Appendix

Case and Hospitalization Data

Two sources of data were used for model fitting. Initially, models were fit to reported cases which were obtained from the Colorado Electronic Disease Reporting System (CEDRS) because hospitalization data were not yet available. CEDRS is updated daily with cases of infection with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) cases reported by private laboratories, public laboratories, hospital infection specialists, and state and local public health officials. CEDRS includes information regarding whether SARS-CoV-2 patients were hospitalized.

Starting in May, models were fit to daily SARS-CoV-2 hospital census data. These data were available daily in real-time from EMResource (https://emresource.juvare.com). The EMResource data appear to undercount hospitalizations in March compared with the reported hospitalizations among case-patients in CEDRS (Appendix Figure 1, top panel). On the basis of discussions with health officials and consistent with the literature, we inferred daily hospital census counts by using case data, assuming hospital length was an average of 8 days, starting with report date for hospitalized cases. Because of this discrepancy, we used inferred hospitalizations from the case data through April 7, and switched to EMResource daily hospital census data starting April 8.

Estimating the Start Date of the Epidemic

The first case of COVID19 reported in Colorado was reported on March 5, 2020, and had a reported symptom onset date of February 18, 2020. The next 3 reported cases had a symptom onset date of February 20, 2020. Assuming a 5.1-day incubation period (1,2), estimates that during the initial phase of the outbreak ≈85% of cases were unreported (3), and the outbreak has
a 5.2–6.4 day doubling time (4,5), we estimated the first cases occurred during January 17–29, 2020.

This estimate was derived by using the number of observed cases on a given day to estimate the number of true cases on the date of observation: \( n_t = n_o / u \), where \( u \) is the fraction of cases unobserved in early phases of the epidemic. We then used the estimated doubling time (T) to estimate the number of generations required to go from 1 case to \( n_t \) such that doubling time, \( G = \log_2(n_t) \). Because observed cases have an approximate exposure date equal to the onset date \( (D_o) \) minus the incubation period \( (I) \), we estimated the date of first introduction as \( D_o - I - T \). We estimated \( D_o \) separately for the 1 case that had an onset date of February 18 and the 3 cases that had an onset date of February 20, by using 5.2 and 6.4 as doubling times, yielding estimated date of first infection during January 17–29.

We note that this estimate is consistent with multiple evidence streams indicating introduction of SARS-CoV-2 in the Pacific Northwest in late January to early February (6,7). Such early introductions are plausible in Colorado, given the high degree of travel into the state during the winter ski season.

**Model Details**

We provide the model (Appendix Figure 2) and model equations. The model is age-structured and has 3 distinct age groups. The probability that a person becomes infected with SARS-CoV-2 is the same for all age groups. Once a susceptible person is infected, he or she moves to the exposed compartment in which they are not yet infectious. After spending time in the exposed compartment, a person can either be infectious and symptomatic or infectious and asymptomatic. We provide probabilities that an exposed person has symptoms and probabilities that a symptomatic person requires hospitalization that is age dependent (Appendix Table 1). Symptomatic persons are assumed to be infectious and presymptomatic for 1 of the 8 days they are infectious. On the basis of evidence suggesting that asymptomatic persons are less infectious than symptomatic persons (11), a parameter is included describing the ratio of infectiousness for symptomatic versus asymptomatic persons. We provide this parameter and other key model parameters (Appendix Table 2). It is assumed no further transmission occurs once the patient enters the hospital.
Model Equations

The susceptible-exposed-infected-recovered model describes the movement of the population of Colorado ($N$) through the following model compartments: susceptible ($S$), exposed ($E$), infectious and symptomatic ($I$), infectious and asymptomatic ($A$) recovered ($R$), hospitalized ($I_h$), needing care in an intensive care unit (ICU) ($I_{ICU}$), recovered from hospitalization or ICU ($R_h$ and $R_{ICU}$), and deaths ($D$). The model is age stratified, such that there are compartments for each of the 3 age strata ($i$) (<30, 30–59, ≥60 years). The subscripts indicate the age strata for each variable. For example, $A_2$ is the number of asymptotically infected persons in the second age class (30–59 years). Similarly, $\sum_{j=1}^{3} I_j$ is the sum of infected persons across the 3 age classes.

Mask wearing ($mask_t$), self-isolation of infectious persons ($sil_t$) and social distancing ($SD_t$) metrics are assumed to change over time, as indicated by the subscript.

\[
\frac{dS_i}{dt} = -\frac{\beta}{N} \star \lambda \star (1 - mask_t \star 0.03) \star (1 - sil_t) \star (1 - SD_t) \star S_i \star \sum_{j=1}^{3} I_j - \frac{\beta}{N} \star (1 - mask_t \star 0.27) \star (1 - SD_t) \star S_i \star \sum_{j=1}^{3} A_j
\]

\[
\frac{dE_i}{dt} = \frac{\beta}{N} \star \lambda \star (1 - mask_t \star 0.03) \star (1 - sil_t) \star (1 - SD_t) \star S_i \star \sum_{j=1}^{3} I_j + \frac{\beta}{N} \star (1 - mask_t \star 0.27) \star (1 - SD_t) \star S_i \star \sum_{j=1}^{3} A_j - \frac{1}{\alpha} \star E_i
\]

\[
\frac{dI_i}{dt} = \frac{pS_i}{\alpha} \star E_i - \frac{I_i}{\gamma}
\]

\[
\frac{dI_h_i}{dt} = \frac{I_i}{\gamma} \star hosp_i - \frac{I_h_i}{losh}
\]

\[
\frac{dI_{ICU_i}}{dt} = \frac{I_i}{\gamma} \star c_{ICU_i} - \frac{I_{ICU_i}}{losc}
\]

\[
\frac{dA_i}{dt} = \frac{(1 - pS_i)}{\alpha} \star E_i - \frac{A_i}{\gamma}
\]

\[
\frac{dR_i}{dt} = \frac{A_i + I_i(1 - hosp_i - c_{ICU_i})}{\gamma}
\]
\[
\frac{dR_h}{dt} = \frac{lh_i}{losh}
\]
\[
\frac{dR_c}{dt} = \frac{(1 - d) \cdot lc_i}{losc}
\]
\[
\frac{dD_i}{dt} = \frac{d \cdot lc_i}{losc}
\]

The effective reproductive number (\(R_e\)) was estimated by using the following equation:

\[
R_e = \frac{\sum_{j=1}^{3} E_{j+t}}{\alpha} \frac{\sum_{j=1}^{3} I_{t-\alpha-1}}{\gamma}
\]

**Model Identifiability**

Analysis was performed using the computation in Omlin et al. (13). In this study, we have fit a social distancing parameter that switches at 4 different timepoints in the course of the hospitalization time series. A natural question is whether or not these 4 parameters are identifiable given the data. In Omlin et al. (13), the authors describe a colinearity index (gamma), which measures how the change in 1 parameter can be compensated by the fraction \(1 - 1/\gamma\) change in other parameters. The gamma threshold for identifiability is conventionally set in the range of 10 to 15 (for \(\gamma = 10\), \(1 - 1/\gamma = 90\%\)). We considered all possible combinations of 1, 2, 3, and 4 social distancing parameters and show the threshold (Appendix Figure 3). There is no issue with identifiability when trying to estimate \(\leq 3\) parameters. The collinearity of identifying all 4 parameters is \(\approx 10\), which is right on the edge of acceptability. In practice, we have not found this to be a problem because when we consider more recent hospitalization data, the previously estimated social distancing estimates do not change appreciably.

**Data and Code**

Model code, parameters, and data used to inform the fitting process can be found at [https://github.com/agb85/CFpaper](https://github.com/agb85/CFpaper)
**References**


**Appendix Table 1.** Estimated probability that an exposed (and latently infected) individual will be symptomatic, and a symptomatic individual will need hospital care*

<table>
<thead>
<tr>
<th>Age group, years</th>
<th>Probability of symptoms, given infection</th>
<th>Probability of non-ICU hospitalization given symptomatic COVID-19</th>
<th>Probability of needing ICU care given symptomatic COVID-19</th>
</tr>
</thead>
<tbody>
<tr>
<td>Colorado specific†</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;30</td>
<td>NA</td>
<td>0.00419</td>
<td>0.004</td>
</tr>
<tr>
<td>30–59</td>
<td>NA</td>
<td>0.056</td>
<td>0.019</td>
</tr>
<tr>
<td>≥60</td>
<td>NA</td>
<td>0.089</td>
<td>0.035</td>
</tr>
<tr>
<td>From the literature‡</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;30</td>
<td>0.18</td>
<td>0.006</td>
<td>0.0003</td>
</tr>
<tr>
<td>30–59</td>
<td>0.47</td>
<td>0.059</td>
<td>0.0045</td>
</tr>
<tr>
<td>≥60</td>
<td>0.79</td>
<td>0.207</td>
<td>0.0808</td>
</tr>
</tbody>
</table>

*The probability that an exposed person becomes symptomatic is based on the literature and adjusted to the age structure of the Colorado population. The probability a symptomatic person requires hospital care is based on Colorado case and hospitalization data and compared with estimates obtained from the literature. COVID-19, coronavirus disease; ICU, intensive care unit.

†The Colorado estimates were based on 2-part optimization by using Colorado hospitalization data and our age-structured susceptible-exposed-infected-recovered model. Colorado-specific estimates were only possible to derive with sufficient data starting in May and were incorporated into the model from then onwards (fits 3 and 4) in place of the literature-derived estimates.

‡Age-dependent probabilities that an infected person is symptomatic are based on estimates of Davies et al. (8). Age-dependent probabilities of hospitalization and ICU need are based on estimates summarized by (9). Because the age compartments of published estimates do not precisely align with our age categories, these estimates are a weighted sum of the age-specific probability of symptoms and the proportion of the population of Colorado within each age-compartment. Colorado demographic data of estimated population distribution in 2020 were based on US census projections. Literature-derived estimates of age-specific hospitalization and ICU need were used for fit 1 and fit 2. Literature-derived age-specific symptomatic fraction was used for all fitting.
### Appendix Table 2. Parameters in model and value estimates and sources

<table>
<thead>
<tr>
<th>Symbol</th>
<th>Description</th>
<th>Estimate</th>
<th>Source</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\beta$</td>
<td>Transmission rate: equivalent to combination of transmission probability and contact rate</td>
<td>Estimated from model</td>
<td>Model</td>
</tr>
<tr>
<td>$\lambda$</td>
<td>Ratio of infectiousness for symptomatic vs. asymptomatic persons. Evidence suggests that asymptomatic persons are less infectious than symptomatic persons (10). However, the comparative infectiousness is unknown.</td>
<td>Estimated from model</td>
<td>Model</td>
</tr>
<tr>
<td>$m_{mask}$</td>
<td>Proportion of population wearing masks</td>
<td>Estimated from model</td>
<td>Model</td>
</tr>
<tr>
<td>$s_{il}$</td>
<td>Reduction in infectious contacts due to symptomatic persons who self-isolate after March 5, 2020</td>
<td>Estimated from model</td>
<td>Model</td>
</tr>
<tr>
<td>SD</td>
<td>Reduction in infectious contacts due to social distancing and other transmission reducing measures</td>
<td>Estimated from model</td>
<td>Model</td>
</tr>
<tr>
<td>$\alpha$</td>
<td>Incubation period</td>
<td>5.1 d</td>
<td>(1,10)</td>
</tr>
<tr>
<td>$p_{S_i}$</td>
<td>Age-specific probability of development of symptoms if infected</td>
<td>0.18 (age 0–29 years), 0.47 (age 30–59 years), 0.79 (age &gt;60 years)</td>
<td>(8)</td>
</tr>
<tr>
<td>$\gamma$</td>
<td>Recovery rate or duration of infectiousness</td>
<td>8 d</td>
<td>(11,12)</td>
</tr>
<tr>
<td>$p_{ID}$</td>
<td>Probability symptomatic cases are identified by state surveillance</td>
<td>Estimated from model</td>
<td>Model</td>
</tr>
<tr>
<td>hosp$_i$</td>
<td>Age-specific probability of non-ICU hospitalization, given symptomatic infection</td>
<td>Appendix Table 1</td>
<td>NA</td>
</tr>
<tr>
<td>losh</td>
<td>Non-ICU hospitalization length of stay</td>
<td>8 d</td>
<td>(9)</td>
</tr>
<tr>
<td>c$c_i$</td>
<td>Age-specific probability of ICU-hospitalization, given symptomatic infection</td>
<td>Appendix Table 1</td>
<td>NA</td>
</tr>
<tr>
<td>losc</td>
<td>ICU hospitalization length of stay</td>
<td>10 d</td>
<td>(9)</td>
</tr>
<tr>
<td>$D$</td>
<td>Probability of death, given ICU hospitalization</td>
<td>0.5</td>
<td>(9)</td>
</tr>
<tr>
<td>$N$</td>
<td>Total population</td>
<td>5,840,795</td>
<td>CDPHE</td>
</tr>
</tbody>
</table>

*CDPHE, Colorado Department of Public Health and Environment; ICU, intensive care unit. NA, not applicable.
Appendix Figure 1. Top: daily number of hospitalized coronavirus disease patients based on Colorado Department of Public Health and Environment (blue lines) and EMResource (https://emresource.juvare.com) hospital census data (red lines). Middle: Severe acute respiratory syndrome coronavirus 2 cases reported to the state health department (Colorado Department of Public Health and Environment) by report date. Bottom: reported number of severe acute respiratory syndrome coronavirus 2 tests performed per day in Colorado.
Appendix Figure 2. Structure of the age-structured susceptible-exposed-infected-recovered model. Model includes separate compartments for the 3 age groups: 0–29 years, 30–59 years, and ≥60 years. ICU, intensive care unit.
Appendix Figure 3. Collinearity plot from identifiability analysis. In this study, we have fit a social distancing parameter that switches at 4 different timepoints in the course of the hospitalization time series.