About the Author

Dr. Schulz is a postdoctoral researcher at University of Veterinary Medicine Hanover, Hanover, Germany. Her primary research interests are the pathogenesis and epidemiology of emerging and vector-borne diseases in the wildlife–livestock and human–animal interfaces.

References

1. Gaudreault NN, Trujillo JD, Carossino M, Meekins DA, Morozov I, Madden DW, et al. SARS-CoV-2 infection, disease and transmission in domestic cats. Emerg Microbes Infect. 2020;9:2322–32. https://doi.org/10.1080/22221751.2020.1833687
2. Halfmann PJ, Hatta M, Chiba S, Masmura T, Fan S, Takeda M, et al. Transmission of SARS-CoV-2 in domestic cats. N Engl J Med. 2020;383:592–4. https://doi.org/10.1056/NEJMc2013400
3. Barrs VR, Peiris M, Tam KW, Law PY, Brackman CJ, To EM, et al. SARS-CoV-2 in quarantined domestic cats from COVID-19 households or close contacts, Hong Kong, China. Emerg Infect Dis. 2020;26:3071–4. https://doi.org/10.3201/eid2612.200786
4. Sit TH, Brackman CJ, Ip SM, Tam KW, Law PY, To EM, et al. Infection of dogs with SARS-CoV-2. Nature. 2020;586:776–8. https://doi.org/10.1038/s41586-020-2334-5
5. Wernike K, Aebischer A, Michelitsch A, Hoffmann D, Freuling C, Balkema-Buschmann A, et al. Multi-species ELISA for the detection of antibodies against SARS-CoV-2 in animals. Transbound Emerg Dis. 2020;Nov 15:1–7. https://doi.org/10.1111/tbed.13926
6. Garigliany M, Van Laere AS, Clercx C, Giet D, Escrion N, Huon C, et al. SARS-CoV-2 natural transmission from human to cat, Belgium, March 2020. Emerg Infect Dis. 2020;26:3069–71. https://doi.org/10.3201/eid2612.200223
7. Yousaf M, Hameed M, Alsoub H, Jamal W, Ahmad M. COVID-19: Prolonged viral shedding in an HIV patient with literature review of risk factors for prolonged viral shedding and its implications for isolation strategies. Clin Case Rep. 2021;9:1397–401. https://doi.org/10.1002/ccr3.3786
8. Ambrosioni J, Blanco JL, Reyes-Urueña JM, Davies MA, Sued O, Marcos MA, et al.; COVID-19 in HIV Investigators. Overview of SARS-CoV-2 infection in adults living with HIV. Lancet HIV. 2021;8:e294–305. https://doi.org/10.1016/S2352-3018(21)00070-9
9. Tarhini H, Reoing A, Bridier-Nahmias A, Rahi M, Lambert C, Martres P, et al. Long term SARS-CoV-2 infectiousness among three immunocompromised patients: from prolonged viral shedding to SARS-CoV-2 superinfection. J Infect Dis. 2021;Feb 8:jia075. https://doi.org/10.1093/infdis/jia075
10. Sailléeau C, Dumarest M, Vanhomwegen J, Delaplace M, Caro V, Kwasiborski A, et al. First detection and genome sequencing of SARS-CoV-2 in an infected cat in France. Transbound Emerg Dis. 2020;67:2324–8. https://doi.org/10.1111/tbed.13659

Address for correspondence: Asisa Volz, Institute of Virology, University of Veterinary Medicine Hanover, Buenteweg 17, 30559 Hanover, Lower Saxony, Germany; email: asisa.volz@tiho-hannover.de

Effects of COVID-19 Vaccination Timing and Risk Prioritization on Mortality Rates, United States

Xutong Wang, Zhanwei Du, Kaitlyn E. Johnson, Remy F. Pasco, Spencer J. Fox, Michael Lachmann, Jason S. McLellan, Lauren Ancel Meyers

Author affiliations: The University of Texas at Austin, Austin, Texas, USA (X. Wang, Z. Du, K.E. Johnson, R.F. Pasco, S.J. Fox, J.S. McLellan, L.A. Meyers); The University of Hong Kong, Hong Kong, China (Z. Du); Hong Kong Science and Technology Park, Hong Kong (Z. Du); Santa Fe Institute, Santa Fe, New Mexico, USA (M. Lachmann, L.A. Meyers)

DOI: https://doi.org/10.3201/eid2707.210118

During rollout of coronavirus disease vaccination, policymakers have faced critical trade-offs. Using a mathematical model of transmission, we found that timing of vaccination rollout would be expected to have a substantially greater effect on mortality rate than risk-based prioritization and uptake and that prioritizing first doses over second doses may be lifesaving.

In December 2020, the US government issued emergency use authorization for two 2-dose severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) vaccines, both estimated to be >94% efficacious in preventing symptomatic coronavirus disease (COVID-19) (1–3). The Advisory Committee on Immunization Practices immediately recommended the prioritization of frontline workers and high-risk subgroups (4). As of February 14, 2021, ≈52 million doses have been administered (5). We used a mathematical model of COVID-19 transmission to evaluate the effects of vaccine timing, risk prioritization, number of doses administered, and uptake rates on population-level mortality rates (Figure).

Focusing on Austin, Texas, USA, we projected COVID-19 deaths over 8 months for both an infection-blocking vaccine that prevents infection upon exposure (assuming 95% reduction in susceptibility in vaccinated persons) and a symptom-blocking vaccine that prevents symptoms upon infection (assuming 95% reduction in symptomatic ratio in vaccinated persons). Vaccination would begin on January 15 or February 15, with 10,000 vaccines administered weekly and allocated to cities pro rata. We compare 3 strategies: no priority groups; 1 of 3 priority groups vaccinated before the general public (adults >65 years of age, adults who have high-risk underlying
conditions, or both); and 10 phases that vaccinate age–risk groups in order of risk for severe COVID-19 outcomes. Stochastic simulations assumed that 7.6% of the overall population of the Austin–Round Rock Metropolitan Statistical Area were immunized by infection before January 15.

If a perfectly risk-prioritized (10-phase) rollout of an infection-blocking vaccine were to begin January 15, we estimated that 52% (95% CI 47%–56%) of deaths would be averted relative to the baseline of no vaccines, assuming 50% uptake, or 56% (95% CI 51%–60%) of deaths averted assuming 90% uptake (Figure, panel A). If rollout were delayed 1 month, 34% (95% CI 28%–40%) of deaths would be averted at 50% uptake, or 38% (95% CI 32%–43%) at 90% uptake. Under low (50%) uptake, prioritization has minimal benefit. Under high uptake (90%), the 10-stage strategy is optimal, followed by prioritizing adults >65 years of age and high-risk younger adults.

Expected differences are magnified with a symptom-blocking vaccine. For a January 15 start and 50% uptake, the risk-prioritized 10-phase strategy would avert 40% (95% CI 35%–45%) of deaths, whereas unprioritized rollout would avert 32% (95% CI 25%–37%). If a single dose with 82% efficacy (1,2) is administered under the 10-phase strategy, we would
expect a 50% (95% CI 45%–54%) reduction in mortality for a symptom-blocking vaccine and 66% (95% CI 63%–70%) reduction for an infection-blocking vaccine (Appendix Table 1, https://wwwnc.cdc.gov/EID/article/27/7/21-0118-App1.pdf).

These projections validate the prioritizing of high-risk groups. In a pessimistic scenario in which a symptom-blocking vaccine rollout began in February 2021 with 50% uptake, prioritizing high-risk adults and adults ≥65 would avert ≈17,000 (95% CI 10–36,000) more deaths in the United States than a nonprioritized campaign. Given the state of the pandemic in early 2021, we expected vaccine delays to cost more lives than either imperfect prioritization or vaccine hesitancy.

The United Kingdom and Belgium have prioritized first doses over second doses (7), in an effort to provide partial immunity to more persons. The United States has publicly resisted this approach, citing the lack of clinical trial data validating the approach (8). We found that providing a single (82% efficacious) dose would be expected to save more lives than the corresponding 2-dose strategy, because partially immunizing a large number confers a greater degree of population-level protection than more fully immunizing half as many. Although a 1-dose campaign may accelerate herd immunity and require far fewer resources than a 2-dose campaign, we strongly caution that additional data and single-dose trials are needed to establish efficacy. If the single-dose efficacy is <82%, then we would expect the difference between a single-dose strategy and the corresponding 2-dose strategy to be smaller. We expect similar reductions in mortality rate from both strategies when the single-dose efficacy is 52% (2) (Appendix Figure 7). We note that low-efficacy vaccines may increase the risk for vaccine-resistant variants (9) and that there may be political, commercial, and societal barriers to shifting priorities mid-campaign (10).

We assumed that vaccines provide lasting immunity and block either infection or symptoms, whereas the reality may be a hybrid of both (Appendix Table 2, Figure 2), along with riskier behavior stemming from pandemic weariness or overconfidence in the vaccination campaign. Our estimates reflect conditions in the United States in early 2021, as cases were surging toward a pandemic peak in the absence of effective mitigation. The estimated public health benefits of vaccines decrease under higher COVID-19 transmission rates that might occur with relaxed mitigation measures, lower levels of immunity before the rollout, or the emergence of more transmissible SARS-CoV-2 variants including B.1.1.7 (Appendix).

Risk prioritization is a valid approach for maximizing the impact of vaccines, but not at the expense of vaccination speed. Our projections suggest 2 immediate strategies: hybrid distributions that combine active outreach to priority groups with passive distribution to the general public; and distribution of single doses to as much of the population as possible, foregoing plans to hold second doses in reserve.

Acknowledgments
We thank Matthew Biggerstaff for critical discussions and parameter guidance.

This research was supported by grants from the US National Institutes of Health (grant no. R01 AI151176) and the US Centers for Disease Control and Prevention (grant no. U01 IP001136) and a donation from Love, Tito’s (the philanthropic arm of Tito’s Homemade Vodka, Austin, TX, USA) to the University of Texas to support the modeling of COVID-19 mitigation strategies.

About the Author
Dr. Wang completed this work as a PhD candidate at the University of Texas at Austin, under the supervision of Lauren Ancel Meyers. Her research interest is on mathematical and statistical modeling of infectious disease dynamics.

References
1. Vaccines and Related Biological Products Advisory Committee. FDA briefing document: Moderna COVID-19 vaccine. December 17, 2020 [cited 2021 May 7]. https://www.fda.gov/media/144434/download
2. Vaccines and Related Biological Products Advisory Committee. FDA briefing document: Pfizer BioNTech COVID-19 vaccine. December 10, 2020 [cited 2021 May 7]. https://www.fda.gov/media/144245/download
3. US Food and Drug Administration. COVID-19 vaccines. [cited 2021 Feb 17]. https://www.fda.gov/emergency-preparedness-and-response/coronavirus-disease-2019-covid-19/covid-19-vaccines
4. Dooling K, McClung N, Chamberland M, Marin M, Wallace M, Bell BP, et al. The Advisory Committee on Immunization Practices’ interim recommendation for allocating initial supplies of COVID-19 vaccine—United States, 2020. MMWR Morb Mortal Wkly Rep. 2020;69:1857–9. https://doi.org/10.15585/mmwr.mm6949e1
5. US Centers for Disease Control and Prevention. Vaccines for COVID-19. 2021 [cited 2021 Feb 15]. https://www.cdc.gov/coronavirus/2019-ncov/vaccines/index.html
6. Tyson A, Johnson C, Funk C. US public now divided over whether to get COVID-19 vaccine. 2020 [cited 2020 Dec 14]. https://www.pewresearch.org/science/2020/09/17/u-s-public-now-divided-over-whether-to-get-covid-19-vaccine/
7. Pancevski B. UK delays second COVID-19 vaccine dose as Europe ponders how to speed up immunization. 2020
SARS-CoV-2 Aerosol Exhaled by Experimentally Infected Cynomolgus Monkeys

Chunmao Zhang,1 Zhendong Guo,1 Zongzheng Zhao,1 Tiecheng Wang, Liang Li, Faming Miao, Cheng Zhang, Yuanguo Li, Yuwei Gao

Author affiliations: College of Veterinary Medicine at Hebei Agricultural University, Baoding, China. (C. Zhang); Military Veterinary Research Institute, Changchun, China (C. Zhang, Z. Guo, Z. Zhao, T. Wang, L. Li, F. Miao, C. Zhang, Y. Li, Y. Gao)

DOI: https://doi.org/10.3201/eid2707.203948

We analyzed size of severe acute respiratory coronavirus 2 (SARS-CoV-2) aerosol particles shed by experimentally infected cynomolgus monkeys. Most exhaled particles were small, and virus was mainly released early during infection. By postinfection day 6, no virus was detected in breath, but air in the isolator contained large quantities of aerosolized virus.

Although airborne transmission of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has been proven possible among humans (1), cats (2), ferrets (3), and Syrian hamsters (4), the relative roles of droplets and aerosols in the airborne transmission of SARS-CoV-2 remain controversial. A recent study showed that coronavirus disease (COVID-19) patients exhaled millions of SARS-CoV-2 particles during early infection stages (5). However, the size distribution of SARS-CoV-2 aerosol particles in exhaled breath of COVID-19 patients is not clear.

To analyze size distribution of SARS-CoV-2 aerosols shed by cynomolgus monkeys, we inoculated 3 monkeys with SARS-CoV-2 via a combination of intranasal, intratracheal, and ocular routes. Monkeys were kept in individual cages placed in an isolator (biosafety housing with HEPA filters and independent ventilation system). The exhaled breath and air in the isolator were collected by a 6-stage Andersen sampler (https://tisch-env.com) at postinfection days 2, 4, and 6, and we quantified the viral RNA copies in samples (Appendix, https://wwwnc.cdc.gov/EID/article/27/7/20-3948-App1.pdf). We also determined size distribution of SARS-CoV-2 particles.

The virus particles monkeys exhaled peaked at postinfection day 2 and ranged from 11,578 to 28,336 RNA copies during a 40-minute period. On average, each monkey exhaled 503 virus particles/min and 209.5 virus particles/L of exhaled breath. At postinfection day 4, the number of exhaled virus particles decreased substantially, ranging from 3,369 to 5,134 RNA copies during a 40-minute period. On average, each monkey exhaled 106 virus particles/min and 44 virus particles/L of breath. At postinfection day 6, no viral RNA was detected in exhaled breath (Figure 1, panel A; Appendix Figure 1). At postinfection days 2, 4, and 6, viral RNA was detected in air within the isolator housing the monkeys; we detected 6,182–13,608 RNA copies during a 30-minute period (Figure 1, panel C).

We measured size distribution of SARS-CoV-2 aerosol particles shed by the monkeys. In exhaled breath of inoculated monkeys and in air in the isolator, viral RNA was detected in all size bins, 0.65–2.1 mm, 2.1–4.7 mm, and >4.7 mm, at postinfection days 2 and 4; most were concentrated in the 2.1–4.7-mm bin (Figure 1, panels B, D; Appendix Tables 1, 2). For exhaled breath, virus particles in each of the 3 size bins accounted for 27.4%, 49.6%, and 23.0% of the total virus copies/40 min, respectively; for air in the isolator, virus particles in each of the 3 size bins accounted for 3.8%, 75.0%, and 21.2% of the total virus copies/30 min, respectively (Appendix Tables 1, 2; Appendix Figure 3). Most virus particles were in the smaller particle size range (0.65–4.7 mm), accounting for 77% to 79% of the total virus particles shed by the monkeys; droplets (>4.7 mm) accounted for ∼21%–23% (Appendix Tables 1, 2; Appendix Figure 3). We tried to isolate

1These authors contributed equally to this article.
Effects of COVID-19 Vaccination Timing and Risk Prioritization on Mortality Rates, United States

Appendix

Hybrid Scenarios: Infection-Blocking and Symptom-Blocking Vaccine

Our model assumes the vaccine is either infection blocking or symptom blocking, while the reality may be somewhere in between. Therefore, we projected coronavirus disease (COVID-19) mortality under a hybrid scenario where 2 doses of the vaccine are 67% efficacious against all infections and 82% efficacious against developing symptoms (Appendix Table 2, Appendix Table 6, Appendix Figure 2).

Stochastic Compartmental Model of COVID-19 Transmission in the Austin-Round Rock Metropolitan Area

Appendix Figure 3 includes a diagram of the model structure. For each vaccine, age and risk group, we build a separate set of compartments to model the transitions between the states: susceptible ($S$), exposed ($E$), pre-symptomatic infectious ($I^p$), symptomatic infectious ($I^y$), asymptomatic infectious ($I^A$), symptomatic infectious that are hospitalized ($I^H$), recovered ($R$), and deceased ($D$). The symbols $S$, $E$, $I^p$, $I^y$, $I^A$, $I^H$, $R$, and $D$ denote the number of persons in that state in the given vaccine/age/risk group and the total size of the vaccine/age/risk group is $N = S + E + I^p + I^y + I^A + I^H + R + D$.

The model for individuals in vaccine group $v$, age group $a$, and risk group $r$ is given by:

$$\frac{dS_{v,a,r}}{dt} = \frac{\sum_{u \in V} \sum_{i \in A} \sum_{j \in K} S_{v,a,r} K_{v,a,r} \beta_0 \phi_{a,i} (\alpha_{u}^{P} I_{u,i,j}^{P} + \alpha_{u}^{A} I_{u,i,j}^{A} + \omega_{u}^{Y} I_{u,i,j}^{Y})}{N_i}$$

$$\frac{dE_{v,a,r}}{dt} = \frac{\sum_{u \in V} \sum_{i \in A} \sum_{j \in K} S_{v,a,r} K_{v,a,r} \beta_0 \phi_{a,i} (\alpha_{u}^{P} I_{u,i,j}^{P} + \alpha_{u}^{A} I_{u,i,j}^{A} + \omega_{u}^{Y} I_{u,i,j}^{Y})}{N_i} - \sigma E_{v,a,r}$$
\[
\frac{dI^A_{v,a,r}}{dt} = (1 - \lambda_{v,a,r} \tau)\sigma E_{v,a,r} - \gamma^A I^A_{v,a,r}
\]
\[
\frac{dI^P_{v,a,r}}{dt} = \lambda_{v,a,r} \tau \sigma E_{v,a,r} - \rho I^P_{v,a,r}
\]
\[
\frac{dI^Y_{v,a,r}}{dt} = \rho I^P_{v,a,r} - (1 - \pi)\gamma^Y I^Y_{v,a,r} - \pi\eta I^Y_{v,a,r}
\]
\[
\frac{dI^H_{v,a,r}}{dt} = \pi\eta I^Y_{v,a,r} - (1 - \nu)\gamma^H I^H_{v,a,r} - \nu\mu I^H_{v,a,r}
\]
\[
\frac{dR_{v,a,r}}{dt} = \gamma^A I^A_{v,a,r} + (1 - \pi)\gamma^Y I^Y_{v,a,r} + (1 - \nu)\gamma^H I^H_{v,a,r}
\]
\[
\frac{dD_{v,a,r}}{dt} = \nu\mu I^H_{v,a,r}
\]

where \( V, A, \) and \( K \) are all possible vaccine, age and risk groups, \( \omega^A, \omega^Y, \omega^P \) are relative infectiousness of the \( I^A, I^Y, I^P \) compartments, respectively, \( \beta_0 \) is baseline transmission rate, \( \phi_{a,i} \) is the mixing rate between age group \( a, i \in A, \gamma^A, \gamma^Y, \gamma^H \) are the recovery rates for the \( I^A, I^Y, I^H \) compartments, respectively, \( \sigma \) is the exposed rate, \( \tau \) is the symptomatic ratio, \( \rho \) is the rate from pre-symptomatic to symptomatic, \( \pi \) is the proportion of symptomatic individuals requiring hospitalization, \( \eta \) is rate at which hospitalized cases enter the hospital following symptom onset, \( \nu \) is mortality rate for hospitalized cases, and \( \mu \) is rate at which terminal patients die.

The transition between each vaccine group \( v, \) is given by

\[
\frac{dX_{U,a,r}}{dt} = -\alpha^i_{a,r} X_{U,a,r}
\]
\[
\frac{dX_{W^i,a,r}}{dt} = \alpha^i_{a,r} X_{U,a,r} - \delta^i X_{W^i,a,r}
\]
\[
\frac{dX_{V^i,a,r}}{dt} = \delta^i X_{W^i,a,r} - \alpha_{a,i}^i \delta X_{V^i,a,r}
\]
\[
\frac{dX_{W^{ii},a,r}}{dt} = \alpha_{a,r}^{ii} \delta X_{V^i,a,r} - \delta^{ii} X_{W^{ii},a,r}
\]
\[
\frac{dX_{V^{ii},a,r}}{dt} = \delta^{ii} X_{W^{ii},a,r}
\]
where $X \in \{ S, E, I^A, I^P, I^Y, I^H, R, D \}$, and $v \in \{ U, W^I, V^I, W^{II}, V^{II} \}$, $\alpha^i$ is vaccination rate, $\delta^i$ is the rate that individuals that receive the first injection gain partial immunity, $\alpha^{ii}$ is second injection adherence rate, $\delta$ is the delay in second dose injection, $\delta^{ii}$ is the rate that individuals that receive the second injection gain immunity.

We model stochastic transitions between compartments using the $\tau$-leap method (3,4) with key parameters given in Appendix Tables 4–11. Assuming that the events at each time-step are independent and do not impact the underlying transition rates, the numbers of each type of event should follow Poisson distributions with means equal to the rate parameters. We thus simulate the model according to the following equations:

\[
S_{v,a,r}(t + 1) - S_{v,a,r}(t) = -P_1
\]

\[
E_{v,a,r}(t + 1) - E_{v,a,r}(t) = P_1 - P_2
\]

\[
I^A_{v,a,r}(t + 1) - I^A_{v,a,r}(t) = \left(1 - \lambda_{v,a,r} \tau\right)P_2 - P_3
\]

\[
I^P_{v,a,r}(t + 1) - I^P_{v,a,r}(t) = \lambda_{v,a,r} \tau P_2 - P_4
\]

\[
I^Y_{v,a,r}(t + 1) - I^Y_{v,a,r}(t) = P_4 - P_5 - P_6
\]

\[
I^H_{v,a,r}(t + 1) - I^H_{v,a,r}(t) = P_6 - P_7 - P_8
\]

\[
D_{v,a,r}(t + 1) - D_{v,a,r}(t) = P_7
\]

\[
R_{v,a,r}(t + 1) - R_{v,a,r}(t) = P_3 + P_5 + P_8
\]

\[
X_{U,a,r}(t + 1) - X_{U,a,r}(t) = -\alpha^i_{a,r}(t)X_{U,a,r}(t)
\]

\[
X_{W^I,a,r}(t + 1) - X_{W^I,a,r}(t) = \alpha^i_{a,r}(t)X_{U,a,r}(t) - P_9
\]

\[
X_{V^I,a,r}(t + 1) - X_{V^I,a,r}(t) = P_9 - P_{10}
\]

\[
X_{W^{II},a,r}(t + 1) - X_{W^{II},a,r}(t) = P_{10} - P_{11}
\]

\[
X_{V^{II},a,r}(t + 1) - X_{V^{II},a,r}(t) = P_{11}
\]

with

\[
P_1 \sim \text{Pois} \left( S_{v,a,r}(t)F_{v,a,r}(t) \right)
\]
\[ P_2 \sim Pois \left( \sigma E_{v,a,r}(t) \right) \]
\[ P_3 \sim Pois \left( \gamma^A I^A_{v,a,r}(t) \right) \]
\[ P_4 \sim Pois \left( \rho I^P_{v,a,r}(t) \right) \]
\[ P_5 \sim Pois \left( (1 - \pi) \gamma^V I^V_{v,a,r}(t) \right) \]
\[ P_6 \sim Pois \left( \pi \eta I^V_{v,a,r}(t) \right) \]
\[ P_7 \sim Pois \left( \nu \mu I^H_{v,a,r}(t) \right) \]
\[ P_8 \sim Pois \left( (1 - \nu) \nu^H I^H_{v,a,r}(t) \right) \]
\[ P_9 \sim Pois \left( \delta^i X^i_{W^i,a,r}(t) \right) \]
\[ P_{10} \sim Pois \left( \alpha^{ii} \delta X^i_{V^{ii},a,r}(t) \right) \]
\[ P_{11} \sim Pois \left( \delta^{ii} X^i_{W^{ii},a,r}(t) \right) \]

and where \( F_{v,a,r} \) denotes the force of infection for individuals in vaccine group \( v \), age group \( a \), and risk group \( r \) and is given by:

\[
F_{v,a,r}(t) = \sum_{u \in V} \sum_{i \in A} \sum_{j \in K} \kappa_{v,a,r} \beta_0 \phi_{a,i} \left( \omega^i_{u,i,j} + \omega^A_{u,i,j} + \omega^V_{u,i,j} \right) / N_i. 
\]

**Model Parameters**

We developed a compartmental model of COVID-19 transmission in a US city, the Austin–Round Rock Metropolitan Statistical Area, using a model period of November 8, 2020–September 17, 2021. We applied various vaccine rollout scenarios and modeled each age-risk subgroup with a separate set of compartments. (Appendix Figure 3). Upon infection, susceptible persons progress to exposed and then to either presymptomatic infectious or asymptomatic infectious. All asymptomatic cases eventually progress to a recovered class where they remain protected from future infection; presymptomatic cases progress to symptomatic then either are hospitalized or recover. Mortality varies by age group and risk group and is assumed to be preceded by hospitalization. Within each compartment, individuals are divided by vaccination status: unvaccinated, newly vaccinated with the first dose, vaccinated with the first dose, newly
vaccinated with the second dose, and fully vaccinated with the second dose. We modeled stochastic transitions between compartments and sample vaccine efficacy parameters from distributions to capture uncertainty, while keeping all other parameters fixed.

**Initial Conditions**

Initial conditions were derived using a COVID-19 healthcare forecasting model that we developed in a partnership with the city of Austin and use to provide daily transmission and healthcare projections on a public dashboard (5). The forecasting model is almost identical to the model in this study, but without vaccines. Specifically, it is an expanded stochastic SEIR model with 8 disease progression compartments, including symptomatic, presymptomatic, asymptomatic patients, and hospitalization. The population is divided into 5 age groups, with different rates of contacts within and between age groups, a high-risk category with each age group, and age- and risk-specific rates of hospitalization. The demographic, health, and mixing parameters are identical to those assumed in this study.

To make daily dashboard projections, we incorporate anonymized local mobility data from SafeGraph (S. Gao et al., unpub. data, http://arxiv.org/abs/2004.04544) into transmission rate. We assume published estimates for all disease progression parameters and calibrate the remaining unknown states and parameters to local COVID-19 hospital admissions and discharge data using iterated filtering made available through the POMP R package (39). The result of the statistical inference is posterior densities for parameters governing the impact of mobility on transmission and the reporting process of hospitalization data, and for hidden states of the model including the number of infected persons (40).

To obtain initial conditions for this study, we used the states in the fitted model based on data through November 7, 2020 (Appendix Table 11).

**Sensitivity Analyses**

**Sensitivity Analysis with Respect to the Effective Reproduction Number \((R_e)\) of the Virus on November 8, 2021**

Our base scenarios assume an effective reproductive number \((R_e)\) of 1.2 (5). Here, we provide projections based on 3 alternative transmission rate scenarios: \(R_e = 1.5\), \(R_e = 1.05\), and \(R_e\)
increasing linearly from 1.2 to 1.8 by May 1, 2021, and remaining at 1.8 through September 17, 2021.

Under the lowest transmission rate scenario \((R_e = 1.05)\), the pace of the epidemic is slower and the expected effect of vaccinations on overall mortality and the duration of the pandemic is greater than in the base case (Appendix Figure 4). Under the moderate transmission scenario \((R_e = 1.5)\), the reverse occurs. The pandemic sweeps through and achieves herd immunity rapidly, leaving little opportunity for vaccines to prevent infections and deaths.

The increasing transmission rate scenario roughly models the emergence and rapid spread of a more transmissible SARS-CoV-2 variant, like B.1.1.7. The pandemic wave reaches a peak at a similar time to the base case, but reaches a much higher prevalence of COVID-19. Overall, vaccination has a lower impact on COVID-19 mortality and the duration of the pandemic wave, but risk-based prioritization and community uptake have similar relative impacts.

Consider the best of the vaccination scenarios: 90% uptake of a perfectly-prioritized rollout of a vaccine that reduces susceptibility beginning on January 15, 2021. The expected COVID-19 deaths averted between January 15 and September 17, 2021 are 60% (95% CI 51%–67%) for \(R_e = 1.05\), 56% (95% CI 51%–60%) for \(R_e = 1.2\), 22% (95% CI 17%–25%) for \(R_e = 1.5\), and 41% for dynamic \(R_e\) scenarios. The expected numbers of COVID-19 deaths per 100,000 population during the peak week are 27 (95% CI 22–33) for \(R_e = 1.05\), 107 (95% CI 100–122) for \(R_e = 1.2\), 386 (95% CI 370–407) for \(R_e = 1.5\); and 439 (95% CI 413–474) for dynamic \(R_e\).

**Sensitivity Analysis with Respect to Prior Immunity and the Prevalence of Infections as of November 8, 2020**

Our base scenarios assume that 7.6% of the population have obtained immunity due to prior infection by early November and that 0.3% of the population is infected on the first day of the simulation (November 8, 2020) (Appendix Table 11) (5). In Figure 5, we provide projections assuming that twice as many persons are infected (0.6%) are infected at the start of each simulation or twice as many persons (15.2%) are immune at the outset. All projections assume the same transmission rate per contact, which was derived to produce a reproduction number of \(R_e = 1.2\) in the absence of prior immunity (Appendix Table 5).
Sensitivity Analysis with Respect to Vaccine Efficacy

Our original analyses assume 95% vaccine efficacy following 2 doses and 82% efficacy following a single dose (Appendix Table 6). However, recent studies have raised concerns that some SARS-CoV-2 vaccines may have reduced efficacy against emerging and future variants, including B.1.351 (41). We provide projections assuming that the vaccine efficacy is reduced by 50% (Appendix Figure 6).

Sensitivity Analysis with Respect to Vaccine Efficacy after a Single Dose

Our baseline analyses assume 82% vaccine efficacy following a single dose (Figure panel C, Appendix Table 6). However, a recent commentary (42) suggests that the efficacy following a single dose of the Pfizer-BioNTech vaccine may be even higher than 82% and provides revised estimates that exclude trial data (35) from the 2 weeks immediately following the first injection, while the body is still building an immune response. Appendix Figure 7 provides projections assuming single-dose efficacies of 52.4% (95% CI 29.5%–68.4%), 68.5% (95% CI 46.5%–81.5%), 82% (95% CI 75.6%–86.9%), and 92.6% (95% CI 69%–98.3%) (42).

Estimation of Age-Stratified Proportion of Population at High Risk for COVID-19 Complications

We estimate age-specific proportions of the population at high risk for complications from COVID-19 based on data for Austin, Texas and Round Rock, Texas from the CDC’s 500 Cities Project (Appendix Figure 8) (16). We assume that high risk conditions for COVID-19 are the same as those specified for influenza by the CDC (13). CDC’s 500 Cities Project provides city-specific estimates of prevalence for several of these conditions among adults (17). The estimates were obtained from the 2015–2016 Behavioral Risk Factor Surveillance System (BRFSS) data using a small-area estimation methodology called multilevel regression and poststratification (14,15). It links geocoded health surveys to high spatial resolution population demographic and socioeconomic data (14).

Estimating High-Risk Proportions for Adults

To estimate the proportion of adults at high risk for complications, we use the CDC’s 500 Cities data, as well as data on the prevalence of HIV/AIDS, obesity, and pregnancy among adults (Appendix Table 12).
The CDC 500 cities dataset includes the prevalence of each condition on its own, rather than the prevalence of multiple conditions (e.g., dyads or triads). Thus, we use separate co-morbidity estimates to determine overlap. Reference about chronic conditions (18) gives US estimates for the proportion of the adult population with 0, 1, or ≥2 chronic conditions, per age group. Using this and the 500 cities data we can estimate the proportion of the population pHRI in each age group in each city with ≥1 chronic condition listed in the CDC 500 cities data (Appendix Table 12) putting them at high risk for flu complications.

**HIV**

We use the data from Table 20a in CDC HIV Surveillance Report (19) to estimate the population in each risk group living with HIV in the United States (last column, 2015 data). Assuming independence between HIV and other chronic conditions, we increase the proportion of the population at high-risk for influenza to account for persons living with HIV but no other underlying conditions.

**Morbid Obesity**

A BMI >40 kg/m² indicates morbid obesity and is considered high risk for influenza. The 500 Cities Project reports the prevalence of obese persons in each city with BMI over 30 kg/m² (not necessarily morbid obesity). We used the data from Table 1 in Sturm and Hattori (20) to estimate the proportion of people with BMI>30 that actually have BMI>40 (across the United States); we then apply this to the 500 Cities obesity data to estimate the proportion of persons who are morbidly obese in each city. Table 1 of Morgan et al. (21) suggests that 51.2% of morbidly obese adults have ≥1 other high-risk chronic condition, and update our high-risk population estimates accordingly to account for overlap.

**Pregnancy**

We estimated the number of pregnant women in each age group and each city, following the methodology in CDC Reproductive Health Report (22). We assume independence between any of the high-risk factors and pregnancy, and further assume that half the population are women.
Estimating High-Risk Proportions for Children

Because the 500 Cities Project reports data for only adults >18 years of age, we took a different approach to estimating the proportion of children at high risk for severe influenza. The 2 most prevalent risk factors for children are asthma and obesity; we also accounted for childhood diabetes, HIV, and cancer.

From Miller et al. (23), we obtained national estimates of chronic conditions in children. For asthma, we assumed that variation among cities will be similar for children and adults. Thus, we used the relative prevalences of asthma in adults to scale our estimates for children in each city. The prevalence of HIV in children are taken from CDC HIV surveillance report (19) and the prevalence of cancer from the CDC cancer research report (24).

We first estimated the proportion of children who had asthma, diabetes, cancer, or HIV (assuming no overlap in these conditions). We estimated city-level morbid obesity in children using the estimated morbid obesity in adults multiplied by a national constant ratio for each age group estimated from Hales et al. (25); this ratio represents the prevalence in morbid obesity in children given the prevalence observed in adults. From Morgan et al. (21), we estimated that 25% of morbidly obese children have another high-risk condition and adjusted our final estimates accordingly.

Resulting Estimates

We compared our estimates for the Austin–Round Rock Metropolitan Area to published national-level estimates (26) of the proportion of each age group with underlying high-risk conditions (Appendix Table 13). The biggest difference was observed in older adults; Austin had a lower proportion at risk for complications for COVID-19 than the national average; for the 25–39 year age group, the high-risk proportion was slightly higher than the national average.

References

1. Texas Department of State Health Services. COVID-19 vaccine allocations. [cited 2021 Feb 4]. https://dshs.texas.gov/coronavirus/immunize/vaccineallocations.aspx

2. U.S. public now divided over whether to get COVID-19 vaccine. 2020 [cited 2020 Dec 14]. https://www.pewresearch.org/science/2020/09/17/u-s-public-now-divided-over-whether-to-get-covid-19-vaccine
3. Keeling MJ, Rohani P. Modeling infectious diseases in humans and animals. Princeton University Press; 2011. 408 p.

4. Gillespie DT. Approximate accelerated stochastic simulation of chemically reacting systems. J Chem Phys. 2001;115:1716–33. https://doi.org/10.1063/1.1378322

5. University of Texas COVID-19 Modeling Consortium. COVID-19 healthcare forecasts: Austin, Texas [cited 2020 Dec 8]. https://covid-19.tacc.utexas.edu/dashboards/austin

6. Austin Independent School District. Calendar of events. [cited 2020 Mar 26]. https://www.austinisd.org/calendar

7. Prem K, Cook AR, Jit M. Projecting social contact matrices in 152 countries using contact surveys and demographic data. PLoS Comput Biol. 2017;13:e1005697. https://doi.org/10.1371/journal.pcbi.1005697 PubMed

8. Mossong J, Hens N, Jit M, Beutels P, Auranen K, Mikolajczyk R, et al. Social contacts and mixing patterns relevant to the spread of infectious diseases. PLoS Med. 2008;5:e74. https://doi.org/10.1371/journal.pmed.0050074 PubMed

9. Diekmann O, Heesterbeek JAP, Roberts MG. The construction of next-generation matrices for compartmental epidemic models. J R Soc Interface. 2010;7:873–85. https://doi.org/10.1098/rsif.2009.0386 PubMed

10. He X, Lau EHY, Wu P, Deng X, Wang J, Hao X, et al. Temporal dynamics in viral shedding and transmissibility of COVID-19. Nat Med. 2020;26:672–5. PubMed https://doi.org/10.1038/s41591-020-0869-5

11. Gudbjartsson DF, Helgason A, Jonsson H, Magnusson OT, Melsted P, Norddahl GL, et al. Spread of SARS-CoV-2 in the Icelandic population. N Engl J Med. 2020;382:2302–15. PubMed https://doi.org/10.1056/NEJMoa2006100

12. Zhang J, Litvinova M, Wang W, Wang Y, Deng X, Chen X, et al. Evolving epidemiology and transmission dynamics of coronavirus disease 2019 outside Hubei province, China: a descriptive and modelling study. Lancet Infect Dis. 2020;20:793–802. PubMed https://doi.org/10.1016/S1473-3099(20)30230-9

13. US Centers for Disease Control and Prevention. People at high risk for flu complications. 2019 [cited 2020 Mar 26]. https://www.cdc.gov/flu/highrisk/index.htm

14. US Centers for Disease Control and Prevention. Behavioral Risk Factor Surveillance System. 2019 [cited 2020 Mar 26]. https://www.cdc.gov/brfss/index.html
15. Zhang X, Holt JB, Lu H, Wheaton AG, Ford ES, Greenland KJ, et al. Multilevel regression and poststratification for small-area estimation of population health outcomes: a case study of chronic obstructive pulmonary disease prevalence using the behavioral risk factor surveillance system. Am J Epidemiol. 2014;179:1025–33. https://doi.org/10.1093/aje/kwu018 PubMed

16. US Centers for Disease Control and Prevention. 500 Cities Project: local data for better health. 2019 [cited 2020 Jun 22]. https://www.cdc.gov/places/about/500-cities-2016-2019/index.html

17. US Centers for Disease Control and Prevention. Health Outcomes | 500 Cities. 2019 [cited 2020 Jun 22]. https://www.cdc.gov/500cities/definitions/health-outcomes.htm

18. Fox S, Duggan M. Who lives with chronic conditions. 2013 [cited 2020 Jun 22]. https://www.pewresearch.org/internet/2013/11/26/part-one-who-lives-with-chronic-conditions/

19. US Centers for Disease Control and Prevention. HIV surveillance report. 2016; 28. http://www.cdc.gov/hiv/library/reports/hiv-surveillance.html

20. Sturm R, Hattori A. Morbid obesity rates continue to rise rapidly in the United States. Int J Obes (Lond). 2013;37:889–91. PubMed https://doi.org/10.1038/ijo.2012.159

21. Morgan OW, Bramley A, Fowlkes A, Freedman DS, Taylor TH, Gargiullo P, et al. Morbid obesity as a risk factor for hospitalization and death due to 2009 pandemic influenza A(H1N1) disease. PLoS One. 2010;5:e9694. PubMed https://doi.org/10.1371/journal.pone.0009694

22. US Centers for Disease Control and Prevention. Estimating the number of pregnant women in a geographic area. [cited 2021 May 10]. https://www.cdc.gov/reproductivehealth/emergency/pdfs/PregnancyEstimatoBrochure508.pdf

23. Miller GF, Coffield E, Leroy Z, Wallin R. Prevalence and costs of five chronic conditions in children. J Sch Nurs. 2016;32:357–64. PubMed https://doi.org/10.1177/1059840516641190

24. American Cancer Society. Cancer facts & figures 2014. 2014 [cited 2020 Jun 22]. https://www.cancer.org/research/cancer-facts-statistics/all-cancer-facts-figures/cancer-facts-figures-2014.html

25. Hales CM, Fryar CD, Carroll MD, Freedman DS, Ogden CL. Trends in obesity and severe obesity prevalence in US youth and adults by sex and age, 2007–2008 to 2015–2016. JAMA. 2018;319:1723–5. PubMed https://doi.org/10.1001/jama.2018.3060

26. Zimmerman RK, Lauderdale DS, Tan SM, Wagener DK. Prevalence of high-risk indications for influenza vaccine varies by age, race, and income. Vaccine. 2010;28:6470–7. PubMed https://doi.org/10.1016/j.vaccine.2010.07.037
27. Martin JA, Hamilton BE, Osterman MJK, Driscoll AK, Drake P. Births: final data for 2017. Nat Vital Stati Rep. 2018;67:1–50.

28. Jatlaoui TC, Boutot ME, Mandel MG, Whiteman MK, Ti A, Petersen E, et al. Abortion surveillance—United States, 2015. MMWR Surveill Summ. 2018;67:1–45. PubMed https://doi.org/10.15585/mmwr.ss6713a1

29. Ventura SJ, Curtin SC, Abma JC, Henshaw SK. Estimated pregnancy rates and rates of pregnancy outcomes for the United States, 1990–2008. Natl Vital Stat Rep. 2012;60:1–21. PubMed

30. US Centers for Disease Control and Prevention. COVID-19 Pandemic planning scenarios. 2020 [cited 2021 Feb 13]. https://www.cdc.gov/coronavirus/2019-ncov/hcp/planning-scenarios.html

31. US Centers for Disease Control and Prevention. Evidence used to update the list of underlying medical conditions that increase a person’s risk of severe illness from COVID-19. 2020 [cited 2021 Jan 15]. https://www.cdc.gov/coronavirus/2019-ncov/need-extra-precautions/evidence-table.html

32. Richardson S, Hirsch JS, Narasimhan M, Crawford JM, McGinn T, Davidson KW, et al.; the Northwell COVID-19 Research Consortium. Presenting characteristics, comorbidities, and outcomes among 5700 patients hospitalized with COVID-19 in the New York City area. JAMA. 2020;323:2052–9. PubMed https://doi.org/10.1001/jama.2020.6775

33. Lewnard JA, Liu VX, Jackson ML, Schmidt MA, Jewell BL, Flores JP, et al. Incidence, clinical outcomes, and transmission dynamics of severe coronavirus disease 2019 in California and Washington: prospective cohort study. BMJ. 2020;369:m1923. PubMed https://doi.org/10.1136/bmj.m1923

34. Tindale L, Stockdale JE, Coombe M, Garlock E, Lau WYV, Saraswat M, et al. Evidence for transmission of COVID-19 prior to symptom onset. eLife. 2020;9:e57149. https://doi.org/10.7554/eLife.57149

35. Vaccines and Related Biological Products Advisory Committee. December 10, 2020 meeting briefing document—FDA. 2020 [cited 2021 May 11]. https://www.fda.gov/media/144245

36. Vaccines and Related Biological Products Advisory Committee December 17, 2020 meeting briefing document addendum—sponsor. 2020 [cited 2021 May 11]. https://www.fda.gov/media/144453

37. US Department of Defense. Operation Warp Speed leaders say 20 million COVID-19 vaccines may be available this month. 2020 [cited 2021 Jan 13].
42. Skowronski DM, De Serres G, Gruber WC. Safety and efficacy of the BNT162b2 mRNA COVID-19 vaccine. [Reply]. N Engl J Med. 2021;384:1576–8. PubMed [https://doi.org/10.1056/NEJMc2036242]

### Appendix Table 1. Percent of deaths from coronavirus disease averted after January 15, 2021 by type of vaccination rollout for infection-blocking or symptom-blocking vaccines*

| Vaccine schema | January 15 rollout | February 15 rollout |
|----------------|---------------------|---------------------|
|                | 50% uptake | 70% uptake | 90% uptake | 50% uptake | 70% uptake | 90% uptake |
| **Infection-blocking 2 dose** | | | | | | |
| No priorities  | 48 (43–53) | 48 (42–54) | 48 (43–53) | 30 (25–36) | 30 (25–35) | 31 (24–36) |
| Adults >65 years of age | 49 (44–54) | 50 (45–55) | 51 (45–55) | 32 (24–37) | 33 (27–38) | 33 (27–39) |
| High-risk adults | 52 (46–56) | 54 (48–58) | 54 (50–59) | 34 (27–39) | 35 (30–40) | 36 (30–41) |
| Adults >65 years of age + high risk | 51 (46–56) | 52 (48–58) | 53 (48–58) | 33 (29–39) | 34 (27–40) | 36 (29–40) |
| 10 Phase | 52 (47–56) | 54 (49–58) | 56 (51–60) | 34 (28–40) | 36 (30–42) | 38 (32–43) |
| **Symptom-blocking 2 dose** | | | | | | |
| No priorities  | 32 (25–37) | 32 (27–38) | 33 (26–38) | 20 (13–26) | 19 (13–25) | 19 (12–25) |
| Adults >65 years of age | 36 (30–42) | 40 (34–44) | 42 (38–46) | 23 (18–29) | 26 (19–32) | 28 (20–33) |
| High-risk adults | 38 (32–43) | 41 (35–47) | 44 (39–49) | 25 (18–30) | 27 (21–33) | 29 (24–34) |
| Adults >65 years of age + high risk | 38 (32–43) | 43 (38–49) | 45 (41–50) | 25 (19–30) | 28 (22–33) | 30 (24–35) |
| 10 Phase | 40 (35–45) | 46 (40–51) | 51 (45–56) | 27 (21–33) | 32 (25–37) | 34 (29–39) |
| **Infection-blocking 1 dose** | | | | | | |
| No priorities  | 65 (61–68) | 65 (62–69) | 65 (63–69) | 44 (39–49) | 44 (40–49) | 44 (40–50) |
| Adults >65 years of age | 66 (62–69) | 66 (63–70) | 68 (63–71) | 45 (41–50) | 47 (42–52) | 47 (43–52) |
| High-risk adults | 66 (62–69) | 68 (64–72) | 70 (66–73) | 46 (41–52) | 48 (43–53) | 49 (43–54) |
| Adults >65 years of age + high risk | 66 (63–70) | 68 (64–71) | 69 (65–73) | 46 (41–51) | 48 (43–53) | 49 (44–54) |
| 10 Phase | 66 (63–70) | 69 (66–73) | 71 (67–74) | 47 (41–51) | 49 (44–54) | 51 (46–55) |
| **Symptom-blocking 1 dose** | | | | | | |
| No priorities  | 46 (39–50) | 50 (45–55) | 51 (47–56) | 31 (25–36) | 33 (27–39) | 34 (28–38) |
| Adults >65 years of age | 48 (42–52) | 54 (49–58) | 57 (52–61) | 33 (28–38) | 37 (32–43) | 39 (33–45) |
| High-risk adults | 49 (43–53) | 55 (49–60) | 58 (54–63) | 34 (29–39) | 38 (33–43) | 41 (35–45) |
| Adults >65 years of age + high risk | 49 (43–53) | 55 (51–60) | 60 (55–64) | 34 (28–39) | 39 (33–43) | 42 (36–46) |
| 10 Phase | 50 (45–54) | 57 (52–62) | 63 (58–67) | 35 (29–40) | 41 (36–46) | 45 (39–50) |

*Values are medians and 95% confidence intervals based on 200 pairs of stochastic simulations. Deaths averted are computed by comparing a simulation of the specified vaccine strategy to a simulation without vaccination.
Appendix Table 2. Percent of deaths from coronavirus disease averted after January 15, 2021 by type of vaccination rollout for hybrid scenarios*.

| Vaccine schema | January 15 rollout | February 15 rollout |
|----------------|---------------------|----------------------|
|                | 50% uptake          | 70% uptake          | 90% uptake          | 50% uptake | 70% uptake | 90% uptake |
| Hybrid 2 dose  |                     |                     |                     |            |            |            |
| No priorities  | 47 (41–51)          | 46 (41–51)          | 47 (41–52)          | 30 (22–35) | 29 (23–35) | 29 (22–35) |
| Adults >65 years of age | 47 (43–53)  | 48 (43–54)          | 50 (43–55)          | 31 (25–36) | 32 (25–37) | 33 (27–38) |
| High-risk adults| 50(43–55)          | 52(46–57)          | 53(47–58)          | 33 (27–38) | 34 (27–39) | 35(29–40)  |
| Adults >65 years of age + high risk | 50 (45–54)  | 51 (46–56)          | 53 (46–57)          | 32 (25–37) | 34 (27–39) | 34 (28–40) |
| 10 Phase       | 50 (45–55)          | 53 (48–58)          | 56 (51–60)          | 34 (29–39) | 36 (30–41) | 37 (31–44) |

Values are medians and 95% confidence intervals based on 200 pairs of stochastic simulations. Deaths averted are computed by comparing a simulation of the specified vaccine strategy to a simulation without vaccination.

Appendix Table 3. Initial conditions, school calendar, and contact rates used in modeling of effects of vaccination by prioritization on mortality.

| Variable                               | Settings                                                                 |
|----------------------------------------|---------------------------------------------------------------------------|
| Initial day of simulation              | 11/8/2020                                                                 |
| Initial number infected                | Based on estimates for Austin, Texas, given Appendix Table 11 (5)         |
| Age-specific and day-specific contact rates' | Home, work, other and school matrices provided in Appendix Tables 7–10    |
|                                        | Typical weekday = home + work + other + school                             |
|                                        | Weekends and holiday weekdays = home + other                              |
| School calendar                        | Austin Independent School District calendar (2019–2020, 2020–2021) (6) |

We assume the age-specific contact rates given in (6), which takes the contact numbers estimated through diary-based POLYMOLY study in Europe (9) and extrapolates to the United States. The values in Appendix Tables 7–10 are the assumed daily contacts between each pair of age groups at home, school, work, and all other places, respectively. These contact matrices are used to adjust the transmission rate between age groups. The accuracy of the contact matrices is limited by (i) possible biases with the original diary-based study (9), (ii) assumptions made when projecting the original study to the U.S (7), and (iii) impacts of COVID-19 policies and perceptions on daily contact patterns.

Appendix Table 4. Epidemiologic parameters for modeling of effects of vaccination by prioritization on mortality*.

| Parameters | Best-guess values | Source |
|------------|-------------------|--------|
| $R_0$: effective reproduction rate | 1.2 | Derived by next-generation matrix method to yield $R_e = 1.2$ without prior immunity (9) |
| $\beta$: baseline transmission rate | 0.0183 | |
| $\gamma^A$: recovery rate on asymptomatic compartment $i$ | 0.1587 | (10) |
| $\gamma^Y$: recovery rate on symptomatic non-treated compartment | 0.25 | (10) |
| $r$: symptomatic proportion (%) | 57 | (11) |
| $\sigma$: exposed rate | 0.3448 | (12) |
| $P$: proportion of infections occurring in pre-symptomatic period (%) | 44 | (10) |
| $\omega^P$: relative infectiousness of infectious individuals in compartment $i^P$ | 1.3669 | $\omega^P = \frac{Y_{HR}}{\eta} + \frac{1 - Y_{HR}}{Y^P} \omega^Y \rho P$ |
| $\omega^A$: relative infectiousness of infectious individuals in compartment $i^A$ | 0.67 | (10) |
| $\rho$: symptom onset rate | 0.43478 | 2.3 d average pre-symptomatic period |
| $IFR$: age-stratified infection fatality ratio (%) | Overall: [0.0016, 0.00495, 0.008428, 1.007711, 3.37149] | Age adjusted from Verity et al., unpublished data, [857] |
| | Low risk: [0.00137, 0.00386, 0.06334, 0.60254, 1.73687] | [857] |
| | High risk: [0.00412, 0.01157, 0.19001, 1.80762, 5.2106] | |
| $YFR$: age-stratified symptomatic fatality ratio (%) | Overall: [0.00281, 0.00868, 0.14785, 1.75458, 5.9149] | $YFR = \frac{IFR}{\tau}$ |
| | Low risk: [0.00241, 0.00677, 0.11112, 1.05709, 3.04713] | |
| | High risk: [0.00722, 0.0203, 0.33336, 3.17127, 9.1414] | |
### Appendix Table 5. Hospitalization parameters

| Parameters | Value | Source |
|------------|-------|--------|
| \( \gamma \): recovery rate in hospitalized compartment | 0.0935 | Austin admissions and discharge data (Avg = 10.96. 95% CI = 9.37 to 12.76) |
| \( \gamma_H \): recovery rate in hospitalized compartment | | |
| \( \gamma_Y \): symptomatic case hospitalization rate (%) | | |
| \( \pi \): rate of symptomatic individuals going to hospital, age-specific | | |
| \( \eta \): symptomatic case hospitalization rate (%) | | |
| \( \mu \): rate of hospitalization based on hospital admission studies | | |
| \( HFR \): hospitalized fatality ratio, age specific (%) | | |
| \( v \): death rate on hospitalized individuals, age specific | | |

### Appendix Table 6. Vaccine parameters

| Parameters | Value | Source |
|------------|-------|--------|
| \( ve \): efficacy of full course vaccination, infection-blocking or symptom blocking | After 1st and before 2nd dose: | (35) |
| \( ve\sim \): Triangular (0.295, 0.524, 0.684) | | |
| \( ve \): vaccine efficacy - single dose only, infection-blocking or symptom blocking | After single dose only: | (35) |
| \( ve\sim \): Triangular (0.898, 0.948, 0.976) | | |
| \( ve \): vaccine efficacy - single dose only, infection-blocking or symptom blocking | Sensitivity analysis | |
| \( ve\sim \): Triangular (0.295, 0.524, 0.684) | | |
| \( ve \): vaccine efficacy - single dose only, infection-blocking or symptom blocking | | |
| \( ve \): vaccine efficacy - single dose only, hybrid model | Efficacy against infection: | (35,36) |
| \( ve\sim \): Triangular (0.69, 0.926, 0.983) | | |
| \( ve \): vaccine efficacy - single dose only, hybrid model | Efficacy against infection: | |
| \( ve\sim \): Triangular (0.926, 1.0, 1.05) | | |
| \( ve \): vaccine efficacy - single dose only, hybrid model | Efficacy against symptomatic infection: | |
| \( ve\sim \): Triangular (0.261, 0.455, 0.603) | | |
| \( vu \): vaccine uptake rate | 0.5, 0.7, 0.9 | 0.5 is based on a survey conducted by PEW Research Center in September (2); 0.7 is based on a survey conducted by PEW Research Center in May (2); 0.9 is hypothetical assumption |

\( a \): second dose return rate

0.8

We assume that the duration of the infectious period is the same for asymptomatic and symptomatic cases.
### Appendix Table 7. Home contact matrix (daily number contacts by age group at home)

| Age Group | 0–4 y | 5–17 y | 18–49 y | 50–64 y | >65 y |
|-----------|-------|--------|---------|---------|-------|
| 0–4 y     | 0.5   | 0.9    | 2.0     | 0.1     | 0.0   |
| 5–17 y    | 0.2   | 1.7    | 1.9     | 0.2     | 0.0   |
| 18–49 y   | 0.2   | 0.9    | 1.7     | 0.2     | 0.0   |
| 50–64 y   | 0.2   | 0.7    | 1.2     | 1.0     | 0.1   |
| >65 y     | 0.1   | 0.7    | 1.0     | 0.3     | 0.6   |

### Appendix Table 8. School contact matrix (daily number contacts by age group at school)

| Age Group | 0–4 y | 5–17 y | 18–49 y | 50–64 y | >65 y |
|-----------|-------|--------|---------|---------|-------|
| 0–4 y     | 1.0   | 0.5    | 0.4     | 0.1     | 0.0   |
| 5–17 y    | 0.2   | 3.7    | 0.9     | 0.1     | 0.0   |
| 18–49 y   | 0.0   | 0.7    | 0.8     | 0.0     | 0.0   |
| 50–64 y   | 0.1   | 0.8    | 0.5     | 0.1     | 0.0   |
| >65 y     | 0.0   | 0.0    | 0.1     | 0.0     | 0.0   |

### Appendix Table 9. Work contact matrix (daily number contacts by age group at work)

| Age Group | 0–4 y | 5–17 y | 18–49 y | 50–64 y | >65 y |
|-----------|-------|--------|---------|---------|-------|
| 0–4 y     | 0.0   | 0.0    | 0.0     | 0.0     | 0.0   |
| 5–17 y    | 0.0   | 0.1    | 0.4     | 0.0     | 0.0   |
| 18–49 y   | 0.0   | 0.2    | 4.5     | 0.8     | 0.0   |
| 50–64 y   | 0.0   | 0.1    | 2.8     | 0.9     | 0.0   |
| >65 y     | 0.0   | 0.0    | 0.1     | 0.0     | 0.0   |

### Appendix Table 10. Others contact matrix (daily number contacts by age group at other locations)

| Age Group | 0–4 y | 5–17 y | 18–49 y | 50–64 y | >65 y |
|-----------|-------|--------|---------|---------|-------|
| 0–4 y     | 0.7   | 0.7    | 1.8     | 0.6     | 0.3   |
| 5–17 y    | 0.2   | 2.6    | 2.1     | 0.4     | 0.2   |
| 18–49 y   | 0.1   | 0.7    | 3.3     | 0.6     | 0.2   |
| 50–64 y   | 0.1   | 0.3    | 2.2     | 1.1     | 0.4   |
| >65 y     | 0.0   | 0.2    | 1.3     | 0.8     | 0.6   |

### Appendix Table 11. Initial states of model compartments*

| Age       | Risk | Value | Source |
|-----------|------|-------|--------|
| 0–4 y     | High | 8,809 |        |
| 0–4 y     | Low  | 121,035 |  (37) and discussion with CDC |
| 5–17 y    | High | 33,884 |        |
| 5–17 y    | Low  | 296,425 |        |
| 18–49 y   | High | 142,443 |        |
| 18–49 y   | Low  | 834,952 |        |
| 50–64 y   | High | 100,333 |        |
| 50–64 y   | Low  | 231,158 |        |
| >65 y     | High | 100,101 |        |
| >65 y     | Low  | 127,788 |        |

*Values indicate the number of individuals in each age-risk group compartment at the start of the simulations.
Appendix Table 12. High-risk conditions for influenza and data sources for prevalence estimation

| Condition                                                                 | Data source                                      |
|---------------------------------------------------------------------------|--------------------------------------------------|
| Cancer (except skin), chronic kidney disease, COPD, coronary heart disease, stroke, asthma, diabetes | CDC 500 Cities (16)                              |
| HIV/AIDS                                                                  | CDC HIV Surveillance report (17)                 |
| Obesity                                                                   | CDC 500 Cities (16), Sturm and Hattori (20), Morgan et al. (21) |
| Pregnancy                                                                 | National Vital Statistics Reports (27) and abortion data (28) |

Appendix Table 13. Comparison between published national estimates and Austin-Round Rock MSA estimates of the percent of the population at high risk for influenza/COVID-19 complications*

| Age group | National estimates (25) | Austin | Pregnant women (proportion of age group) |
|-----------|-------------------------|--------|-----------------------------------------|
| 0–6 mo    | NA                      | 6.8    | –                                       |
| 6 mo–4 y  | 6.8                     | 7.4    | –                                       |
| 5–9 y     | 11.7                    | 11.6   | –                                       |
| 10–14 y   | 11.7                    | 13.0   | –                                       |
| 15–19 y   | 11.8                    | 13.3   | 1.7                                     |
| 20–24 y   | 12.4                    | 10.3   | 5.1                                     |
| 25–34 y   | 15.7                    | 13.5   | 7.8                                     |
| 35–39 y   | 15.7                    | 17.0   | 5.1                                     |
| 40–44 y   | 15.7                    | 17.4   | 1.2                                     |
| 45–49 y   | 15.7                    | 17.7   | –                                       |
| 50–54 y   | 30.6                    | 29.6   | –                                       |
| 55–60 y   | 30.6                    | 29.5   | –                                       |
| 60–64 y   | 30.6                    | 29.3   | –                                       |
| 65–69 y   | 47.0                    | 42.2   | –                                       |
| 70–74 y   | 47.0                    | 42.2   | –                                       |
| >75 y      | 47.0                    | 42.2   | –                                       |

*Proportions for Austin exclude pregnancy. NA, not available.

Appendix Figure 1. COVID-19 vaccine doses allocated to the Austin-Round Rock Metropolitan Statistical Area in comparison to the model assumption (1). Red represents doses manufactured by Moderna and green represents vaccines manufactured by Pfizer-BioNTech. Black points indicate the rollout assumed by the model.
Appendix Figure 2. Projected COVID-19 mortality in the Austin-Round Rock Metropolitan Statistical Area for November 8, 2020–September 17, 2021 under various vaccine rollout scenarios. A) COVID-19 deaths averted after January 15, 2021 for a hybrid type of vaccination protection under combinations of: vaccine uptake, either 50% or 90% (x-axis); rollout dates, either January 15 (circles) or February 15 (triangles); and risk prioritization, either no priority (gray), prioritize all adults over 65 y (light blue), adults with high-risk comorbidities (medium blue), or the combination of the two (dark blue), or a 10-phase risk-ordered strategy (green) that sequentially vaccinates >65 y high risk, 50–64 y high risk, >65 y low risk, 18–49 y high risk, 50–64 y low risk, 18–49 y low risk, 0–4 y high risk, 5–17 y high risk, 0–4 y low risk, 5–17 y low risk. Points and whiskers indicate the median and 95% CI across 200 paired stochastic simulations. B) Weekly incident COVID-19 deaths per 100,000 assuming intermediate (70%) uptake (black) or under a 10-phase risk-based rollout of a 95% efficacious infection-blocking vaccine, starting either January 15 (orange) or February 15 (purple). The brown line assumes that only first doses are administered starting January 15. Solid lines and shading indicate the median and 95% CI across 200 stochastic simulations.
Appendix Figure 3. Compartmental model of COVID-19 transmission in the Austin-Round Rock Metropolitan Statistical Area for November 8, 2020–September 17, 2021 under various vaccine rollout scenarios. Each age-risk subgroup is modeled with a separate set of compartments. Upon infection, susceptible individuals (S) progress to exposed (E) and then to either pre-symptomatic infectious (I\textsuperscript{p}) or asymptomatic infectious (I\textsuperscript{A}). All asymptomatic cases eventually progress to a recovered class where they remain protected from future infection (R); pre-symptomatic cases progress to symptomatic (I\textsuperscript{Y}) then are either hospitalized (I\textsuperscript{H}) or recover. Mortality (D) varies by age group and risk group and is assumed to be preceded by hospitalization. Within each compartment, individuals are divided by vaccination status: unvaccinated (U), newly vaccinated with the first dose (W\textsuperscript{i}), vaccinated with the first dose (V\textsuperscript{i}), newly vaccinated with the second dose (W\textsuperscript{ii}), and fully vaccinated with the second dose (V\textsuperscript{ii}). We model stochastic transitions between compartments and sample vaccine efficacy parameters from distributions to capture uncertainty, while keeping all other parameters fixed.
Appendix Figure 4. 1. Projected COVID-19 mortality in the Austin-Round Rock Metropolitan Statistical Area for November 8, 2020–September 17, 2021 under different vaccination scenarios, assuming either $R_e = 1.05$, $R_e = 1.2$, $R_e = 1.5$, or $R_e$ increases linearly from 1.2 to 1.8 by May 1, 2021, and remains at 1.8 thereafter. A) COVID-19 deaths averted after January 15, 2021 under combinations of: vaccine uptake, either 50% (top) or 90% (bottom); type of protection, either symptom blocking (left) or infection blocking (right); rollout dates, either January 15 (circles) or February 15 (triangles); and risk prioritization, either no priority (gray), prioritize all adults over 65 y (light blue), adults with high-risk comorbidities (medium blue), or the combination of the two (dark blue), or a 10-phase risk-ordered strategy (green) that sequentially vaccinates >65 y high risk, 50–64 y high risk, >65 y low risk, 18–49 y high risk, 50–64 y low risk, 18–49 y low risk, 0–4 y high risk, 5–17 y high risk, 0–4 y low risk, 5–17 y low risk. Points and whiskers indicate the median and 95% CI across 200 paired stochastic simulations. B) Weekly incident COVID-19 deaths per 100,000 population assuming intermediate (70%) uptake (2) without vaccine (black) or under a 10-phase risk-based rollout of a 95% efficacious infection-blocking vaccine, starting either January 15 (orange) or February 15 (purple), for the indicated $R_e$ scenario. The brown lines assume that only first doses are administered starting January 15. Solid lines and shading indicate the median and 95% CI across 200 stochastic simulations.
Appendix Figure 5. Projected COVID-19 mortality in the Austin-Round Rock Metropolitan Statistical Area for November 8, 2020–September 17, 2021 under different vaccine rollout scenarios, assuming a higher initial prevalence (0.6%) or prior immunity (15.2%). A) COVID-19 deaths averted after January 15, 2021 under combinations of: vaccine uptake, either 50% (top) or 90% (bottom); type of protection, either symptom blocking (left) or infection blocking (right); initial conditions (x-axis); rollout dates, either January 15 (circles) or February 15 (triangles); and risk prioritization (colors). Points and whiskers indicate the median and 95% CI across 200 paired stochastic simulations. B) Weekly incident COVID-19 deaths per 100,000 assuming intermediate (70%) uptake (2) without vaccine (black) or under a 10-phase risk-based rollout of a 95% efficacious infection-blocking, starting either January 15 (orange) or February 15 (purple), under the original scenario (top), double initial prevalence (middle), and double prior immunity (bottom). The brown line assumes that only first doses are administered starting January 15. Solid lines and shading indicate the median and 95% CI across 200 stochastic simulations.
Appendix Figure 6. Projected COVID-19 in the Austin-Round Rock Metropolitan Statistical Area for November 8, 2020–September 17, 2021 under different vaccine rollout scenarios, assuming a 50% reduction in vaccine efficacy. A) COVID-19 deaths averted after January 15, 2021 under combinations of: vaccine uptake, either 50% (top) or 90% (bottom); type of protection, either symptom blocking (left) or infection blocking (right); vaccine efficacy, either original or half of original (x-axis); rollout dates, either January 15 (circles) or February 15 (triangles); and risk prioritization (colors). Points and whiskers indicate the median and 95% CI across 200 paired stochastic simulations. B) Weekly incident COVID-19 deaths per 100,000 assuming intermediate (70%) uptake (2) without vaccine (black) or under a 10-phase risk-based rollout of a 95% efficacious infection-blocking, starting either January 15 (orange) or February 15 (purple) with original vaccine efficacy (top) or 50% lower vaccine efficacy (bottom). The brown line assumes that only first doses are administered starting January 15. Solid lines and shading indicate the median and 95% CI across 200 stochastic simulations.
Appendix Figure 7. Projected COVID-19 mortality in the Austin-Round Rock Metropolitan Statistical Area for November 8, 2020–September 17, 2021 under different assumptions regarding the efficacy of a single dose. In both panels, the graphs from top to bottom assume single-dose efficacies of 52%, 69%, 82%
A) COVID-19 deaths averted after January 15, 2021 under combinations of: type of protection, either symptom blocking (reducing severity) or infection blocking (reducing susceptibility); 1-dose (hollow) or 2-dose (solid); and risk prioritization (colors). All projections assume intermediate (70%) vaccine uptake (2) starting January 15. Points and whiskers indicate the median and 95% CI across 200 paired stochastic simulations. B) Weekly incident COVID-19 deaths per 100,000 population assuming intermediate (70%) uptake (2) without vaccine (black) or under a ten-phase risk-based rollout of a 95% efficacious infection-blocking, starting either January 15 (orange) or February 15 (purple). The brown line assumes that only first doses are administered starting January 15. Solid lines and shading indicate the median and 95% CI across 200 stochastic simulations.

Appendix Figure 8. Demographic and risk composition of the Austin-Round Rock Metropolitan Statistical Area. Bars indicate age-specific population sizes, separated by low risk, high risk, and pregnant. High risk is defined as persons with cancer, chronic kidney disease, COPD, heart disease, stroke, asthma, diabetes, HIV/AIDS, and morbid obesity, as estimated from the CDC 500 Cities Project (16), reported HIV prevalence (19), and reported morbid obesity prevalence (20,21), corrected for multiple conditions. The population of pregnant women is derived using the CDC’s method combining fertility, abortion and fetal loss rates (27–29).