



Original Investigation

Projecting COVID-19 Mortality as States Relax Nonpharmacologic Interventions

Benjamin P. Linas, MD, MPH; Jade Xiao, BE; Ozden O. Dalgic, PhD; Peter P. Mueller, PhD; Madeline Adee, MPH; Alec Aaron, MPH; Turgay Ayer, PhD; Jagpreet Chhatwal, PhD

Abstract

IMPORTANCE A key question for policy makers and the public is what to expect from the COVID-19 pandemic going forward as states lift nonpharmacologic interventions (NPIs), such as indoor mask mandates, to prevent COVID-19 transmission.

OBJECTIVE To project COVID-19 deaths between March 1, 2022, and December 31, 2022, in each of the 50 US states, District of Columbia, and Puerto Rico assuming different dates of lifting of mask mandates and NPIs.

DESIGN, SETTING, AND PARTICIPANTS This simulation modeling study used the COVID-19 Policy Simulator compartmental model to project COVID-19 deaths from March 1, 2022, to December 31, 2022, using simulated populations in the 50 US states, District of Columbia, and Puerto Rico. Projected current epidemiologic trends for each state until December 31, 2022, assuming the current pace of vaccination is maintained into the future and modeling different dates of lifting NPIs.

EXPOSURES Date of lifting statewide NPI mandates as March 1, April 1, May 1, June 1, or July 1, 2022.

MAIN OUTCOMES AND MEASURES Projected COVID-19 incident deaths from March to December 2022.

RESULTS With the high transmissibility of current circulating SARS-CoV-2 variants, the simulated lifting of NPIs in March 2022 was associated with resurgences of COVID-19 deaths in nearly every state. In comparison, delaying by even 1 month to lift NPIs in April 2022 was estimated to mitigate the amplitude of the surge. For most states, however, no amount of delay was estimated to be sufficient to prevent a surge in deaths completely. The primary factor associated with recurrent epidemics in the simulation was the assumed high effective reproduction number of unmitigated viral transmission. With a lower level of transmissibility similar to those of the ancestral strains, the model estimated that most states could remove NPIs in March 2022 and likely not see recurrent surges.

CONCLUSIONS AND RELEVANCE This simulation study estimated that the SARS-CoV-2 virus would likely continue to take a major toll in the US, even as cases continued to decrease. Because of the high transmissibility of the recent Delta and Omicron variants, premature lifting of NPIs could pose a substantial threat of rebounding surges in morbidity and mortality. At the same time, continued delay in lifting NPIs may not prevent future surges.

JAMA Health Forum. 2022;3(4):e220760. doi:10.1001/jamahealthforum.2022.0760

Key Points

Question What is the expected trend in COVID-19 mortality if US states were to lift nonpharmacologic interventions (NPIs) at different times over the remainder of 2022?

Findings In this simulation modeling study, lifting NPIs was likely to result in rebounding epidemics regardless of the delay in lifting. The degree of population-level immunity was associated with the size of the rebounding peak in incident deaths.

Meaning This simulation study found no path to the end of the COVID-19 pandemic that avoided difficult trade-offs between prolonged NPIs and increased COVID-19 mortality following their removal.

+ Supplemental content

Author affiliations and article information are listed at the end of this article.

Open Access. This is an open access article distributed under the terms of the CC-BY License.

Introduction

The emergency authorization and dissemination of SARS-CoV-2 vaccines starting in late 2020 fundamentally changed the epidemiology of the COVID-19 pandemic. Leading up to the summer of 2021, nearly every state enjoyed falling case rates while simultaneously relaxing nonpharmacologic interventions (NPIs), such as mask mandates and restrictions on social gatherings. This dissociation of social mobility and COVID-19 case rates was a categorical shift that implied a pending end to the pandemic. Unfortunately, around this time, the Delta variant entered circulation in the US and quickly became the dominant SARS-CoV-2 strain. The period of falling case rates was ended by the “fourth wave,” even in states that had achieved relatively high levels of vaccination. Next, even as the Delta variant was still spreading rapidly, the Omicron surge arrived, driving case rates to the highest levels seen in the pandemic and forcing some jurisdictions to reinstate mitigation measures.

The return of NPIs and ongoing need to integrate COVID-19 risk into everyday decision-making have greatly added to pandemic fatigue in the US. Now, as the Omicron wave begins to recede, many states are once again lifting mandatory NPIs, including indoor capacity limits and guidance on social distancing. In particular, local decision makers face the difficult decision of when to lift mask mandates. One of the most important questions currently on the minds of citizens, public health officials, and policy makers is: when can we safely lift restrictions?

We used the COVID-19 Policy Simulator,¹ a compartmental model of SARS-CoV-2 transmission and COVID-19 disease in the US, to project rates of hospitalization and death over the course of the 2022 calendar year assuming different dates of lifting NPIs.

Methods

Model Overview

Our model is an extension of the traditional SEIR model, which partitions a population into compartments representing mutually exclusive disease states: susceptible, exposed, infected, recovered, and deceased. In this model, the flow of people between compartments was assumed to obey a system of deterministic ordinary differential equations. The time step was set 1 day to be compatible with data sources reporting daily data. The model was calibrated to historical trends in daily incident deaths up to February 20, 2022. The **Table** displays the values and data sources for select model parameters. A full model specification is provided in the eAppendix in the [Supplement](#). Where applicable, we followed the Consolidated Health Economic Evaluation Reporting Standards (CHEERS) framework for communicating results. The study was exempt from institutional review board review because it used only publicly available data and was not human participants research.

Age Stratification

We stratified the population into 2 age groups: younger than 65 years (lower risk) and 65 years and older (higher risk), assuming that the size of these subpopulations was constant over the simulation period. Age stratification allowed the model to capture differential vaccination trends and COVID-19 mortality between the age groups.

Mortality

We estimated COVID-19–associated mortality using an infection fatality rate (IFR). The IFR is age specific: 0.5% for the younger than 65 years age group and 3.0% for the 65 years and older age group.

Vaccination

To reflect 2-dose administration guidelines of the COVID-19 messenger RNA vaccines, we stratified the disease states by vaccination status: 0 doses (unvaccinated), 1 dose (partially vaccinated), and 2 doses (fully vaccinated). The third vaccine, the viral vector vaccine, approved for a single-dose

regimen, was omitted from the model owing to its accounting for only 3.7% of all administered doses in the US as of June 2021.¹¹ Because there are no data on vaccination status at the time of infection, we assumed doses were allocated proportionally to the susceptible and recovered compartments over the historical time horizon. The vaccine reduces both susceptibility to infection and mortality risk. After the first and second vaccine doses, the probability of contracting the virus was reduced by 46% and 92%, respectively; similarly, the IFR was reduced by 48% and 37%, respectively (see eAppendix in the Supplement for derivation). Note that it is the conditional probability of death that is higher after the second dose than after the first dose, such that the overall reduction in COVID-19 mortality is 72% and 95%, respectively. Vaccine effectiveness was assumed to decrease as the Delta and Omicron variants enter circulation. We defined *effective immunity* as the sum of the proportion of the populations in the unvaccinated, partially vaccinated, and fully vaccinated susceptible states, weighted by their susceptibility to infection.

Transmission

For a susceptible individual, the rate of exposure to the virus was dependent on the individual's risk group, vaccination status, time-varying effective reproduction number, and the size of the infected subpopulation. We estimated an age-stratified matrix of contact patterns among and between the age groups (Table).

Table. Values of Select Parameters Used in the COVID-19 Policy Simulator Model

Parameter	Estimate	Notes	Reference
Fixed parameters			
Size of the subpopulations <65 y (lower risk) and ≥65 y (higher risk)	State dependent	NA	US Census Bureau, ² 2021
Contact matrix			
LL	0.93	Aggregate columns and rows into age groups <65 y and ≥65 y, then normalize so that rows sum to 1.	Prem et al, ³ 2017
LH	0.07		
HL	0.48		
HH	0.52		
Period, d			
Latent	5.5	NA	Xin et al, ⁴ 2019
Infectious	10	NA	Byrne et al, ⁵ 2020
Mean (exponentially distributed) duration of natural and vaccine-conferred immunity, mo	16	NA	Townsend et al, ⁶ 2021
Effective reproduction number when all NPIs are removed	5.0	NA	Liu and Rocklöv, ⁷ 2021
Calibrated parameters			
Time-varying effective reproduction number	0.5-6.0	Widely varying by location and SARS-CoV-2 variant	Liu and Rocklöv, ⁷ 2021
Initial number of infectious people at the start of the simulation (March 15, 2020)	100-10 000	Calibrated and divided proportionally into the low-risk and high-risk groups	NA
Variant-dependent parameters (see eAppendix in the Supplement for derivation and changes associated with the Delta and Omicron variants)			
Baseline, %			
IFR of the low-risk/high-risk group	0.1/3.0	These values chosen to approximate the CDC's estimated total infections ⁸	Based on this meta-analysis ⁹
Reduction in susceptibility to infection after the 1st/2nd vaccine dose	46/92	NA	Dagan et al, ¹⁰ 2021
Reduction in IFR after the 1st/2nd vaccine dose	48/37	It is the conditional probability of death that is higher after the second dose than after the first dose. If a fully vaccinated individual contracts a breakthrough infection despite 92% reduction in susceptibility, it is plausible that they are particularly vulnerable and have a smaller reduction in mortality risk conditional on infection compared with a partially vaccinated individual who contracts a breakthrough infection.	Dagan et al, ¹⁰ 2021

Abbreviations: CDC, Centers for Disease Control and Prevention; IFR, infection fatality rate; HH, high-high risk; HL, high-low risk; LH, low-high risk, LL, low-low risk; NA, not applicable; NPI, nonpharmacologic intervention.

Waning Immunity

An individual who has recovered from natural infection would experience a period of natural immunity before transitioning back into the susceptible state. A fully vaccinated susceptible individual would be protected for the duration of vaccine-conferred immunity before transitioning back into the partially susceptible state. Finally, because individuals with natural immunity who are subsequently vaccinated have been reported to exhibit “unusually potent immune responses,”¹² a fully vaccinated recovered individual was assumed to possess 2 “layers” of immunity, shedding first their natural immunity then their vaccine-conferred immunity. At present, there are no certain estimates of the mean duration of natural and vaccine-conferred immunity. We used the results of a study that examined the immune responses to evolutionarily similar viruses to estimate their time to reinfection under endemic conditions.⁶ Reinfection by endemic SARS-CoV-2 was expected to occur between 3 months and 5 years after peak antibody response, with a median of 16 months.

Booster Shots

It was assumed that, once vaccinated, an individual would never shed their immunity completely (within the time frame of the simulation), and a fully vaccinated individual who has shed their vaccine-conferred immunity would be indistinguishable from a partially vaccinated individual. Thus, the model differentiated between the subpopulation that was willing to receive booster shots and the subpopulation that was unwilling to be vaccinated. Fully vaccinated individuals would wane into the partially vaccinated state and would be “boosted” back into the fully vaccinated state.

Scenario Analysis

We projected epidemiologic trends assuming that current rates of infection and vaccination would continue until the date of lifting NPIs, which are the beginning of each calendar month from March to July 2022. We allowed the most recent calibrated value of the effective reproduction number to persist until the lifting date, after which it was increased to the assumed value of 5.0, similar to the basic reproduction number of the Delta variant, representing unmitigated transmission of the virus. We also present projections assuming a lower value of the effective reproduction number of 3.0, similar to the transmissibility of the ancestral strains.⁷

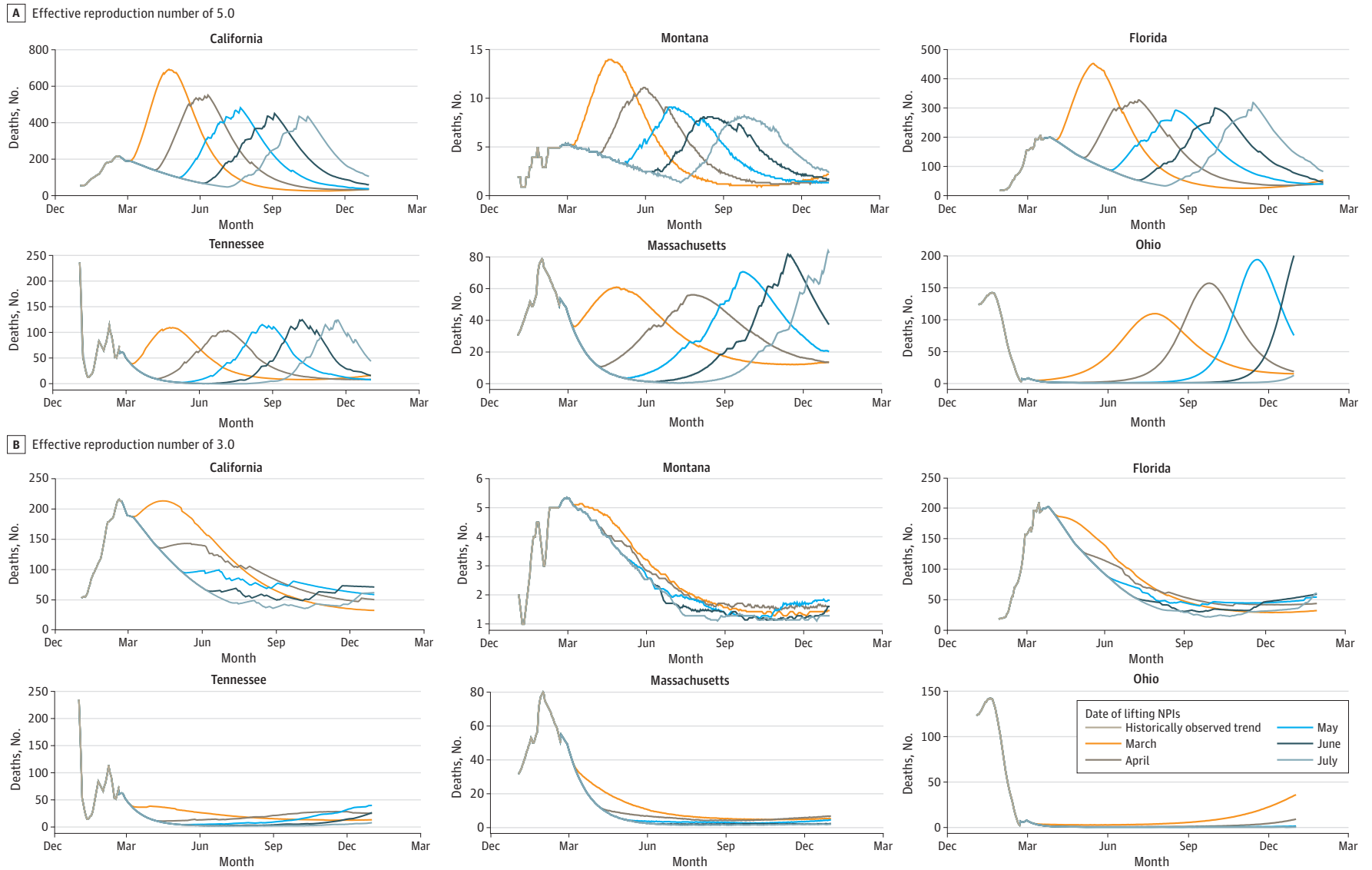
Outcomes and Statistical Analysis

The primary outcome was projections of COVID-19 incident deaths during the remainder of the 2022 calendar year in the 50 US states, the District of Columbia, and Puerto Rico. In addition, we calculated the Spearman rank correlation coefficient as a measure of the correlation between population-level immunity at the time of lifting NPIs and the height of the rebounding surges in COVID-19 deaths. Analyses were performed in February and March 2022 using R, version 4.0.3 (R Foundation for Statistical Computing).

Results

Simulation outcomes indicate that in almost every state, lifting NPIs in 2022 would lead to a substantial rebound in COVID-19 deaths, with peak incident deaths rivaling those seen at the peak of the Omicron surge if lifting occurred in March 2022. Beyond that, however, delaying lifting would not benefit every state equally. In California, Montana, North Carolina, and Oregon, incremental 1-month delays in lifting was estimated to mitigate the amplitude of the rebounding peak in incident deaths. In contrast, the predicted peaks in Florida, Illinois, Michigan, Tennessee, and Washington are similar in size regardless of the timing of lifting, indicating that prolonging restrictions would not meaningfully reduce the disease burden. Moreover, in Massachusetts, New Jersey, New York, and Ohio, delaying lifting was estimated to increase the subsequent peak in incident deaths. Panel A in the **Figure** presents projections for select states in each group of qualitative outcomes. A complete set of state projections can be found in the eAppendix in the [Supplement](#).

Figure. Projected COVID-19 Incident Deaths



Model-based projections of COVID-19 deaths in 2022 following the lifting of nonpharmacologic interventions in California, Montana, Florida, Tennessee, Massachusetts, and Ohio, assuming an effective reproduction number of 5.0 (A) and 3.0 (B).

Heterogeneity in the magnitude of the resurgent epidemic across states was driven by the assumption of a single value of the effective reproduction number when no NPIs were in place. It is plausible that smaller states would never reach the level of transmission implied by an effective reproduction number of 5.0 owing to lower population density and activity; hence, the results may be overestimating the severity of their outlook. Panel B in the Figure presents projections with a lower reproduction number of 3.0, in which case the majority of states could lift restrictions with minimal COVID-19 repercussions.

Heterogeneity in response was also associated with differential levels of immunity from both natural infection and vaccination. The combination of waning immunity and falling rates of infection and vaccination means that the net change in population-level immunity would eventually become negative, such that longer delays in lifting could be associated with larger rebounding epidemics. Therefore, current levels of immunity are crucial determinants of the outcomes of returning to higher levels of transmission. We define *effective immunity* as the sum of the proportion of the populations in the unvaccinated, partially vaccinated, and fully vaccinated susceptible states, weighted by their susceptibility to infection. The Spearman rank correlation coefficient between peak incident deaths (as a percentage of the total population) following the lifting of restrictions on March 1, 2022, and effective immunity on February 20, 2022, was -0.88 ($P < 2.2 \times 10^{-16}$). This highly significant and strongly negative correlation suggests that immunity to infection may be associated with a reduction in the severity of the ensuing epidemic following the lifting of NPIs.

Discussion

We used the COVID-19 Policy Simulator to forecast the number of COVID-19 deaths in each of the 50 US states, the District of Columbia, and Puerto Rico through the 2022 calendar year pending relaxation of NPIs. This analysis could potentially aid state public health officials in evaluating the costs and benefits of lifting NPIs and the timing thereof.

The analysis demonstrates the importance of the timing of lifting NPIs. Premature lifting was estimated to result in recurrent epidemic surges in every almost state. At the same time, a delay of even 1 month was estimated to result in marked reductions to the peak of the mortality curve and the burden on US hospitals. Unfortunately, in most states, no critical moment was identified after which it would be possible to lift NPIs without expecting to see a rebounding surge in deaths. The message that there is no "magic moment" to lift restrictions is important for both sides of the current masking debates in the US. Those opposed to mask mandates should recognize the adverse health outcomes related to relaxing transmission mitigation measures. Any argument to remove such restrictions must address the trade-off and explicitly argue for lifting restrictions within a cost-benefit framework examining the cost of restrictions vs the cost of COVID-19 mortality. At the same time, those who favor maintaining NPIs must recognize that "just a little longer" will not suffice. There is likely no amount of additional waiting time in any state after which removing NPIs will not lead to a rise in morbidity and mortality. The same logic and goals that drive mitigation today will persist, emphasizing the need for mitigation in the future.

A difficult trade-off lies on the horizon. The decision need not be made today, and there is ample evidence that a March 2022 lifting date would have been too soon in many states. However, whenever states do remove NPIs, they will face the same difficult decision regarding the trade-off between increased COVID-19 mortality and the freedoms of returning to a prepandemic norm.

We also estimate that the highly transmissible Delta and Omicron variants will likely continue to take a major toll on the US. The simulations reveal that it is the high transmissibility of these recent variants that sustains the pandemic. With a lower level of transmission similar to that of the ancestral strains, the burden of rebounding morbidity and mortality would be substantially lower. Were this the case, it would likely be possible to remove NPIs at the beginning of the second quarter of 2022.

Limitations

This study has important limitations. First, we use simulation modeling, which includes all caveats that past performance does not ensure future performance. The COVID-19 Policy Simulator closely replicates historical trends in COVID-19 cases and deaths in all states, but it cannot forecast trends introduced by entirely new dynamics, such as new SARS-CoV-2 variants. Second, the true level of transmission following the lifting of NPIs is uncertain but is obviously a key driver of the outcomes of the analysis. We have presented outcomes with a pessimistic and an optimistic value of the effective reproduction number to allow readers to make their own assessments. Third, the model does not incorporate interstate travel. Predictions may be biased in states that typically experience a high level of travel from other states that have differing levels of COVID-19 cases. Fourth, the model assumes that when NPIs are removed, the virus returns to the level of transmissibility expected in the complete absence of mitigation measures. In reality, individuals may voluntarily continue to wear masks and practice social distancing, which could mitigate the severity of rebounding epidemics.

Conclusions

This study used simulation modeling to project COVID-19 deaths in each of the 50 US states, the District of Columbia, and Puerto Rico assuming different timing of the lifting of NPIs. We estimated substantial heterogeneity in outcomes between states, that the timing of lifting NPIs is important, that even short delays in lifting could have a big impact, but that there is likely no amount of delay after which it would be completely safe to remove NPIs. Policy makers should consider the findings of this analysis as they monitor their state's progress during the COVID-19 pandemic, project a suitable time to end restrictions, begin to discuss the conditions that must be met before declaring the pandemic over, and keep the public informed by making public health plans both safe and explicit. Ongoing vaccination efforts will help to contain the COVID-19 pandemic in 2022.

ARTICLE INFORMATION

Open Access: This is an open access article distributed under the terms of the [CC-BY License](#). © 2022 Linas BP et al. *JAMA Health Forum*.

Accepted for Publication: March 8, 2022.

Published: April 1, 2022. doi:[10.1001/jamahealthforum.2022.0760](https://doi.org/10.1001/jamahealthforum.2022.0760)

Corresponding Author: Benjamin P. Linas, MD, MPH, Boston Medical Center, 801 Massachusetts Ave, Crosstown Bldg, Room 2007, Boston, MA 02118 (benjamin.linas@bmc.org).

Author Affiliations: Boston Medical Center, Boston, Massachusetts (Linas); Boston University Schools of Medicine and Public Health, Boston, Massachusetts (Linas); H. Milton Stewart School of Industrial and Systems Engineering, Georgia Institute of Technology, Atlanta (Xiao, Ayer); Value Analytics Labs, Boston, Massachusetts (Dalgic); Massachusetts General Hospital Institute for Technology Assessment, Boston (Mueller, Adee, Aaron, Chhatwal); Harvard Medical School, Boston, Massachusetts (Chhatwal).

Author Contributions: Ms Xiao had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis. Dr Linas and Ms Xiao contributed equally to the analysis and are co-first authors. Drs Ayer and Chhatwal contributed equally to the analysis and are co-senior authors.

Concept and design: Linas, Xiao, Mueller, Ayer, Chhatwal.

Acquisition, analysis, or interpretation of data: All authors.

Drafting of the manuscript: Linas, Xiao, Adee, Ayer, Chhatwal.

Critical revision of the manuscript for important intellectual content: Linas, Xiao, Dalgic, Mueller, Aaron, Ayer, Chhatwal.

Statistical analysis: Linas, Xiao, Mueller, Adee, Ayer, Chhatwal.

Obtained funding: Chhatwal.

Administrative, technical, or material support: Dalgic, Adee, Aaron.

Supervision: Linas, Ayer, Chhatwal.

Conflict of Interest Disclosures: Dr Linas reported receiving grants from the National Institute on Drug Abuse, the US Centers for Disease Control and Prevention, and the National Science Foundation during the conduct of the study. Mr Dalgic reported that Value Analytics Labs provided consulting services for Janssen Pharmaceuticals outside the submitted work. Dr Ayer reported being a managing partner at Value Analytics Labs, a health care analytics firm, outside the submitted work. Dr Chhatwal reported receiving grants from the National Science Foundation and Rockefeller Foundation during the conduct of the study; and receiving personal fees from Value Analytics Labs outside the submitted work. No other disclosures were reported.

Funding/Support: This work was supported by National Science Foundation awards 2035360 and 2035361, and the Rockefeller Foundation COVID-19 Modeling Accelerator.

Role of the Funder/Sponsor: The funders had no role in the design and conduct of the study; collection, management, analysis, and interpretation of the data; preparation, review, or approval of the manuscript; and decision to submit the manuscript for publication.

Additional Information: Statistical code is available from the authors on request to the corresponding author.

REFERENCES

1. Chhatwal J, Dalgic O, Mueller P, et al. PIN68 COVID-19 simulator: an interactive tool to inform COVID-19 intervention policy decisions in the United States. *Value Health*. 2020;23:S556. doi:10.1016/j.jval.2020.08.909
2. US Census Bureau. SC-EST2020-AGESEX-CIV: annual estimates of the civilian population by single year of age and sex for the United States, states, and the District of Columbia: April 1, 2010 to July 1, 2020. Accessed March 21, 2022. <https://www2.census.gov/programs-surveys/popest/technical-documentation/file-layouts/2010-2020/sc-est2020-agesex-civ.pdf>
3. Prem K, Cook AR, Jit M. Projecting social contact matrices in 152 countries using contact surveys and demographic data. *PLoS Comput Biol*. 2017;13(9):e1005697. doi:10.1371/journal.pcbi.1005697
4. Xin H, Li Y, Wu P, et al. Estimating the latent period of coronavirus disease 2019 (COVID-19). *Clin Infect Dis*. Published online August 28, 2021. doi:10.1093/cid/ciab746
5. Byrne AW, McEvoy D, Collins AB, et al. Inferred duration of infectious period of SARS-CoV-2: rapid scoping review and analysis of available evidence for asymptomatic and symptomatic COVID-19 cases. *BMJ Open*. 2020;10(8):e039856. doi:10.1136/bmjopen-2020-039856
6. Townsend JP, Hassler HB, Wang Z, et al. The durability of immunity against reinfection by SARS-CoV-2: a comparative evolutionary study. *Lancet Microbe*. 2021;2(12):e666-e675. doi:10.1016/S2666-5247(21)00219-6
7. Liu Y, Rocklöv J. The reproductive number of the Delta variant of SARS-CoV-2 is far higher compared to the ancestral SARS-CoV-2 virus. *J Travel Med*. 2021;28(7):taab124. doi:10.1093/jtm/taab124
8. Bevand M. Covid19-Age-Stratified-Ifr. Accessed March 1, 2022. <https://github.com/mbevand/covid19-age-stratified-ifr>
9. Centers for Disease Control and Prevention. Estimated COVID-19 burden. Accessed March 1, 2022. <https://www.cdc.gov/coronavirus/2019-ncov/cases-updates/burden.html>
10. Dagan N, Barda N, Kepten E, et al. BNT162b2 mRNA COVID-19 vaccine in a nationwide mass vaccination setting. *N Engl J Med*. 2021;384(15):1412-1423. doi:10.1056/NEJMoa2101765
11. Mathieu E, Ritchie H, Ortiz-Ospina E, et al. A global database of COVID-19 vaccinations. *Nat Hum Behav*. 2021;5(7):947-953. doi:10.1038/s41562-021-01122-8
12. Crotty S. Hybrid immunity. *Science*. 2021;372(6549):1392-1393. doi:10.1126/science.abj2258

SUPPLEMENT.

eAppendix
eReferences

Supplemental Online Content

Linas BP, Xiao J, Dalgic OO, et al. Projecting COVID-19 mortality as states relax nonpharmacologic interventions. *JAMA Health Forum*. 2022;3(4):e220760.
doi:10.1001/jamahealthforum.2022.0760

eAppendix

eReferences

This supplemental material has been provided by the authors to give readers additional information about their work.

eAppendix.

A1. Introduction

The *COVID-19 Policy Simulator* uses a mathematical model to simulate the COVID-19 pandemic at the national and state levels in the United States. The model is calibrated to historical trends in daily incident deaths and updated weekly as new data arises and the pandemic situation evolves. The online tool allows users to simulate the disease trajectory under different non-pharmaceutical interventions (NPIs) with varying timing and intensity. Forecasts are made for total cases, diagnosed cases, active cases, deaths, hospital bed occupancy, and ICU bed occupancy. Since May 2020, we have contributed weekly to the Centers for Disease Control and Prevention (CDC) COVID-19 Forecast Hub.¹

A2. Development Timeline

eTable 1 lists major model updates and the dates on which they were introduced.

eTable 1. Major model updates.

Date	Update
February 2021	Vaccine rollout
August 2021	Age-stratification to incorporate age-stratified vaccine data and differential mortality of age groups
October 2021	Lower vaccine effectiveness due to Delta variant from August 1, 2021
December 2021	Waning (natural and vaccine-conferred) immunity
January 2022	Lower vaccine effectiveness due to Omicron variant from December 1, 2021

A3. Model Overview

Our model is an extension of the traditional susceptible-infected-recovered (SIR) model,² which partitions a population into compartments representing mutually exclusive disease states. At any time t , the variables $S(t)$, $E(t)$, $I(t)$, $R(t)$, and $D(t)$ denote the number of people in the susceptible, exposed, infected, recovered, and deceased compartments respectively. The flow of people between compartments is assumed to obey a system of deterministic ordinary differential equations. We let $\Delta t = 1$ to be compatible with data sources reporting daily data.

Age stratification

We stratify the population into two age groups, <65 years (low-risk) and ≥ 65 years (high-risk), with the subscript $a \in \{L, H\}$. The total population in age group a , denoted by N_a , is assumed to be constant over the simulation period.

Vaccination

To reflect administration guidelines of the Pfizer-BioNTech and Moderna vaccines,ⁱ we stratify the disease states by vaccination status. The subscript $v \in \{0, 1, 2\}$ denote the number of vaccine doses received under the recommended two-dose regime. The third vaccine, Janssen, approved for a single-dose regime, is omitted from the model due to its accounting for only 3.7% of all administered doses in the U.S. as of October 31, 2021.ⁱⁱ Since there is no data on vaccination status at the time of infection, we assume doses are allocated proportionally to the susceptible and recovered compartments over the

historical time horizon,ⁱⁱⁱ i.e., if $V_{a,1}(t)$ and $V_{a,2}(t)$ are the actual number of first and second doses administered to age group a on day t , the proportion of the v -dose susceptible and recovered compartments, $v \in \{0, 1\}$, moving into the corresponding $(v + 1)$ -dose compartments on day t is

$$\alpha_{a,v+1}(t) = \min \left\{ 1, \frac{V_{a,v+1}(t - 12)}{S_{a,v}(t) + R_{a,v}(t)} \right\}.$$

The time lag of 12 days accounts for the delay between receiving a vaccine dose and the beginning of protection.³ The implicit assumption is that a susceptible person does not become infected in the 12 days after receiving a dose. The vaccine reduces both susceptibility to infection and mortality risk. After v vaccine doses, the probability of contracting the virus is reduced by $100 \times e_v^I\%$, with $0 = e_0^I \leq e_1^I, e_2^I \leq 1$; similarly, the infection fatality rate is reduced by $100 \times e_v^D\%$, with $0 = e_0^D \leq e_1^D, e_2^D \leq 1$.

Transmission

For a susceptible individual in age group a who has received v vaccine doses, the rate of exposure to the virus is given by

$$\lambda_{a,v} = (1 - e_v^I) \mathcal{R}(t) \gamma \sum_{a' \in \{L, H\}} c_{a,a'} \frac{I_{a',0}(t) + I_{a',1}(t) + I_{a',2}(t)}{N_{a'}},$$

where $\mathcal{R}(t)$ is the time-varying effective reproduction number. We model $\mathcal{R}(t)$ as a step function with breakpoints at the beginning of each calendar month over the historical time horizon to capture the effect of NPIs enforced during this period. The coefficients $c_{a,a'}$ are the elements of the contact matrix with row sums normalized to 1, so that $c_{a,a'}$ is the proportion of contacts per day of age group a that are with age group a' . When a susceptible individual contracts the virus, they enter the exposed state and remain there for the duration of the latent period with a mean of $1/\kappa$ days. After that, they transition to the infected state and remain there for the duration of the infectious period with a mean of $1/\gamma$ days. Finally, the infected individual will either die with probability $\delta_{a,v} = (1 - e_v^D)\delta_a$, where δ_a is the baseline infection fatality rate for age group a , or recover with probability $(1 - \delta_{a,v})$.

Waning immunity

An individual who has recovered from natural infection ($R_{a,v}$) enjoys a period of natural immunity with a mean of $1/\omega_n$ days before transitioning back into the susceptible state ($S_{a,v}$). A fully vaccinated susceptible individual ($S_{a,2}$) is protected for the duration of vaccine-conferred immunity with a mean of $1/w_v$ days before transitioning back into the partially susceptible state ($S_{a,1}$). Finally, since individuals with natural immunity who are subsequently vaccinated have been reported to exhibit “unusually potent immune responses”,⁴ a fully vaccinated recovered individual ($R_{a,2}$) is assumed to possess two ‘layers’ of immunity, shedding first their natural immunity then their vaccine-conferred immunity.

Booster shots

It is assumed that, once vaccinated, an individual will never shed their immunity completely (within the time frame of the simulation), and a fully vaccinated individual who has shed their vaccine-conferred

immunity is indistinguishable from a partially vaccinated individual. Thus, the model differentiates between the subpopulation that is willing to receive booster shots and the subpopulation that is unwilling to be vaccinated. Fully vaccinated individuals wane into the partially vaccinated state and are ‘boosted’ back into the fully vaccinated state.

Differential equation formulation

In summary, our model is described by the following system of equations, where (t) has been dropped for notational simplicity:

$$\begin{aligned}
 \dot{S}_{a,0} &= -(\lambda_{a,0} + \alpha_{a,1})S_{a,0} + \omega R_{a,0}, \\
 \dot{E}_{a,0} &= \lambda_{a,0}S_{a,0} - \kappa E_{a,0}, \\
 \dot{I}_{a,0} &= \kappa E_{a,0} - \gamma I_{a,0}, \\
 \dot{R}_{a,0} &= (1 - \delta_{a,0})\gamma I_{a,0} - (\alpha_{a,1} + \omega)R_{a,0}, \\
 \\
 \dot{S}_{a,1} &= -(\lambda_{a,1} + \alpha_{a,2} + \alpha_{a,3})S_{a,1} + \alpha_{a,1}S_{a,0} + \omega(R_{a,1} + \alpha'_{a,3}S_{a,2}), \\
 \dot{E}_{a,1} &= \lambda_{a,1}S_{a,1} - \kappa E_{a,1}, \\
 \dot{I}_{a,1} &= \kappa E_{a,1} - \gamma I_{a,1}, \\
 \dot{R}_{a,1} &= (1 - \delta_{a,1})\gamma I_{a,1} + \alpha_{a,1}R_{a,0} - (\alpha_{a,2} + \alpha_{a,3} + \omega)R_{a,1}, \\
 \\
 \dot{S}_{a,2} &= -\lambda_{a,2}S_{a,2} + (\alpha_{a,2} + \alpha_{a,3})S_{a,1} + \omega\alpha'_{a,3}(R_{a,2} - S_{a,2}), \\
 \dot{E}_{a,2} &= \lambda_{a,2}S_{a,2} - \kappa E_{a,2}, \\
 \dot{I}_{a,2} &= \kappa E_{a,2} - \gamma I_{a,2}, \\
 \dot{R}_{a,2} &= (1 - \delta_{a,2})\gamma I_{a,2} + (\alpha_{a,2} + \alpha_{a,3})R_{a,1} - \omega\alpha'_{a,3}R_{a,2}, \\
 \\
 \dot{D} &= \delta_{a,0}\gamma I_{a,0} + \delta_{a,1}\gamma I_{a,1} + \delta_{a,2}\gamma I_{a,2}.
 \end{aligned}$$

The initial conditions are $(I_{a,0}(0), S_{a,0}(0)) = (I_{a,0}^{\text{init}}, N_a - I_{a,0}^{\text{init}})$ and zero for all other variables. eTables 2 and 3 display the values or ranges for all model parameters and their references.

eTable 2. Estimates of fixed and calibrated parameters.

Parameter	Estimate	Reference and notes
<i>Fixed parameters</i>		
N_L	State-dependent	U.S. Census Bureau: SC-EST2020-AGESEX-CIV: POPEST2019_CIV
N_H		
$[c_{L,L}, c_{L,H}; c_{H,L}, c_{H,H}]$	[0.93, 0.07; 0.48, 0.52]	⁵ Aggregate columns and rows into age groups <65 years and ≥65 years, then normalize so that rows sum to 1.
κ	1/5.5	⁶
γ	1/10	⁷
ω_n, ω_v	16 months	⁸

		Reinfection by endemic SARS-CoV-2 is expected to occur between 3 months and 5 years after peak antibody response, with a median of 16 months.
Calibrated parameters		
$\mathcal{R}(t)$	0.5–6.0	9, 10, 11, 12 Widely varying by location and SARS-CoV-2 variant.
I^{init}	100–10,000	Divided proportionally into: <ul style="list-style-type: none"> $I_L^{\text{init}} = \frac{N_L}{N_L + N_H} I^{\text{init}}$, $I_H^{\text{init}} = \frac{N_H}{N_L + N_H} I^{\text{init}}$.

eTable 3. Evolution of estimates of variant-dependent parameters.

Parameter	Estimate	Reference and notes
Baseline values		
δ_L	0.001	Based on this meta-analysis . These values chosen to approximate the CDC 's "Estimated Total Infections".
δ_H	0.030	
$1 - e_1^I$	0.54	¹³ Table 2: "Documented Infection" at "14 to 20 days after first dose": $(1 - \text{RR})\% = 46$. Derivation: $1 - e_1^I = 1 - 0.46$.
$1 - e_2^I$	0.08	¹³ Table 2: "Documented Infection" at "7 days after second dose to end of follow-up": $(1 - \text{RR})\% = 92$. Derivation: $1 - e_2^I = 1 - 0.92$.
$1 - e_1^D$	0.52	¹³ Table 2: "Death" at "14 to 20 days after first dose": $(1 - \text{RR})\% = 72$. Derivation: $(1 - e_1^I)(1 - e_1^D) = 1 - 0.72$.
$1 - e_2^D$	0.63	Vaccine clinical trials report 95% efficacy against death. Derivation: $(1 - e_2^I)(1 - e_2^D) = 1 - 0.95$. Note that it is the <i>conditional probability</i> of death that is higher after the second dose than after the first dose. If a fully vaccinated individual contracts a breakthrough infection despite 92% reduction in susceptibility, it is plausible that they are particularly vulnerable and have a smaller reduction in mortality risk <i>conditional on infection</i> compared to a partially vaccinated individual who contracts a breakthrough infection.
Delta variant (parameter values change at a linear rate as the variant saturates over August 2021)		
$1 - e_2^I$	0.13	¹⁴ "Two dose vaccine effectiveness was 86.7% (95% confidence interval 84.3% to 88.7%) against infection with the delta variant, ..." Derivation: $(1 - e_2^I) = 1 - 0.87$.
$1 - e_1^I$	0.90	Scaled up proportionally to e_2^I . Derivation: $1 - e_1^I = 0.54 \times (0.13/0.08)$.
δ_L	0.0023	¹⁵ "Increased risk with the Delta variant was more pronounced at ... 133% (95% CI 54%–231%) for death." Derivation: $\delta_L = 0.001 \times 2.33$.
δ_H	0.0700	Derivation: $\delta_H = 0.030 \times 2.33$.

		We assume that the increased mortality of Delta is a consequence of higher IFR only, not lower vaccine effectiveness against death after infection.
<i>Omicron variant (parameter values change at a linear rate as the variant saturates over December 2021)</i>		
$1 - e_2^I$	0.30	UKHSA Report : Figure 2B
δ_L	0.0008	UKHSA Report: “The risk of hospital admission from emergency departments with Omicron was approximately one-third of that for Delta (Hazard Ratio 0.33, 95% CI: 0.30 to 0.37).”
δ_H	0.0233	Same as above.

A4. Calibration and Numerical Solution

We calibrate the model to historical daily incident deaths. The system of ordinary differential equations is solved numerically using Euler’s method (R package `deSolve`).¹⁶ The calibration method is generalized simulated annealing (R package `GenSA`) with the sum of squared errors as the objective function.¹⁷ To account for uncertainty in the calibrated values, we repeat the calibration process 100 times with different initial solutions, resulting in 100 unique sets of parameter values and fitted curves. At each time point, we take the median as the point estimate and compute the 90% coverage simulation band.

eTable 4 displays our input data and their references.

eTable 4. Input data.

Data	Source	Reference
COVID-19 cases and deaths	JHU CSSE	¹⁸
Vaccine administration	Our World in Data and CDC	¹⁹ and CDC

A5. Forecasting

We make forecasts by allowing the model to continue running past the historical time horizon. Diagnosed cases and hospital and ICU occupancy are not accounted for in the SEIR model. We estimate these in a post-processing step as follows.

Diagnosed cases

We assume the future diagnosis rate remains at the latest estimated value, i.e., the number of incident diagnosed cases on the last day of data divided by the number of incident total cases on the last day of data.

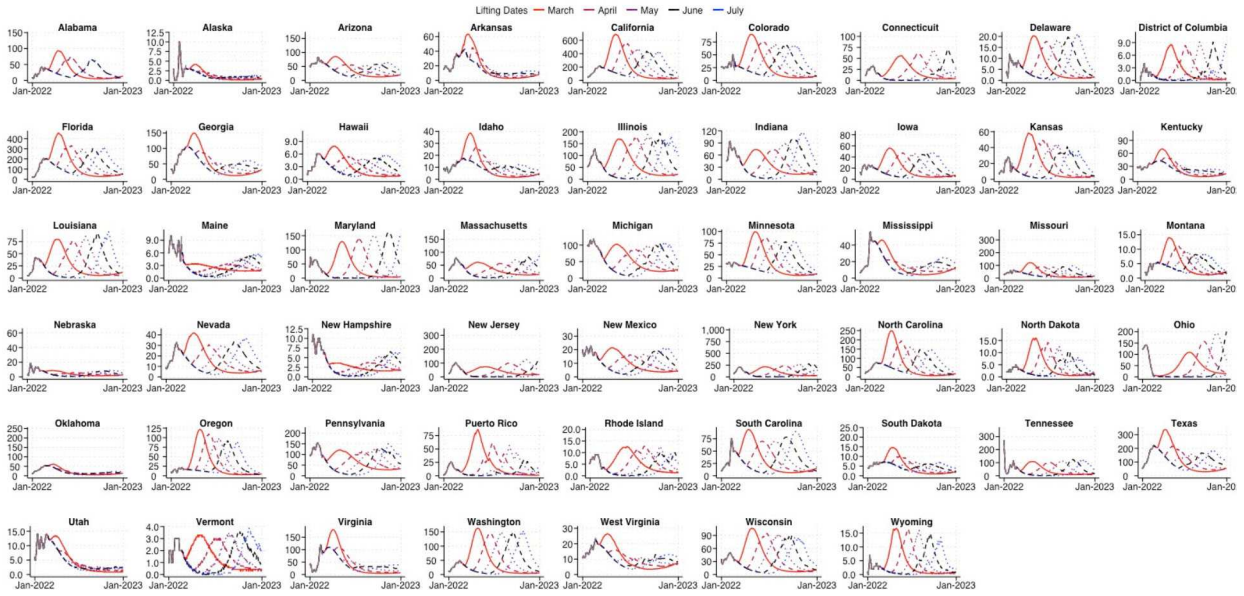
Hospital and ICU bed occupancy

We back-calculate hospital and ICU bed occupancy from incident deaths assuming an average time to death from hospital and ICU admission of 16 and 10 days respectively.²⁰ Starting in August, we forecast occupancy data provided by the [U.S. Department of Health and Human Services](#). Note that the data does not include all hospitals in any given state so our forecasts do not estimate the total demand for hospital and ICU beds.

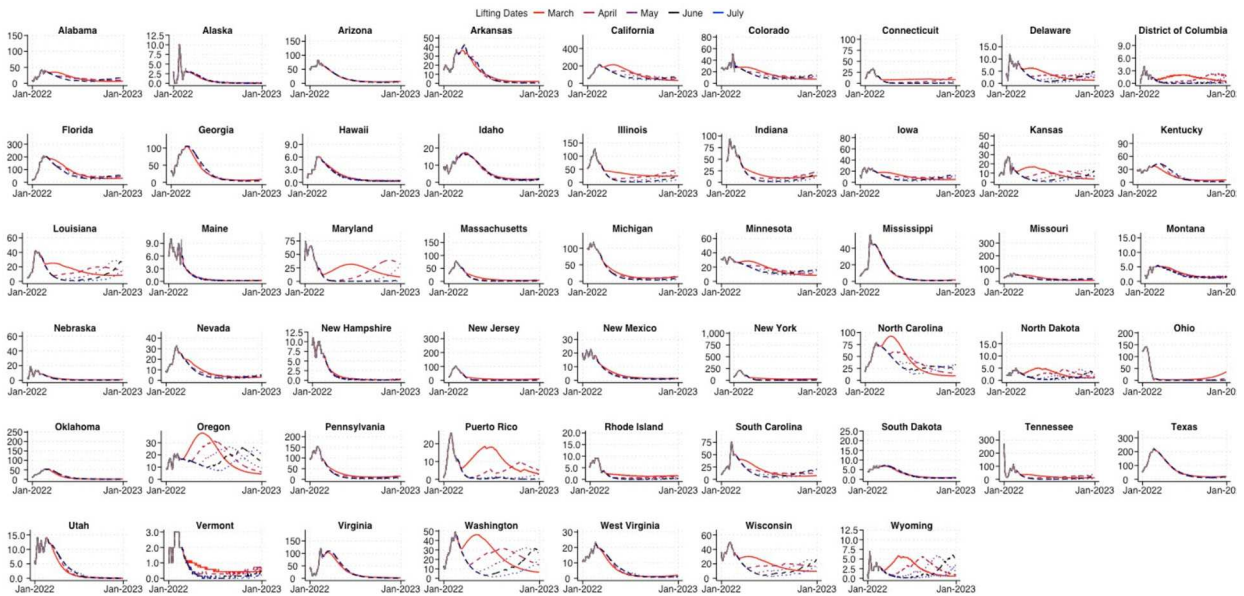
A6. Non-Pharmaceutical Interventions

The *Simulator* offers four types of non-pharmaceutical interventions:

1. *Current interventions*: The effective reproduction number remains at the latest calibrated value.
2. *Lockdown*: The effective reproduction number is 0.3, the estimated value in Wuhan during the strict lockdown of the city starting in March 2020.²¹
3. *Stay-at-home orders*: The effective reproduction number is the lowest calibrated value attained during the period from March to July 2020.
4. *Minimal restrictions*: The effective reproduction number is the basic reproduction number, which changes with the proportions of the circulating variants of concern.



(a)



(b)

Figure 1. Model-based projections of COVID-19 deaths following the lifting of NPIs in each state, assuming an effective reproduction number of 5.0 (eFigure 1a) and 3.0 (eFigure 1b).

References

1. Cramer, E. Y. *et al.* Evaluation of individual and ensemble probabilistic forecasts of COVID-19 mortality in the US. *medRxiv* 2021.02.03.21250974 (2021) doi:10.1101/2021.02.03.21250974.
2. Kermack, W. O., McKendrick, A. G. & Walker, G. T. A contribution to the mathematical theory of epidemics. *Proc. R. Soc. Lond. Ser. Contain. Pap. Math. Phys. Character* **115**, 700–721 (1927).
3. Mahase, E. Covid-19: Pfizer vaccine efficacy was 52% after first dose and 95% after second dose, paper shows. *BMJ* **371**, m4826 (2020).
4. Crotty, S. Hybrid immunity. *Science* **372**, 1392–1393 (2021).
5. Prem, K., Cook, A. R. & Jit, M. Projecting social contact matrices in 152 countries using contact surveys and demographic data. *PLOS Comput. Biol.* **13**, e1005697 (2017).
6. Xin, H. *et al.* Estimating the Latent Period of Coronavirus Disease 2019 (COVID-19). *Clin. Infect. Dis.* (2021) doi:10.1093/cid/ciab746.
7. Byrne, A. W. *et al.* Inferred duration of infectious period of SARS-CoV-2: rapid scoping review and analysis of available evidence for asymptomatic and symptomatic COVID-19 cases. *BMJ Open* **10**, e039856 (2020).
8. Townsend, J. P. *et al.* The durability of immunity against reinfection by SARS-CoV-2: a comparative evolutionary study. *Lancet Microbe* (2021) doi:10.1016/S2666-5247(21)00219-6.
9. Hilton, J. & Keeling, M. J. Estimation of country-level basic reproductive ratios for novel Coronavirus (SARS-CoV-2/COVID-19) using synthetic contact matrices. *PLOS Comput. Biol.* **16**, e1008031 (2020).
10. Billah, M. A., Miah, M. M. & Khan, M. N. Reproductive number of coronavirus: A systematic review and meta-analysis based on global level evidence. *PLOS ONE* **15**, e0242128 (2020).
11. Ke, R., Romero-Severson, E., Sanche, S. & Hengartner, N. Estimating the reproductive number R_0 of SARS-CoV-2 in the United States and eight European countries and implications for vaccination. *J. Theor. Biol.* **517**, 110621 (2021).
12. Liu, Y. & Rocklöv, J. The reproductive number of the Delta variant of SARS-CoV-2 is far higher compared to the ancestral SARS-CoV-2 virus. *J. Travel Med.* **28**, (2021).
13. Dagan, N. *et al.* BNT162b2 mRNA Covid-19 Vaccine in a Nationwide Mass Vaccination Setting. *N. Engl. J. Med.* **384**, 1412–1423 (2021).
14. Bruxvoort, K. J. *et al.* Effectiveness of mRNA-1273 against delta, mu, and other emerging variants of SARS-CoV-2: test negative case-control study. *BMJ* **375**, e068848 (2021).
15. Fisman, D. N. & Tuite, A. R. Evaluation of the relative virulence of novel SARS-CoV-2 variants: a retrospective cohort study in Ontario, Canada. *CMAJ Can. Med. Assoc. J. J. Assoc. Medicale Can.* **193**, E1619–E1625 (2021).
16. Soetaert, K., Petzoldt, T. & Setzer, R. W. Package deSolve: Solving Initial Value Differential Equations in R. 52.
17. Xiang, Y., Gubian, S., Suomela, B. & Hoeng, J. Generalized Simulated Annealing for Global Optimization: The GenSA Package. *R J.* **5**, 13 (2013).
18. Dong, E., Du, H. & Gardner, L. An interactive web-based dashboard to track COVID-19 in real time. *Lancet Infect. Dis.* **20**, 533–534 (2020).
19. Mathieu, E. *et al.* A global database of COVID-19 vaccinations. *Nat. Hum. Behav.* 1–7 (2021) doi:10.1038/s41562-021-01122-8.

20. Ferguson, N. *et al.* *Report 9: Impact of non-pharmaceutical interventions (NPIs) to reduce COVID19 mortality and healthcare demand*. 20 <http://spiral.imperial.ac.uk/handle/10044/1/77482> (2020) doi:10.25561/77482.
21. Pan, A. *et al.* Association of Public Health Interventions With the Epidemiology of the COVID-19 Outbreak in Wuhan, China. *JAMA* **323**, 1915–1923 (2020).

ⁱ See <https://www.cdc.gov/coronavirus/2019-ncov/vaccines/second-shot.html> for details.

ⁱⁱ See <https://ourworldindata.org/grapher/covid-vaccine-doses-by-manufacturer?country=~USA> for details.

ⁱⁱⁱ The CDC advises against vaccination while under active infection. See <https://www.cdc.gov/coronavirus/2019-ncov/vaccines/faq.html> for details.