health care. One of CDC’s Morbidity and Mortality Weekly Reports found that vaccination against COVID-19 was lower in rural communities than in urban communities between December 2020 and April 2021 (Murthy, 2021).

Figure 1 shows the monthly COVID-19 mortality rates for the counties grouped within each of the nine U.S. Census Divisions (U.S. Department of Commerce Economics and Statistics Administration & U.S. Census Bureau, 2000). The figure summarizes each division’s median mortality rates weighted by county population size. Quantile ranges are included to illustrate the range of mortality rates observed, again weighted by county population size. Weighting based on county population size is used as opposed to equal weighting for calculating the median and quantile ranges so that in divisions where low population counties predominate, mortality rate estimates that are noisy due to small population counts are not weighted equally to more reliable mortality rate estimates from larger population counties. Notably, the mortality associated with the early surge of cases starting in NYC and spreading through New York, New Jersey, and Massachusetts is visible in the Middle Atlantic and New England division figures. The figure also illustrates how the first peak in the mortality time-series for states in the Midwest (West North Central, East North Central) occurred later, in late 2020 and going into early 2021.

### The Social Determinants of COVID-19 Mortality

- Insert section on prior modeling results
- Insert discussion motivating the need for spatiotemporal modeling

## Methods

### Data Sources

The following variables were retrieved at the county level:

- Counts of COVID-19 deaths (The New York Times, 2021).
- Population size estimates for 2020 from the U.S. Census (Redistricting File Public-Law 94-171 Dataset) (US Census Bureau, 2021).
- Median age, median household income, racial/ethnic composition, population density, percent below the federal poverty line, and number of households with high (\$100k+)/low (<\$25k) household income by racial/ethnic group from the 2014-2019 5-year American Community Survey (US Census Bureau, 2020) through the `tidyverse` R package (Walker & Herman, 2022).
- Votes cast in the 2020 presidential election (`mit_county_2022?`)

### Generalized Additive Models

Traditional generalized linear models fit a regression model using the functional relation  $g(\mu_i) = \mathbf{X}_i\beta$  where Examples of common link functions include the identity function for linear regression, log for Poisson regression, and the logit function for logistic regression.

Generalized additive models (GAMs) improve upon generalized linear models by allowing for the fitting of smooth functions that transform the  $x$  variables. This is a convenient means to include nonlinear relationships

between the outcome and predictor variables. Whereas a generalized linear model may have looked like

$$g(\mu_i) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_3 x_{i3} \dots$$

a generalized additive model could look like

$$g(\mu_i) = \mathbf{A}_i \theta + f_1(x_{i1}) + f_2(x_{i2}) + f_3(x_{i3}, x_{i4}) + \dots$$

where  $\mu_i \equiv \mathbb{E}(Y_i)$ ,  $\mathbf{A}_i$  is a row of the model matrix for any strictly non-parametric model components,  $\theta$  is the corresponding parameter vector, and the  $f_j$  are smooth functions of the covariates  $x_k$  (Wood, 2017).

## Results

### Evaluating Spatial and Temporal Autocorrelation

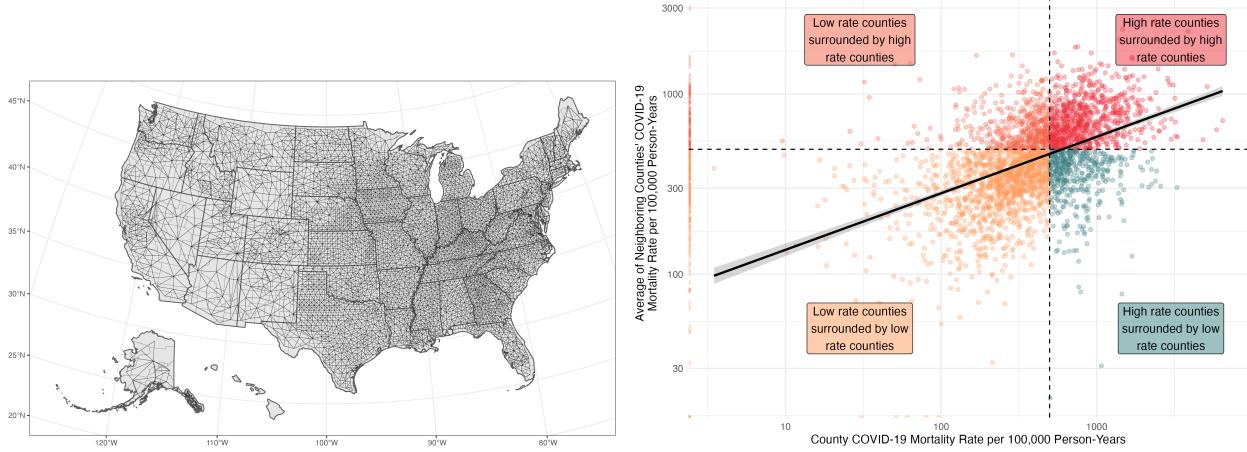


Figure 2: A) The spatial neighbor graph for counties in the US. In this graph, county centroids are connected if the two counties are adjacent to one another (i.e., share a boundary). This and similar graphs that represent neighbor relationships or geographic proximity are often used in modeling contexts to create spatial weights that account for which regions help to explain a given region's observed data.

(#fig:figure county neighboring graph)

### Regression Modeling Results

We used GAMs to describe the following:

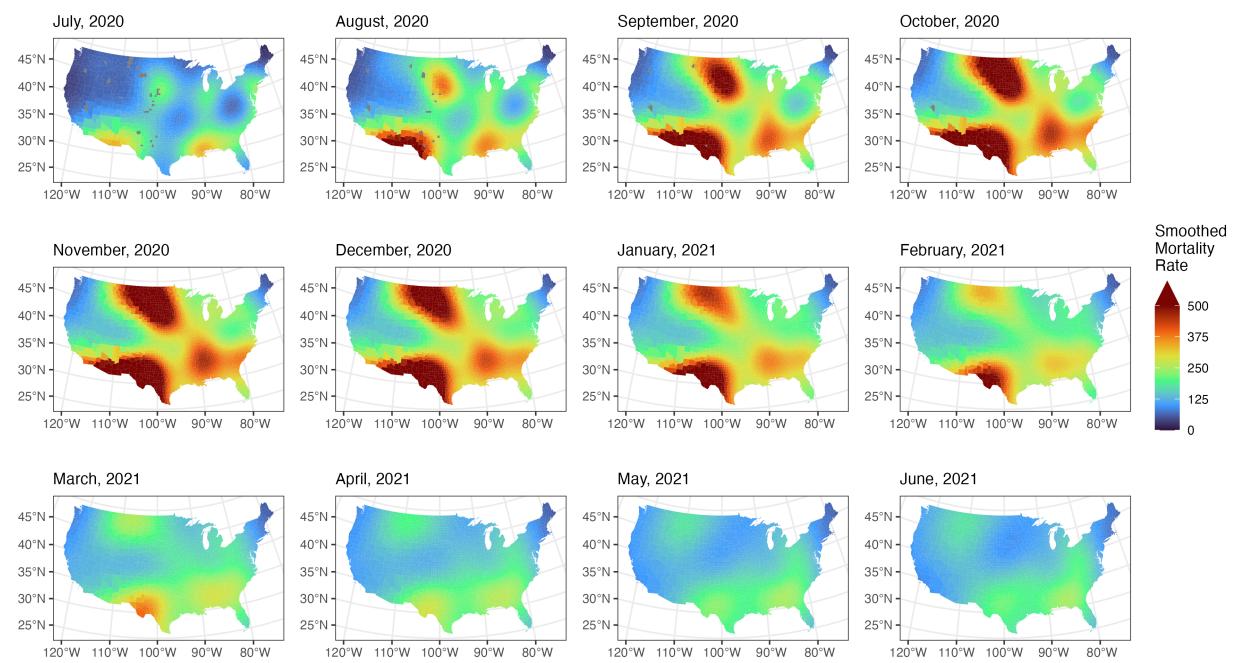
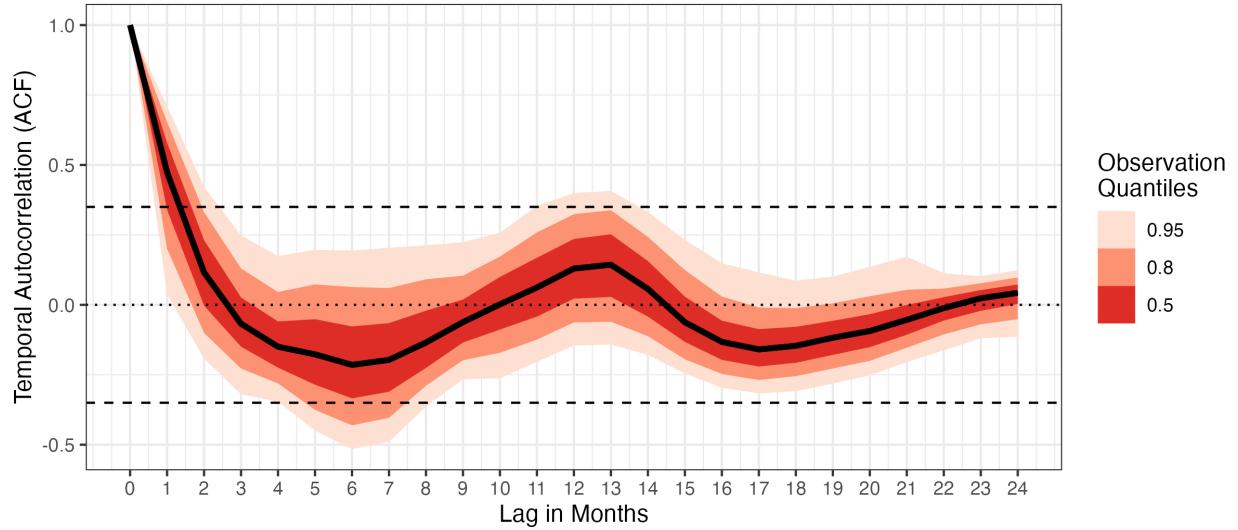
- the changing associations between individual area-based measures and COVID-19 mortality over time;
- the changing associations between bivariate area-based measures and COVID-19 mortality over time;
- associations with area-based measures and COVID-19 mortality after controlling for spatio-temporal autocorrelation.

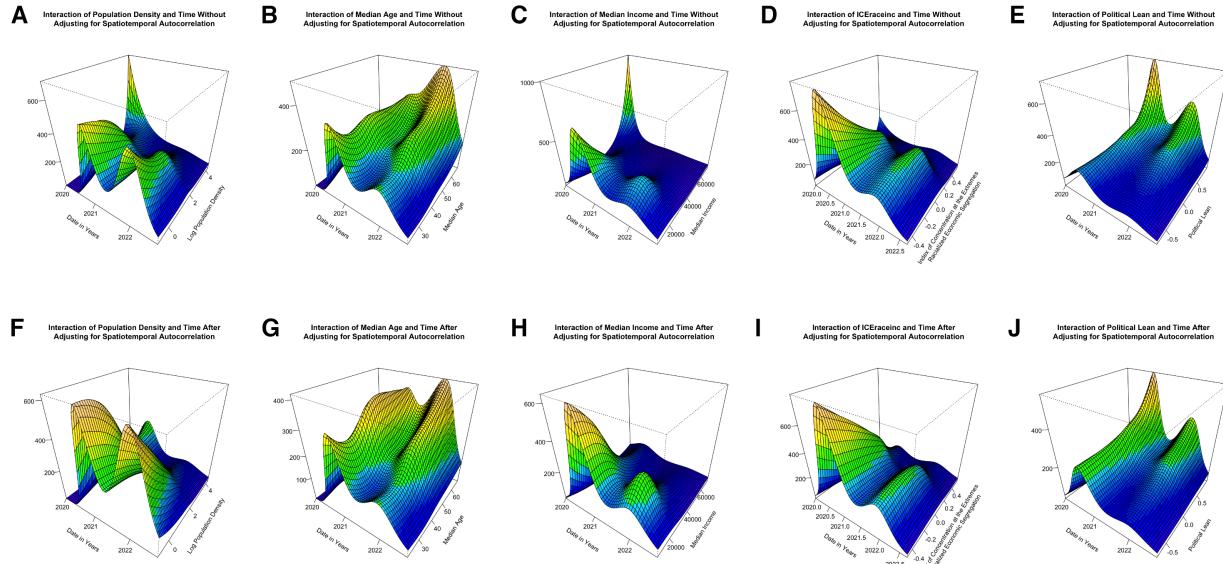
Additionally, as sensitivity analyses, random forest models were fit to compare with the coefficient estimates from the GAMs.

## Discussion

### Limitations

- Missing data
- Ecological fallacy





- Age effects
- Fixed population denominators -> real-time denominator estimates
- Vaccination and mobility effects

## Conclusions

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