# Appendix 1:

SEIR Model Vetting Plots

Scenario: 95% Mask Use With Mandates

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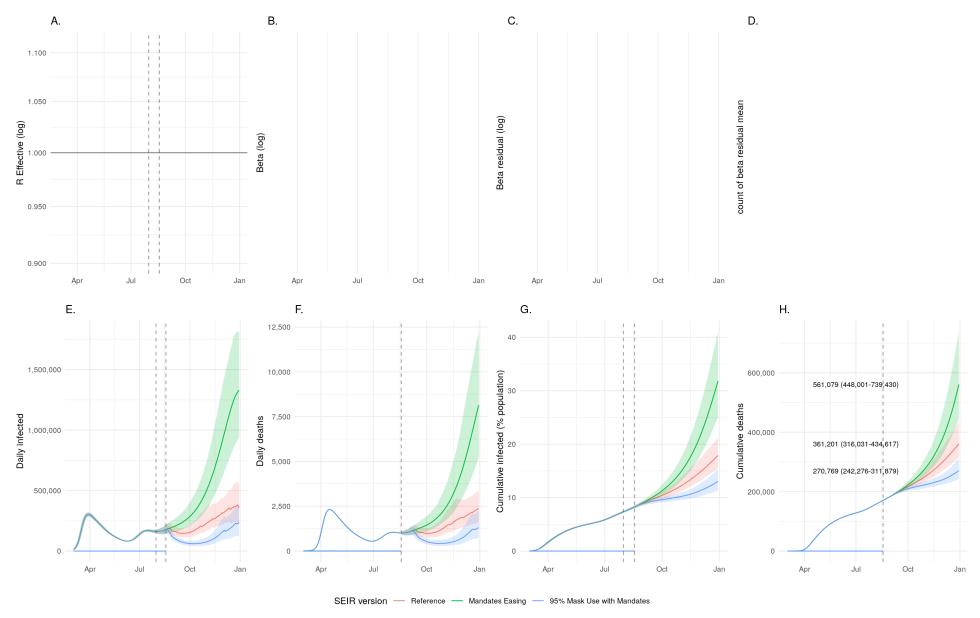
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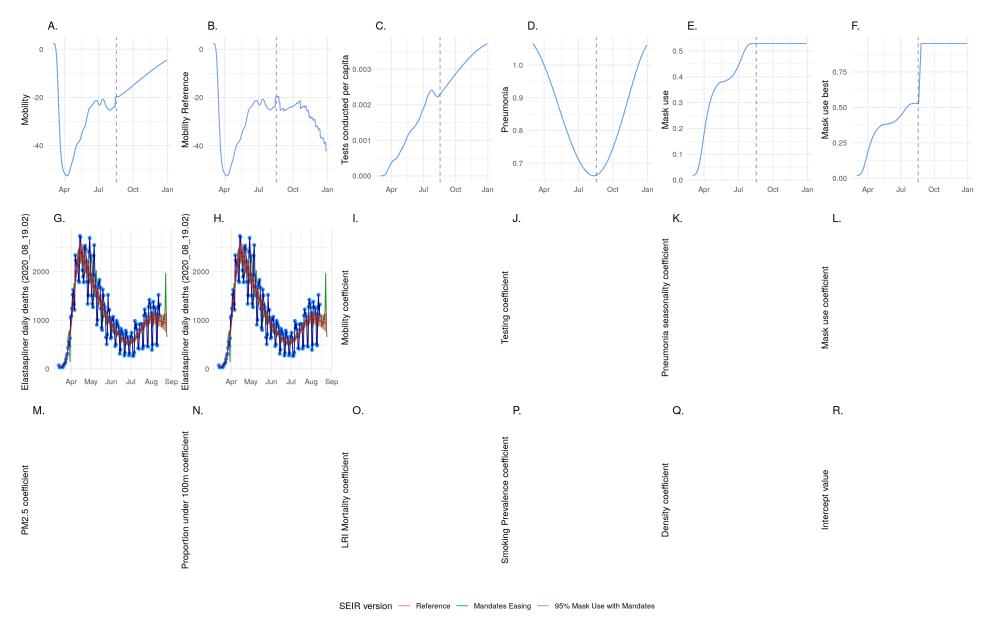
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## 1 United States of America: SEIR fit comparison



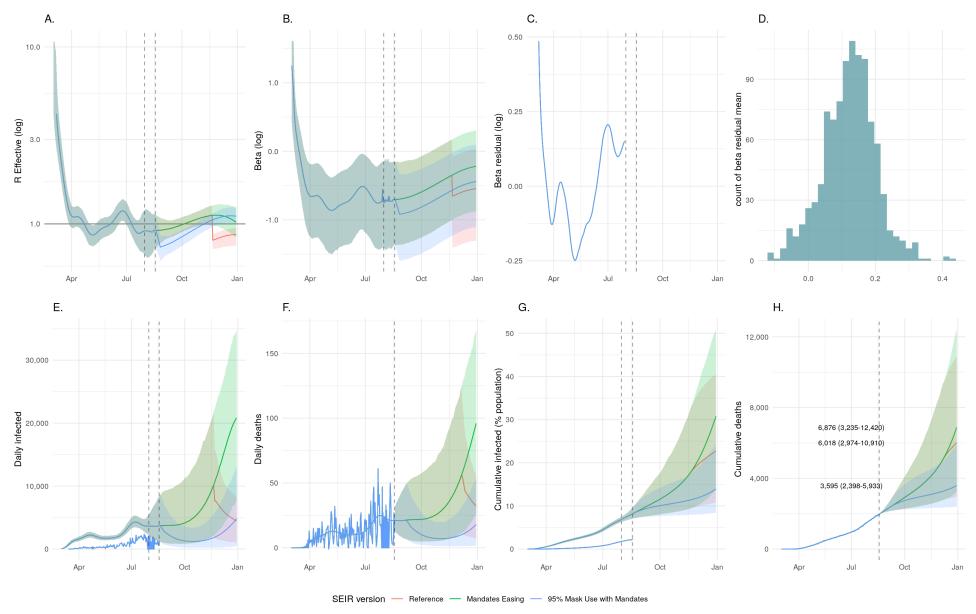
United States of America: SEIR fit comparison. Panels A-D display values that are not directly calculated for aggregate locations. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 2 United States of America: Covariate fits and regression coefficients



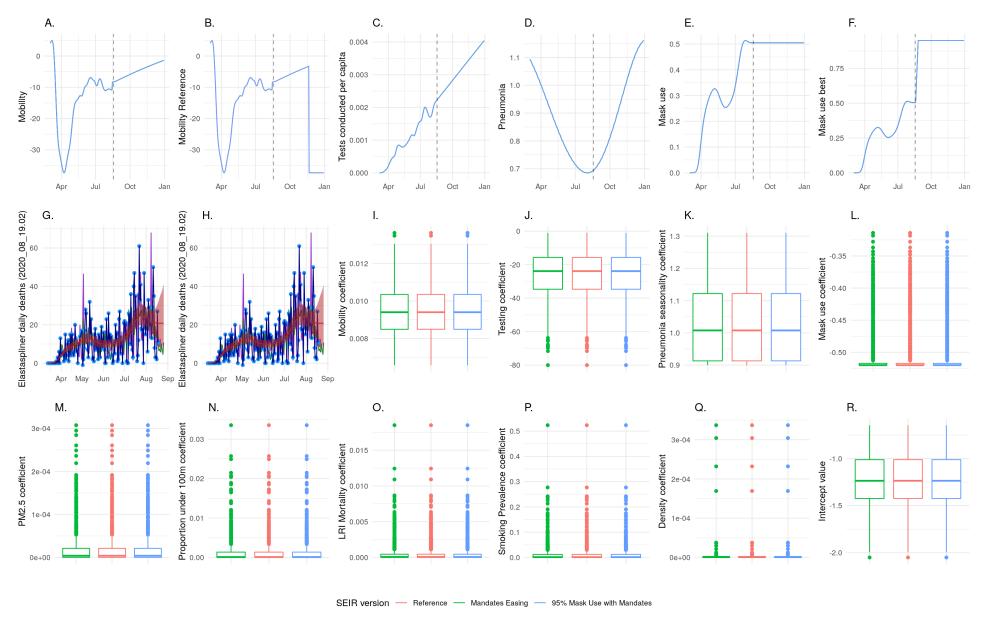
United States of America: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). Panels I-R display coefficients for a regression fit to log(beta), which is not applicable to aggregate locations.

#### 3 Alabama: SEIR fit comparison



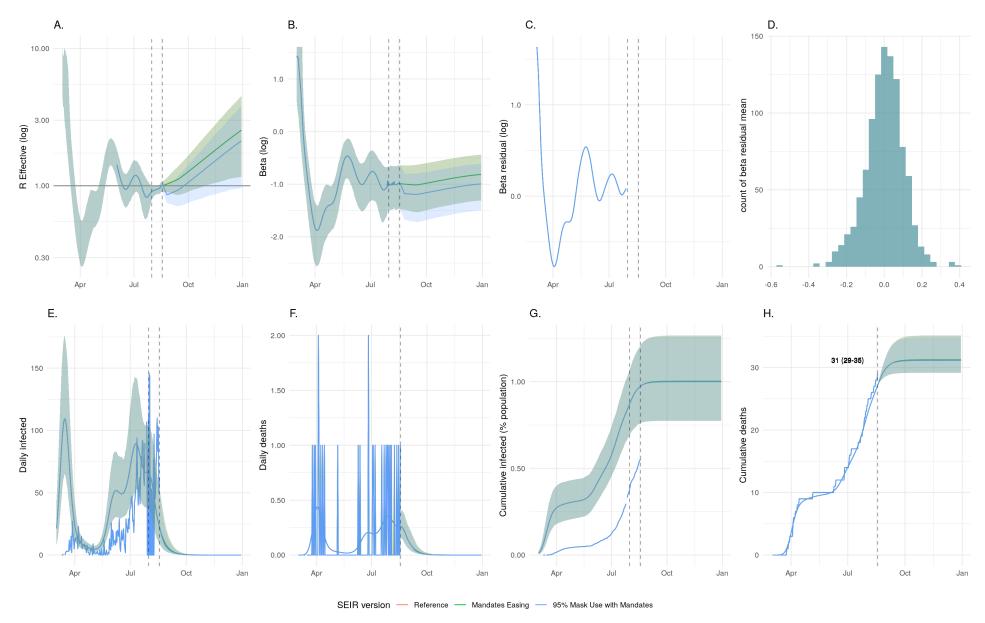
Alabama: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 4 Alabama: Covariate fits and regression coefficients



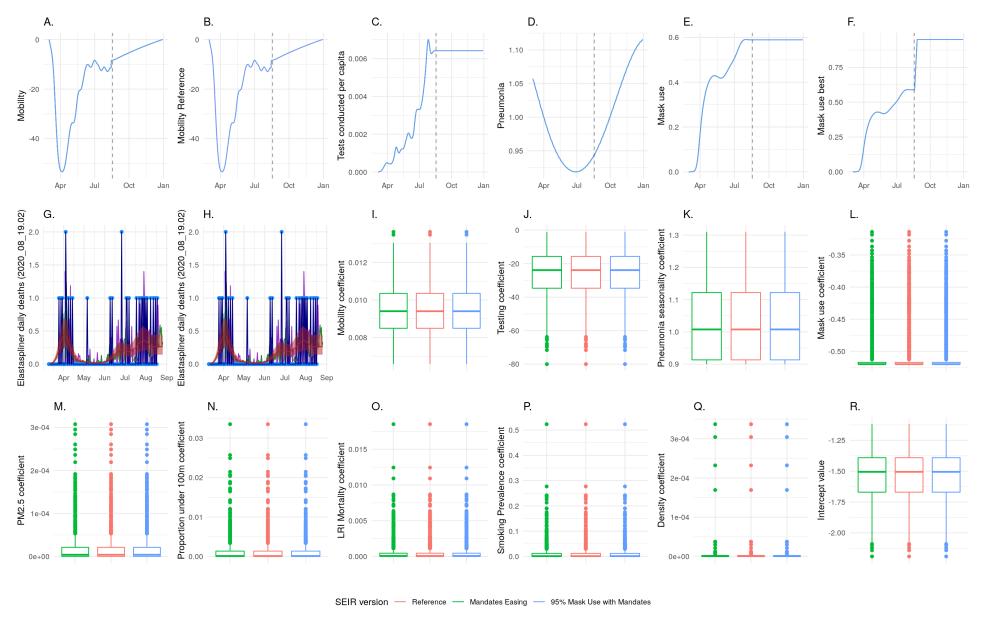
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#### 5 Alaska: SEIR fit comparison



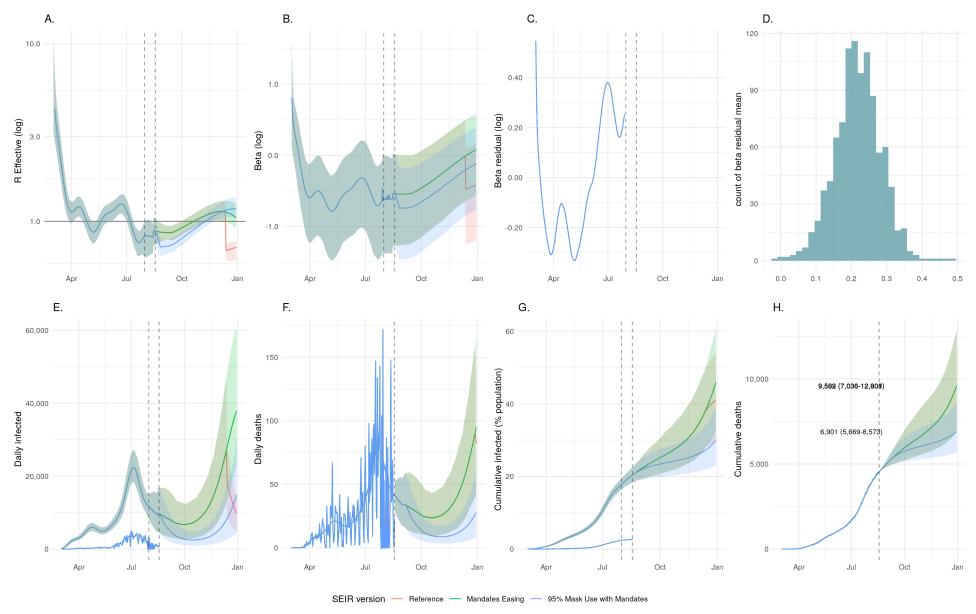
Alaska: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 6 Alaska: Covariate fits and regression coefficients



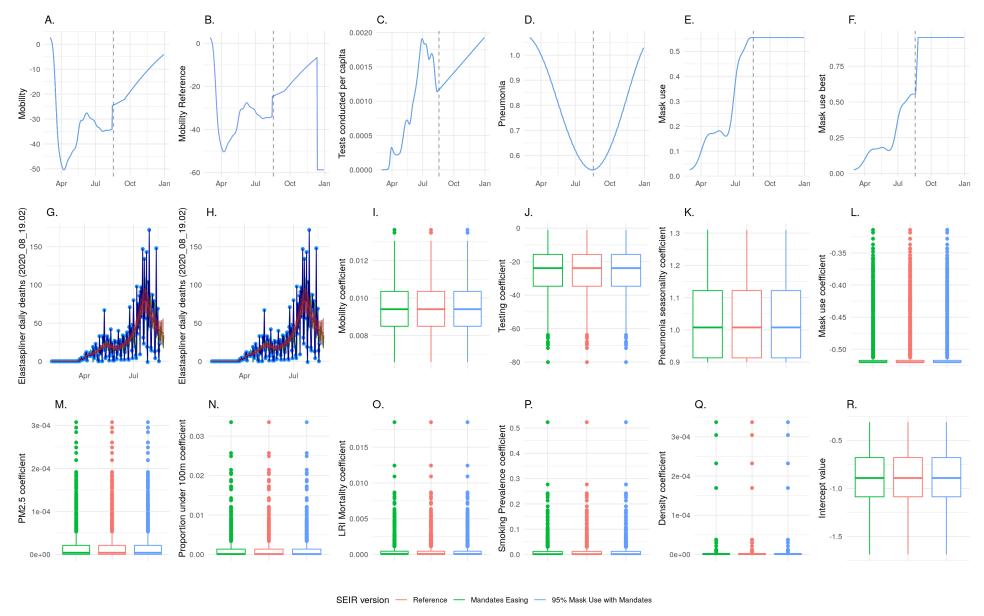
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# 7 Arizona: SEIR fit comparison



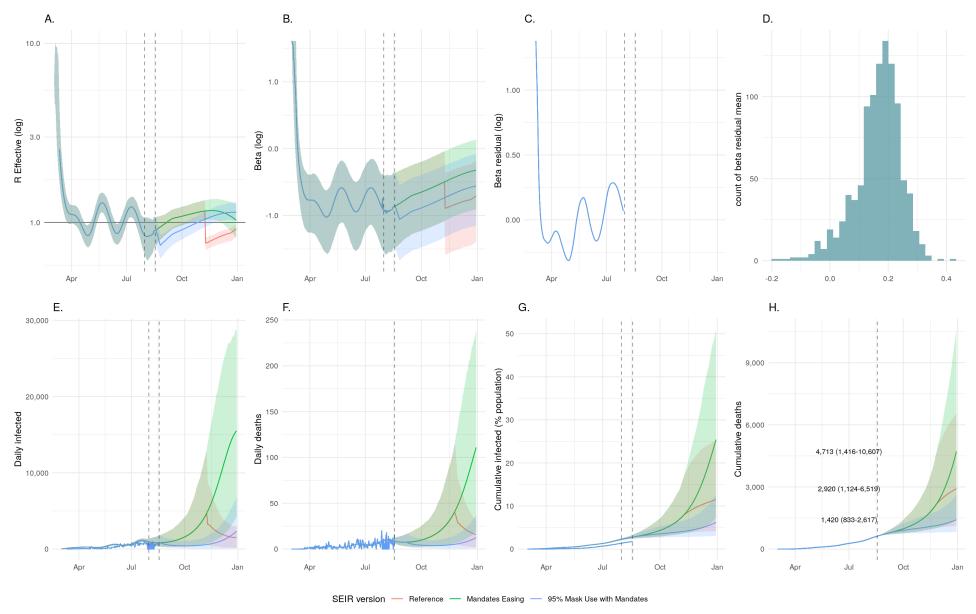
Arizona: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 8 Arizona: Covariate fits and regression coefficients



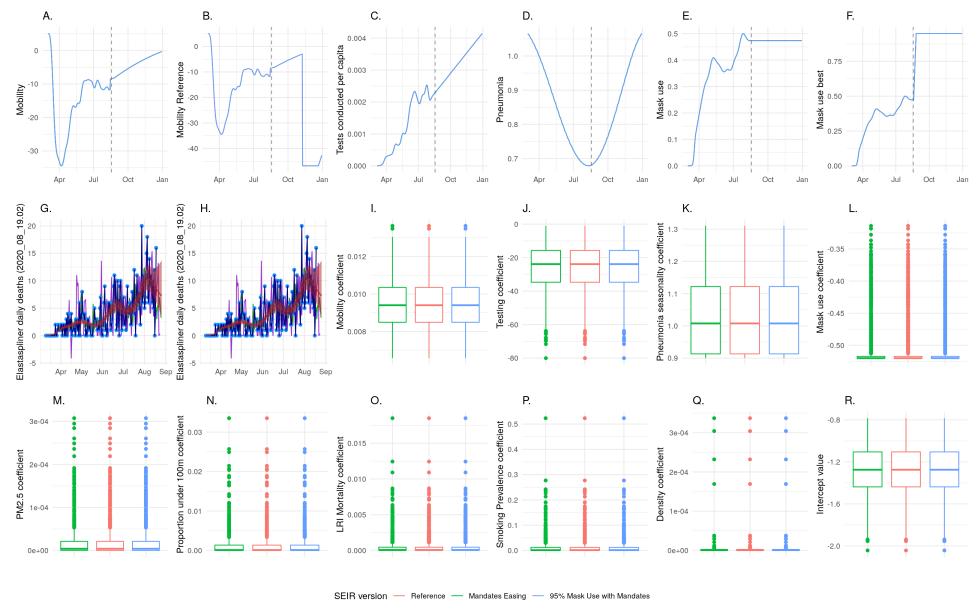
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## 9 Arkansas: SEIR fit comparison



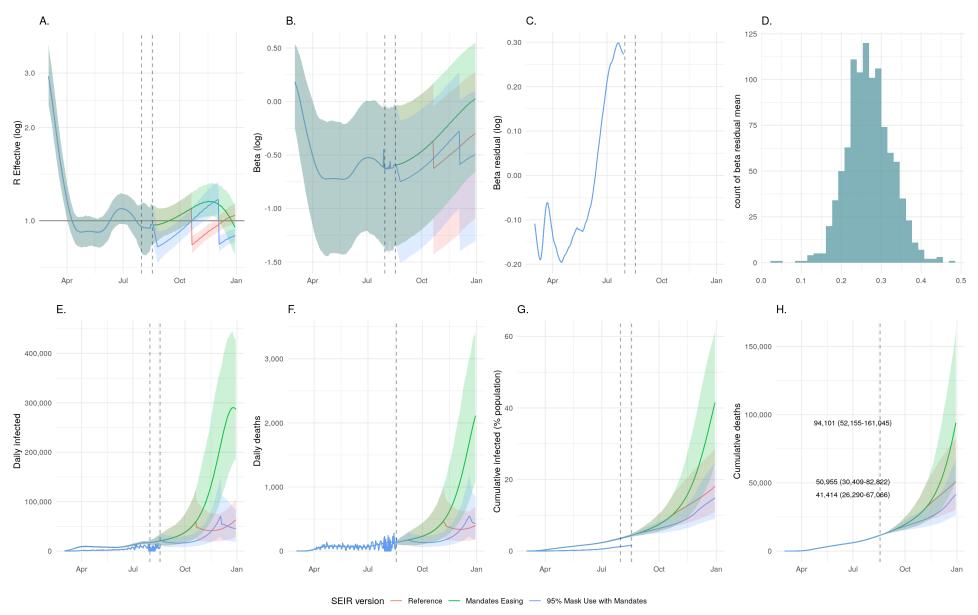
Arkansas: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 10 Arkansas: Covariate fits and regression coefficients



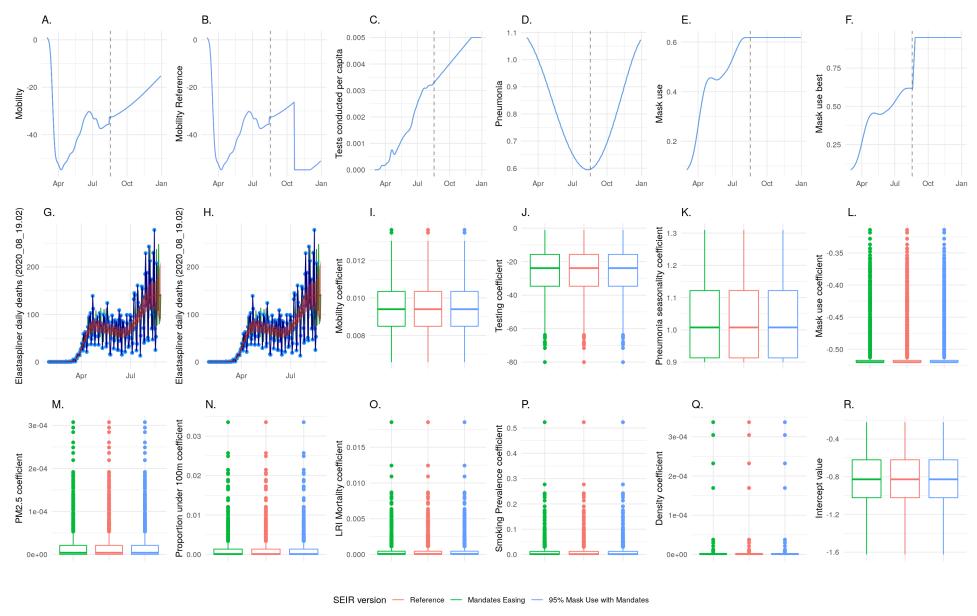
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#### 11 California: SEIR fit comparison



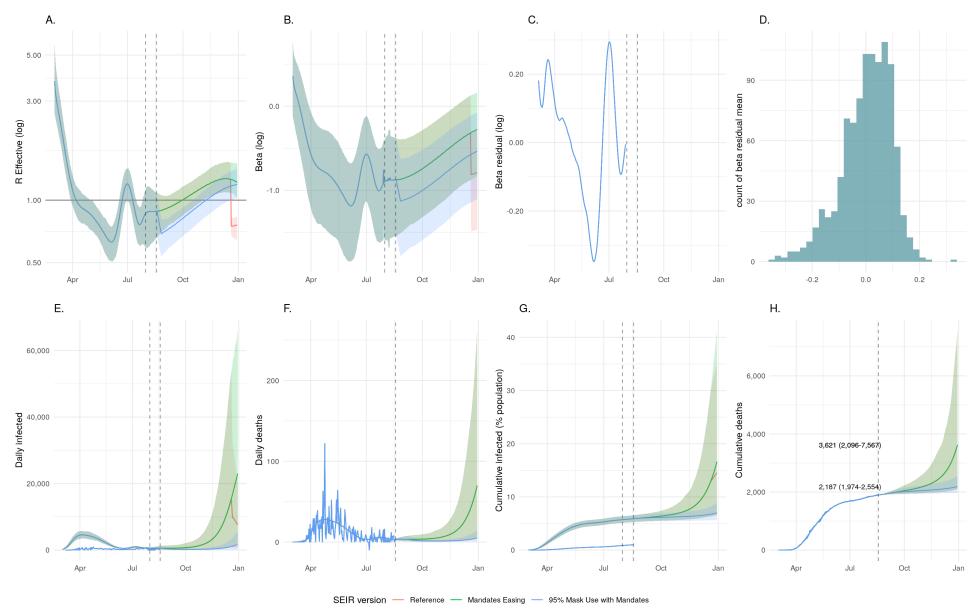
California: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 12 California: Covariate fits and regression coefficients



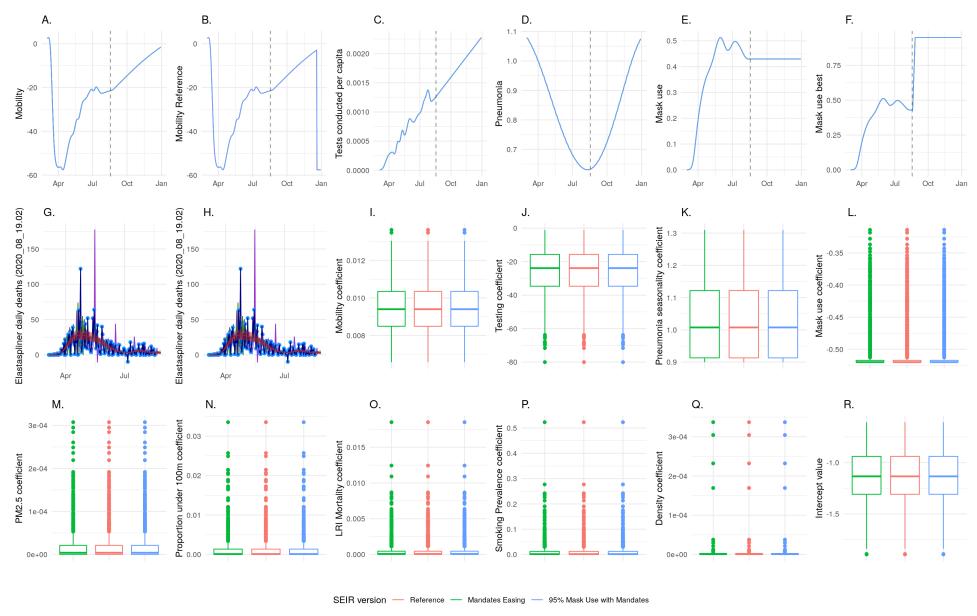
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#### 13 Colorado: SEIR fit comparison



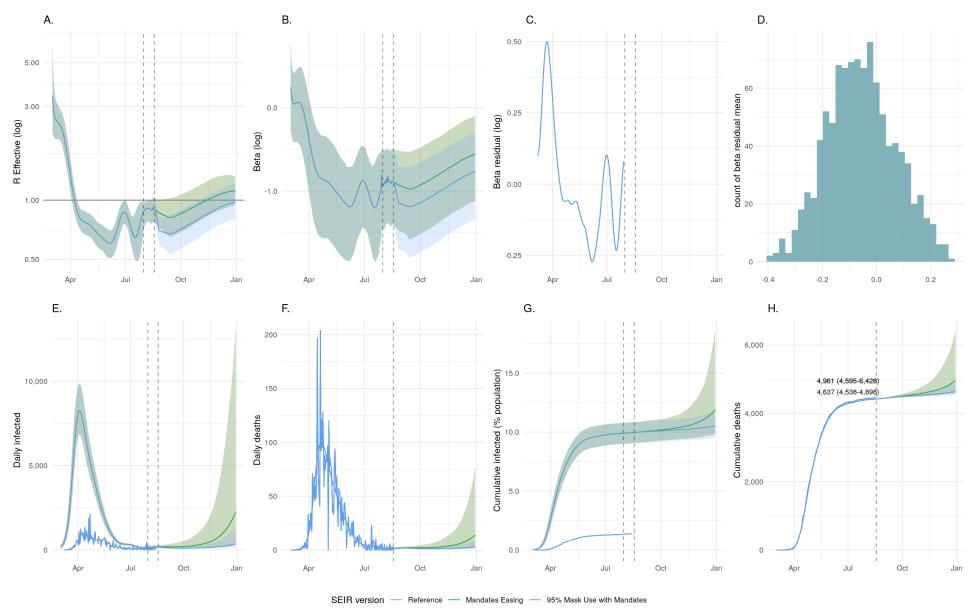
Colorado: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 14 Colorado: Covariate fits and regression coefficients



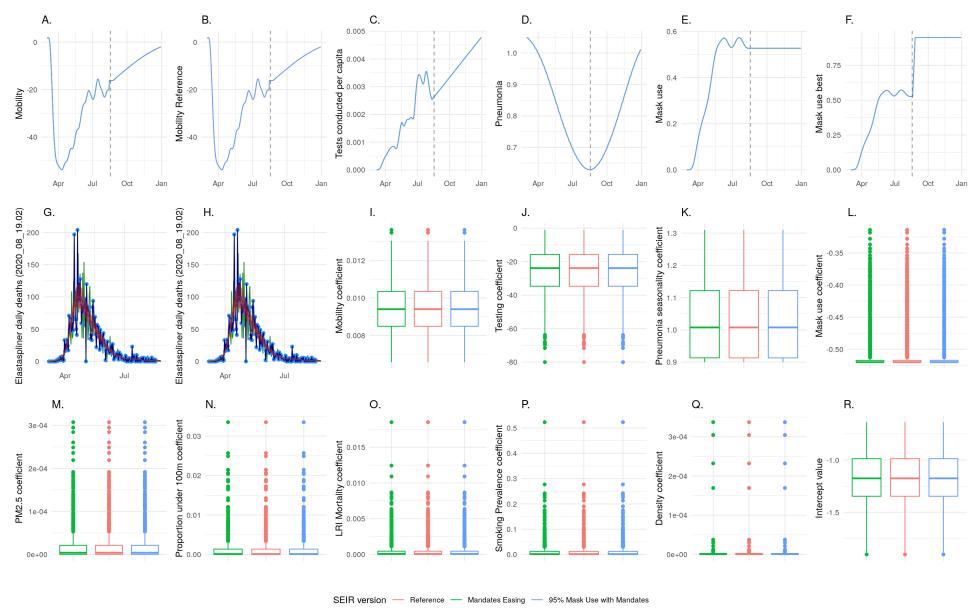
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#### 15 Connecticut: SEIR fit comparison



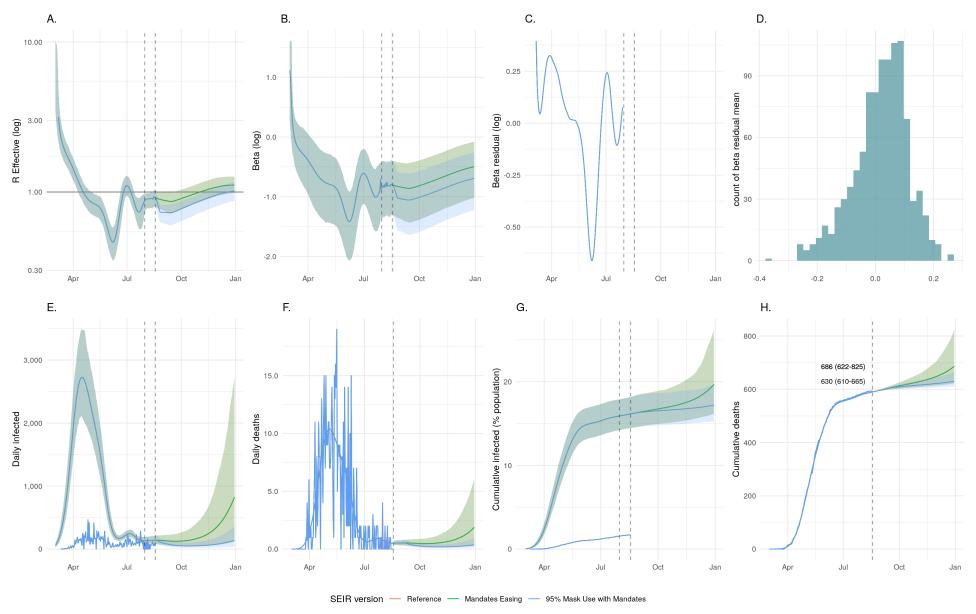
Connecticut: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 16 Connecticut: Covariate fits and regression coefficients



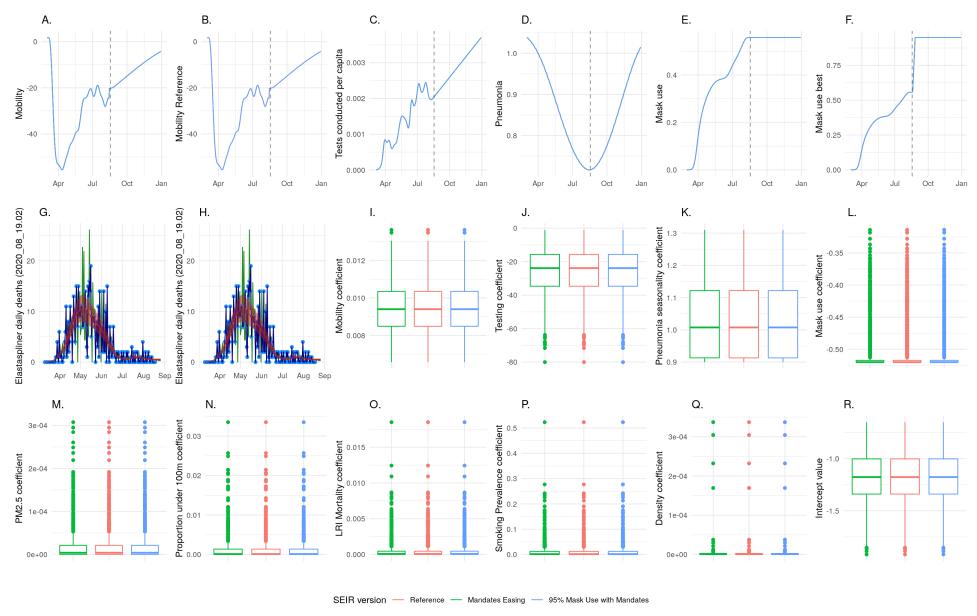
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#### 17 Delaware: SEIR fit comparison



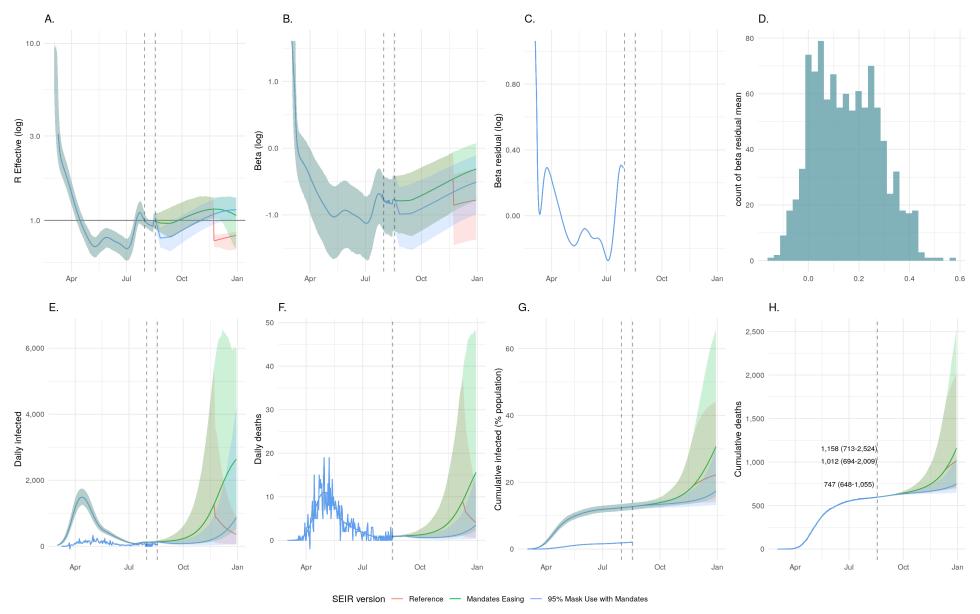
**Delaware: SEIR fit comparison. A**: predicted R effective for each model through December 31. **B**: predicted SEIR beta parameter. **C**: residual of predicted beta and the observed value calculated directly from infection data over time. **D**: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. **E**: predicted daily infections from each model through December 31. **F**: predicted daily deaths from each model through December 31. **G**: predicted cumulative infections through December 31, as a proportion of the total population. **H**: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 18 Delaware: Covariate fits and regression coefficients



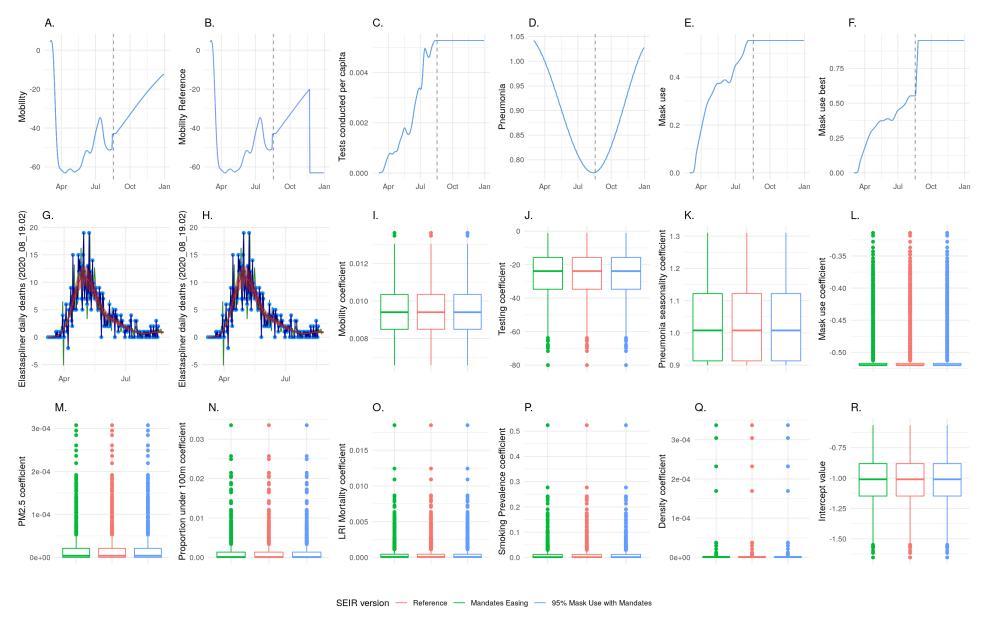
**Delaware:** Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

#### 19 District of Columbia: SEIR fit comparison



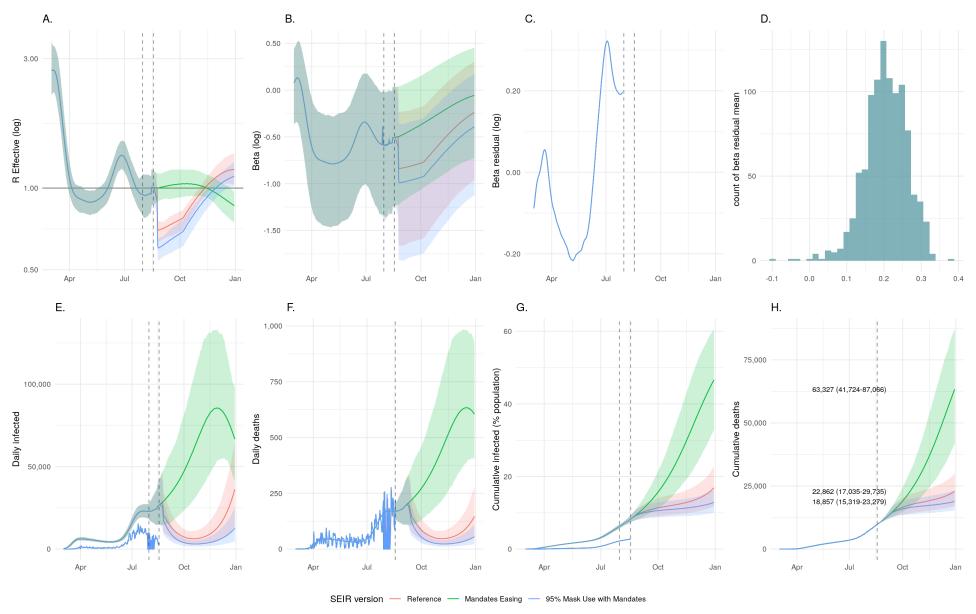
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#### 20 District of Columbia: Covariate fits and regression coefficients



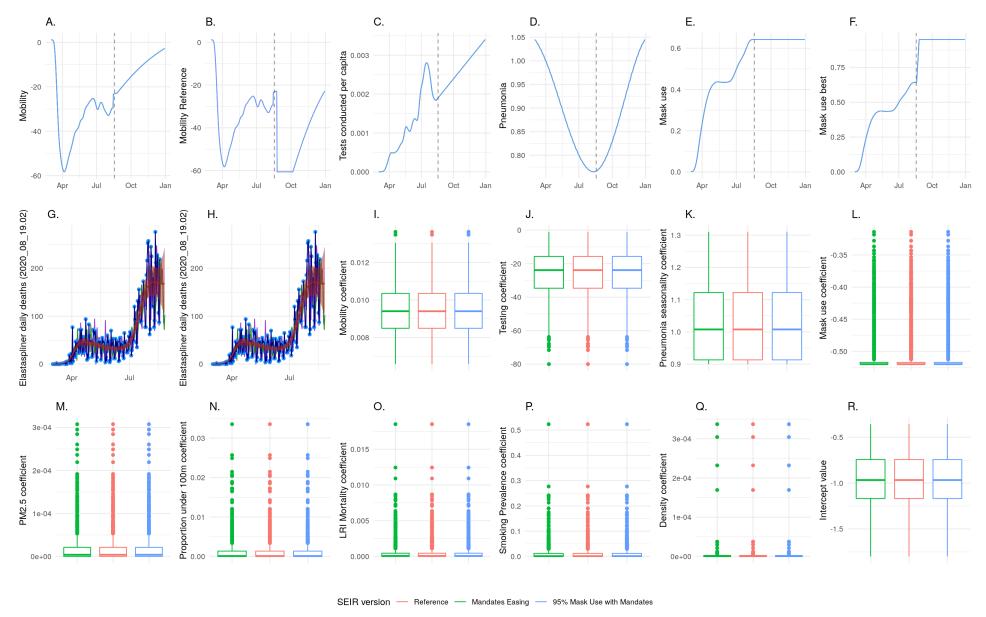
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#### 21 Florida: SEIR fit comparison



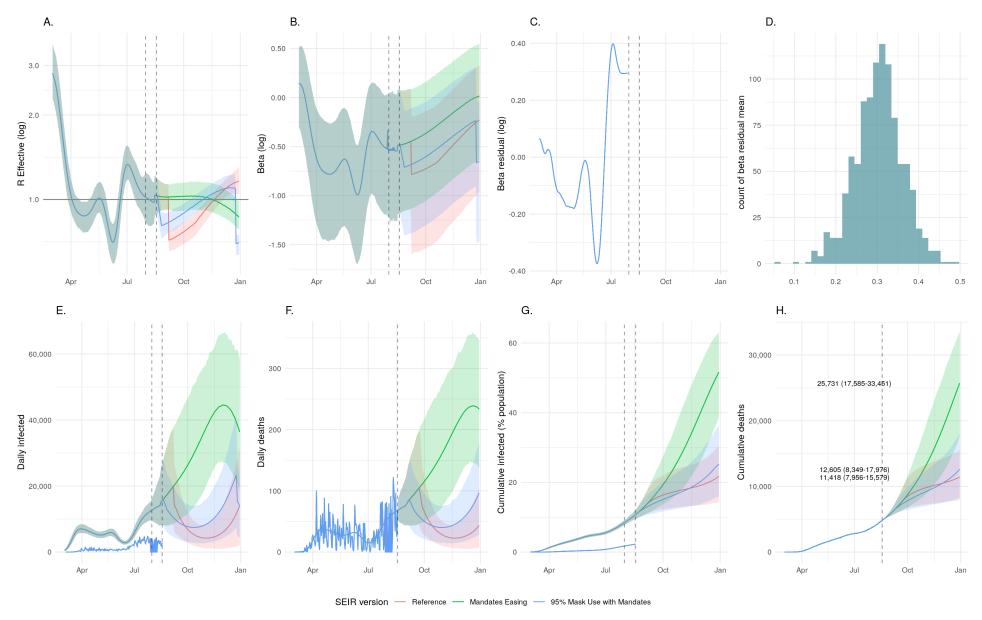
Florida: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 22 Florida: Covariate fits and regression coefficients



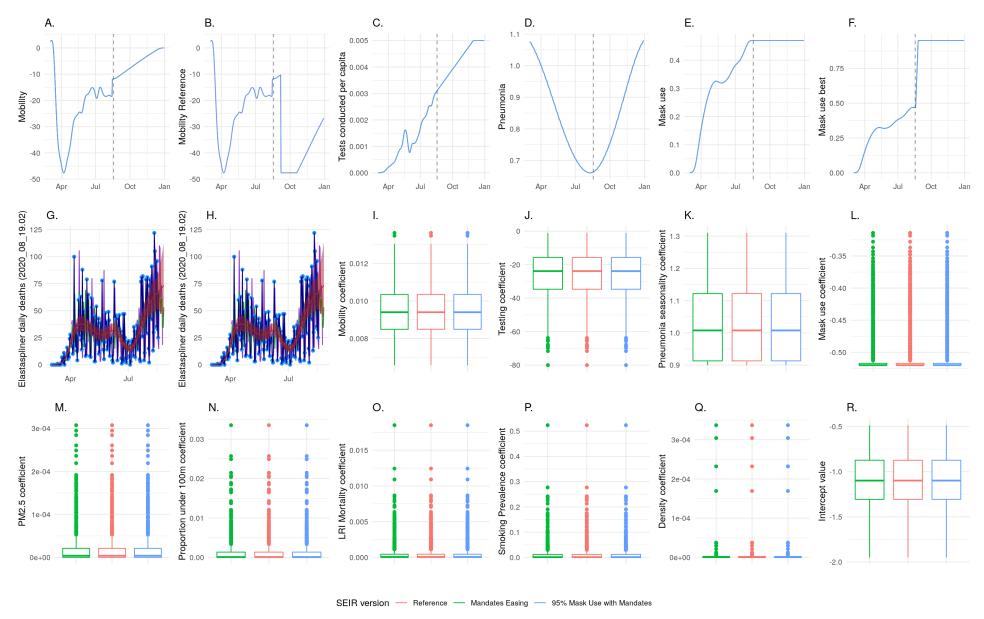
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#### 23 Georgia: SEIR fit comparison



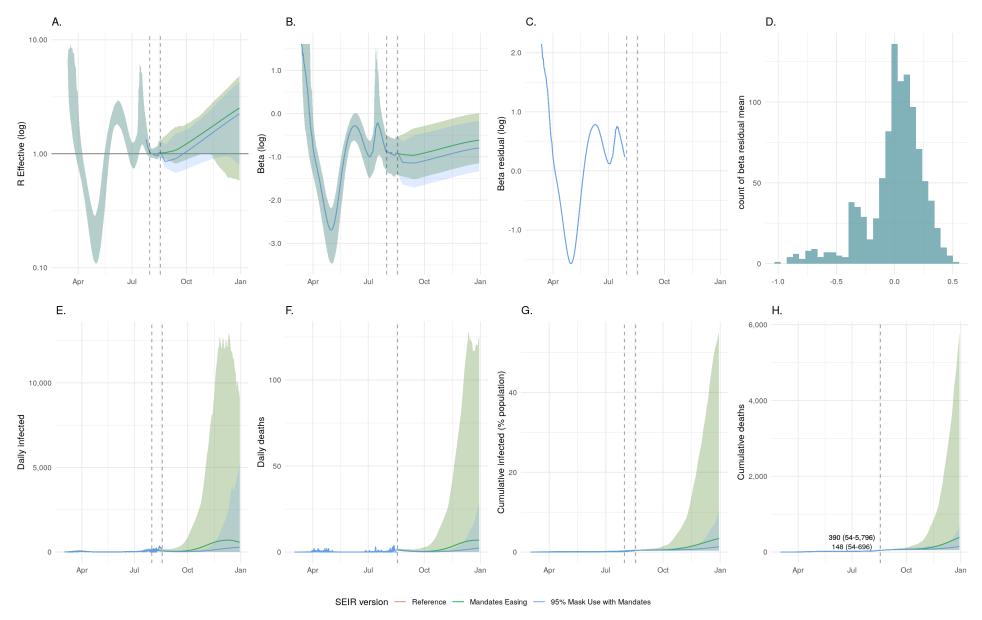
Georgia: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 24 Georgia: Covariate fits and regression coefficients



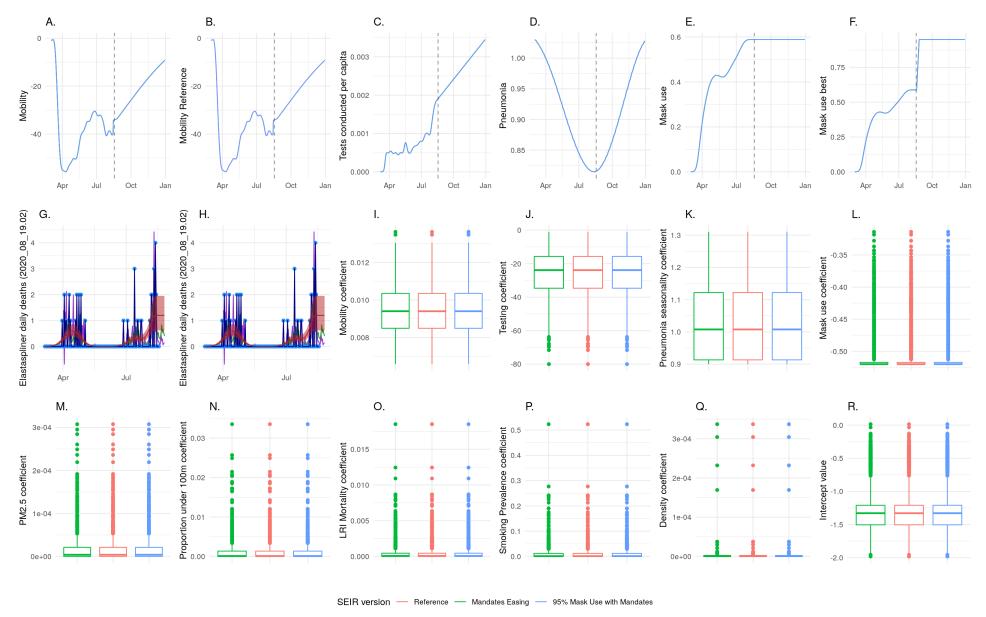
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#### 25 Hawaii: SEIR fit comparison



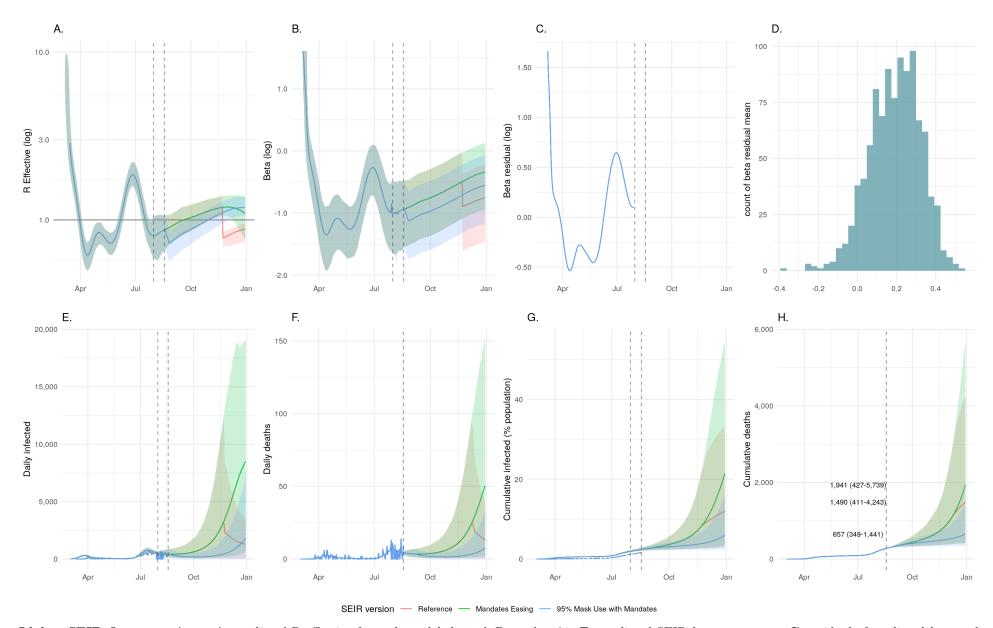
Hawaii: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 26 Hawaii: Covariate fits and regression coefficients



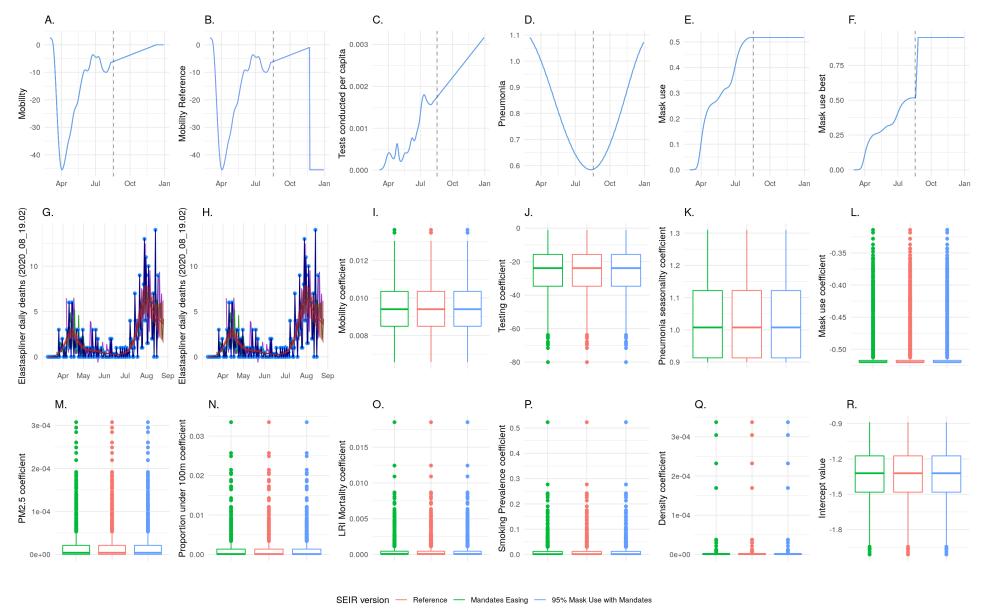
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#### 27 Idaho: SEIR fit comparison



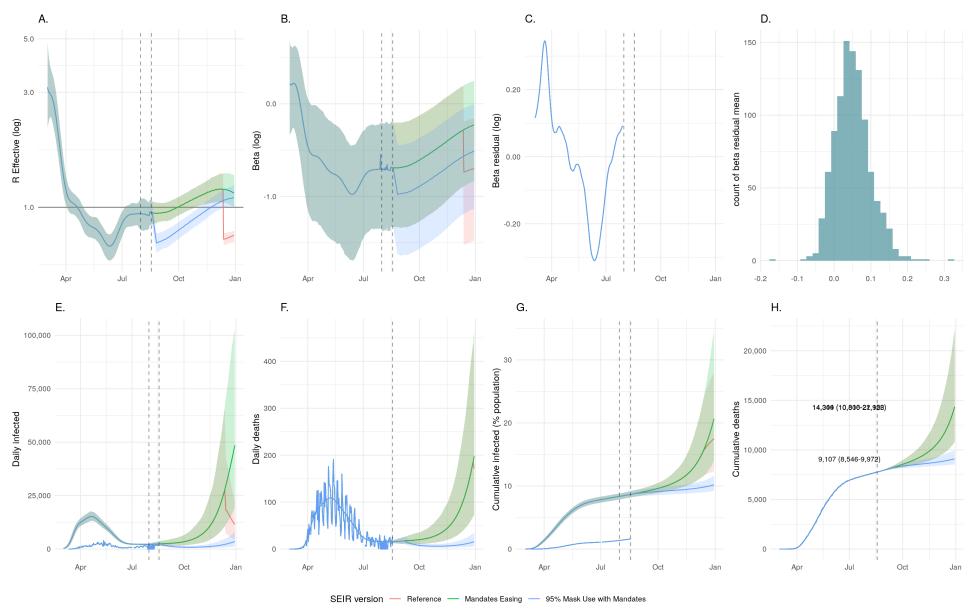
Idaho: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

# 28 Idaho: Covariate fits and regression coefficients



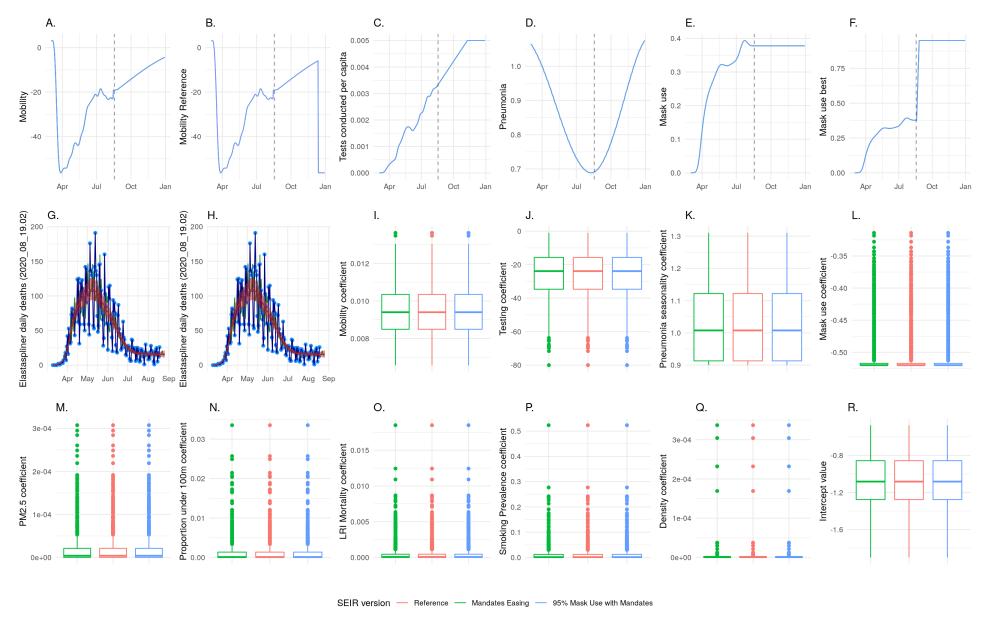
Idaho: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

#### 29 Illinois: SEIR fit comparison



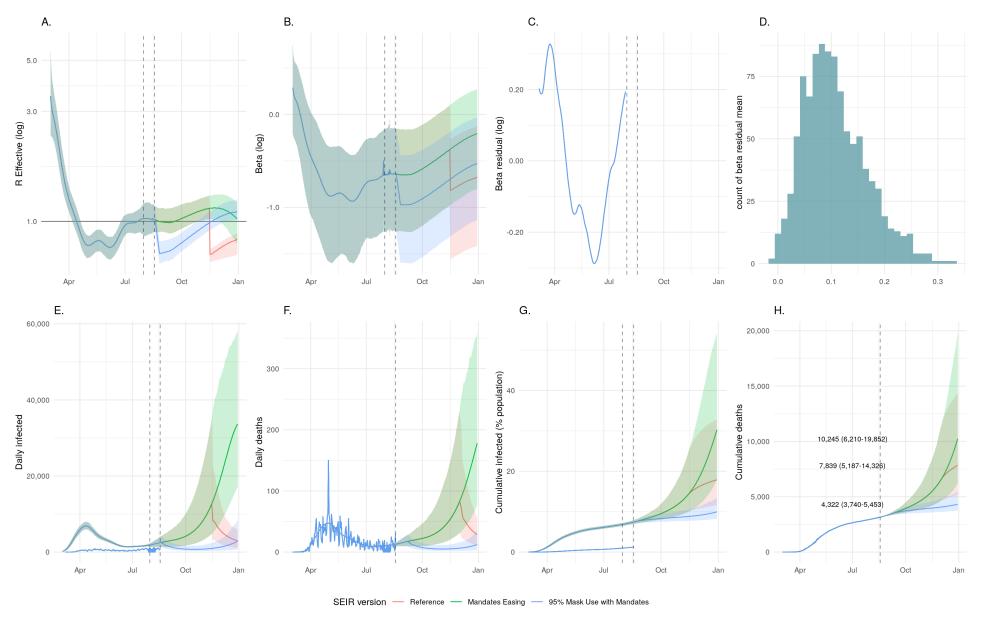
Illinois: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 30 Illinois: Covariate fits and regression coefficients



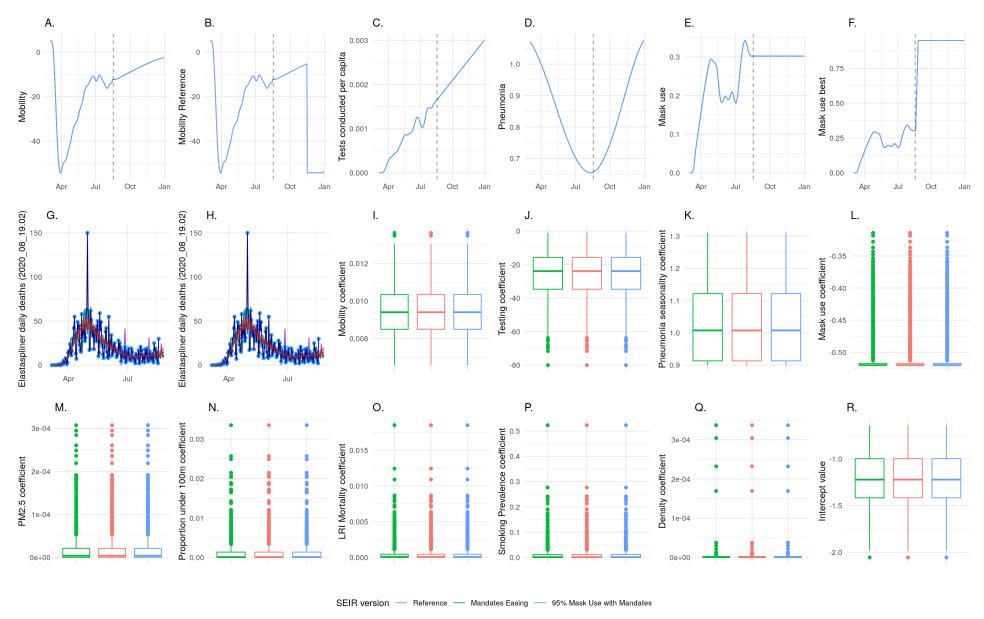
Illinois: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 31 Indiana: SEIR fit comparison



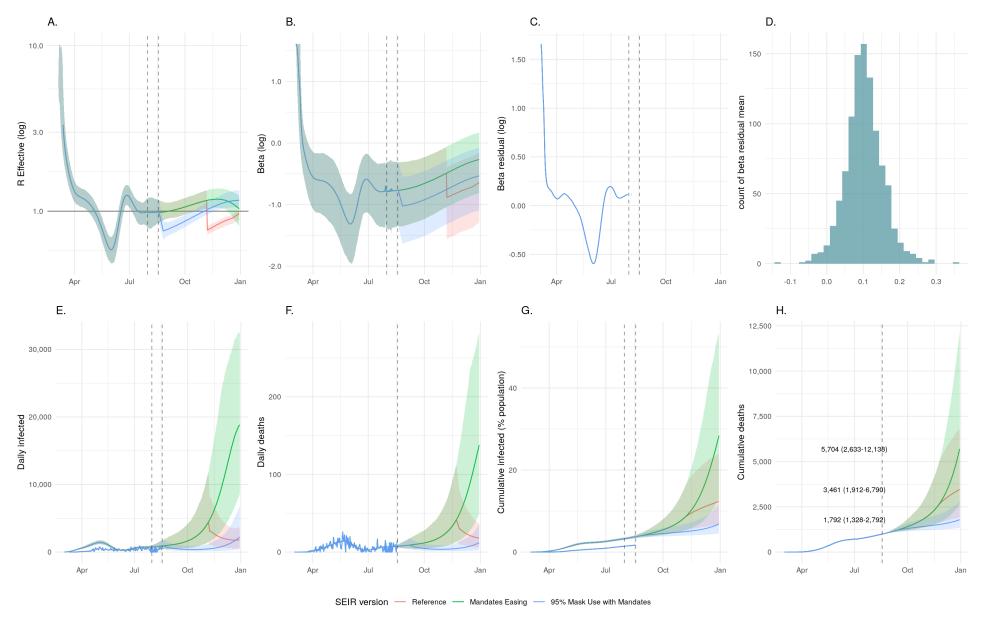
Indiana: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 32 Indiana: Covariate fits and regression coefficients



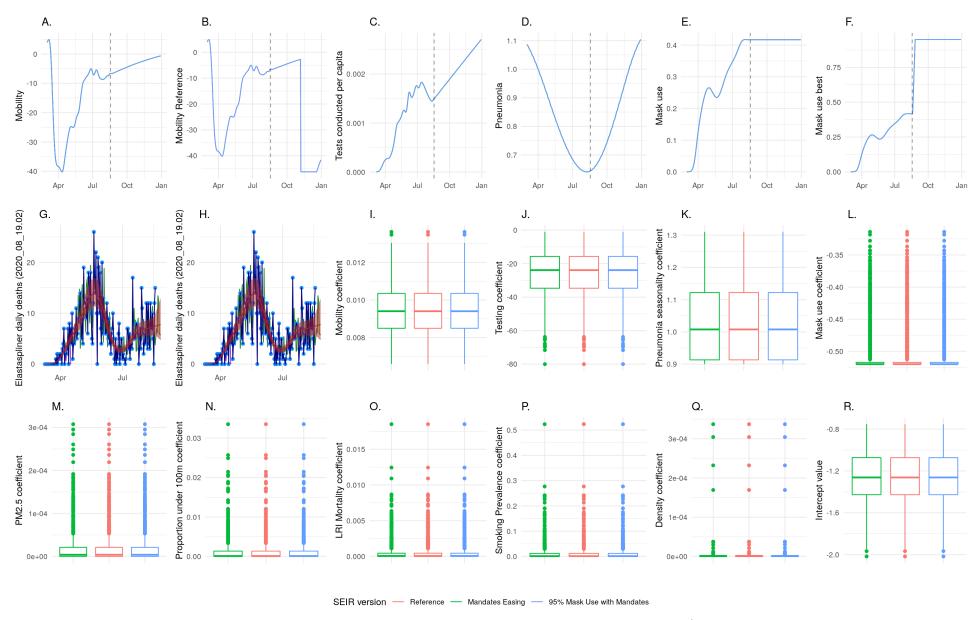
Indiana: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 33 Iowa: SEIR fit comparison



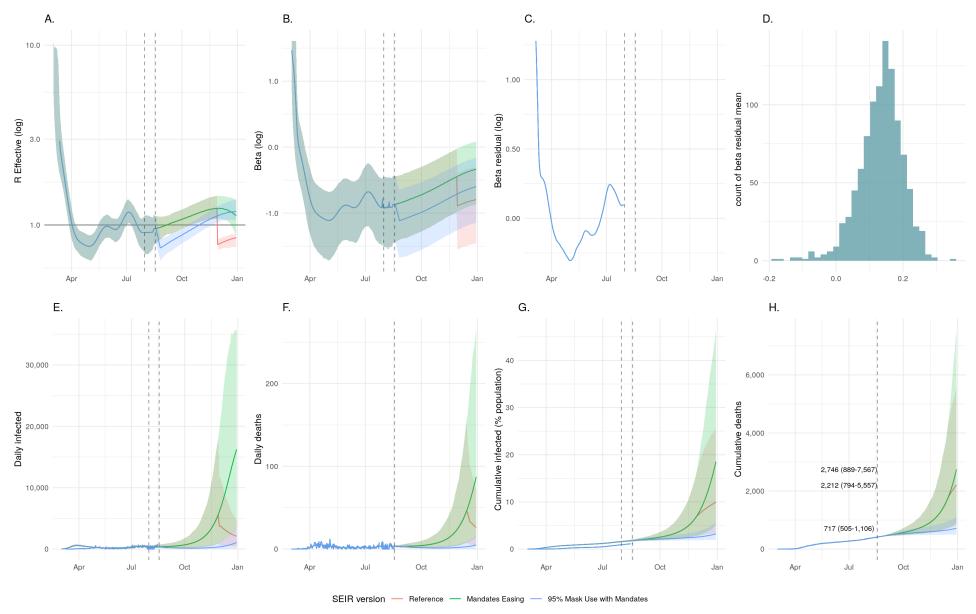
Iowa: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 34 Iowa: Covariate fits and regression coefficients



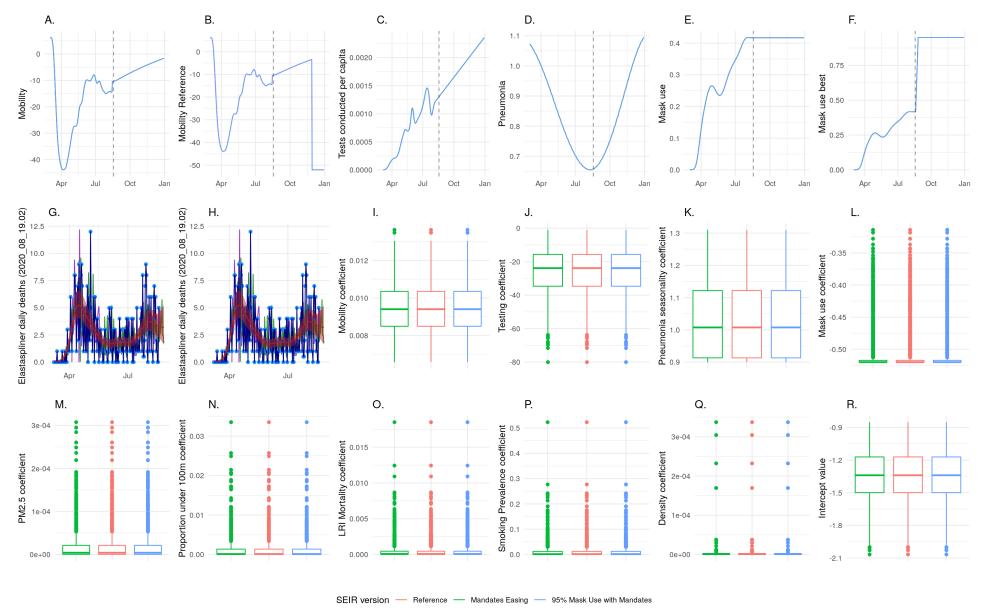
Iowa: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 35 Kansas: SEIR fit comparison



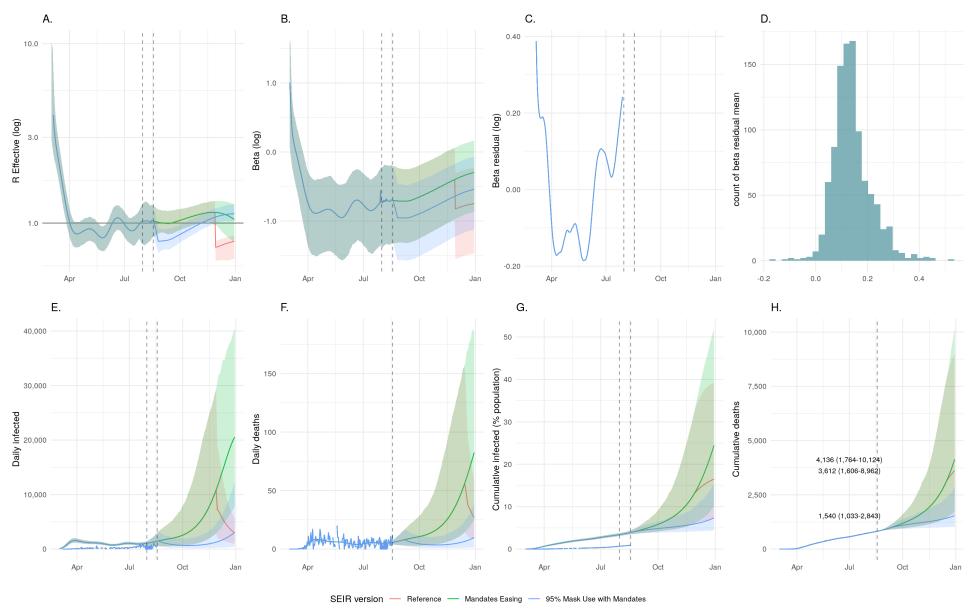
Kansas: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 36 Kansas: Covariate fits and regression coefficients



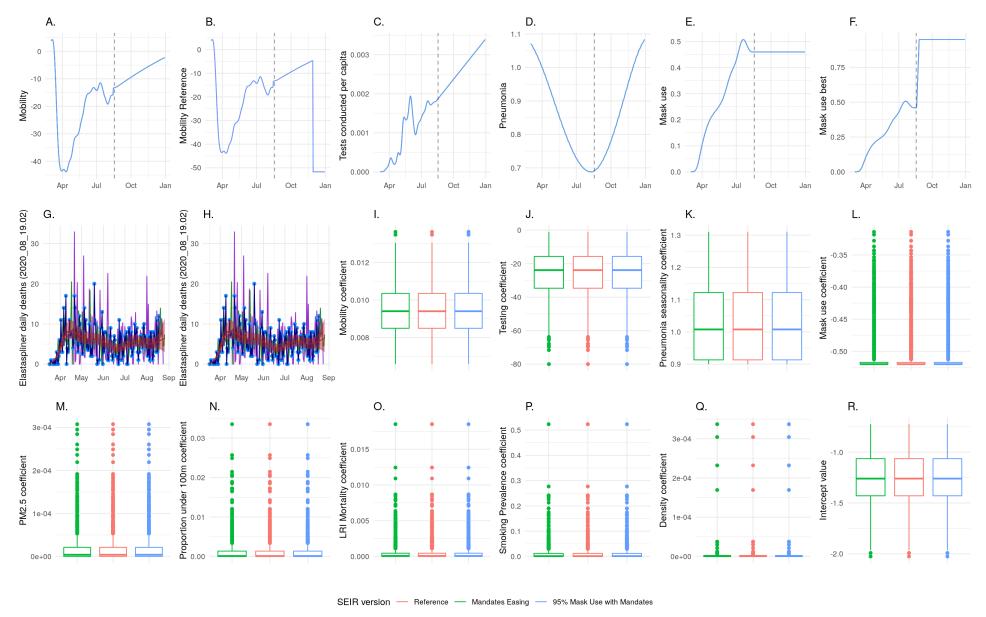
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# 37 Kentucky: SEIR fit comparison



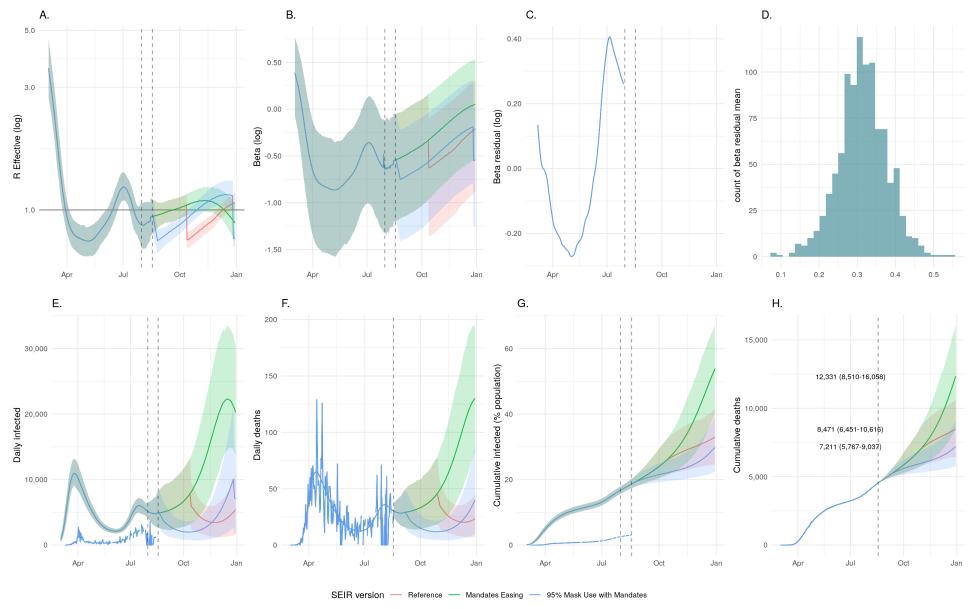
**Kentucky: SEIR fit comparison. A**: predicted R effective for each model through December 31. **B**: predicted SEIR beta parameter. **C**: residual of predicted beta and the observed value calculated directly from infection data over time. **D**: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. **E**: predicted daily infections from each model through December 31. **F**: predicted daily deaths from each model through December 31. **G**: predicted cumulative infections through December 31, as a proportion of the total population. **H**: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 38 Kentucky: Covariate fits and regression coefficients



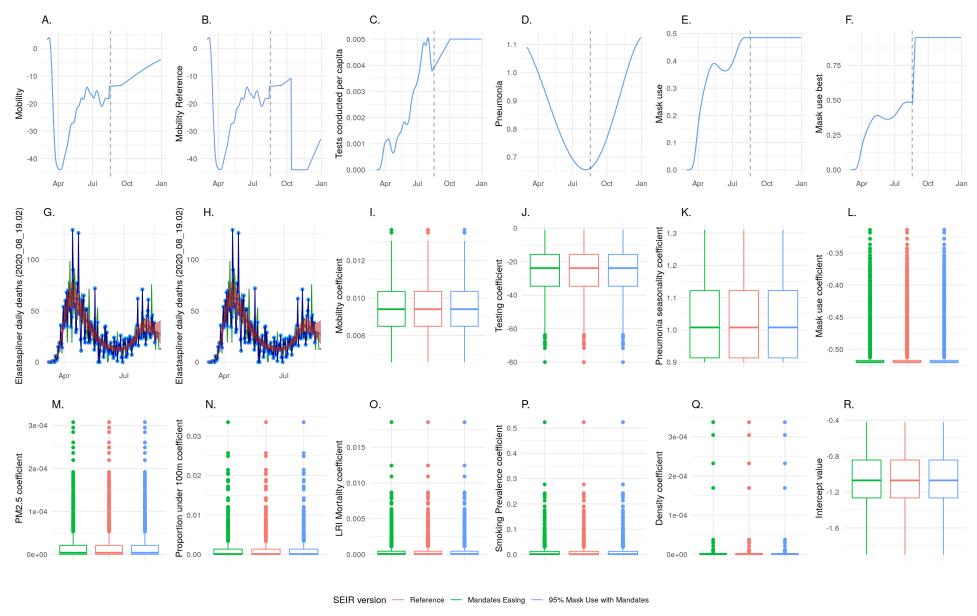
Kentucky: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 39 Louisiana: SEIR fit comparison



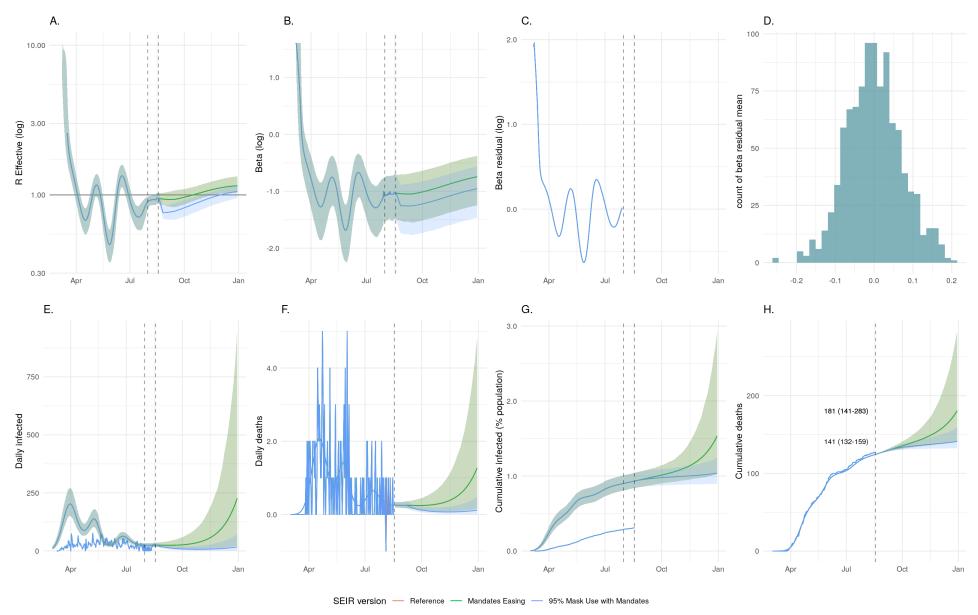
Louisiana: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 40 Louisiana: Covariate fits and regression coefficients



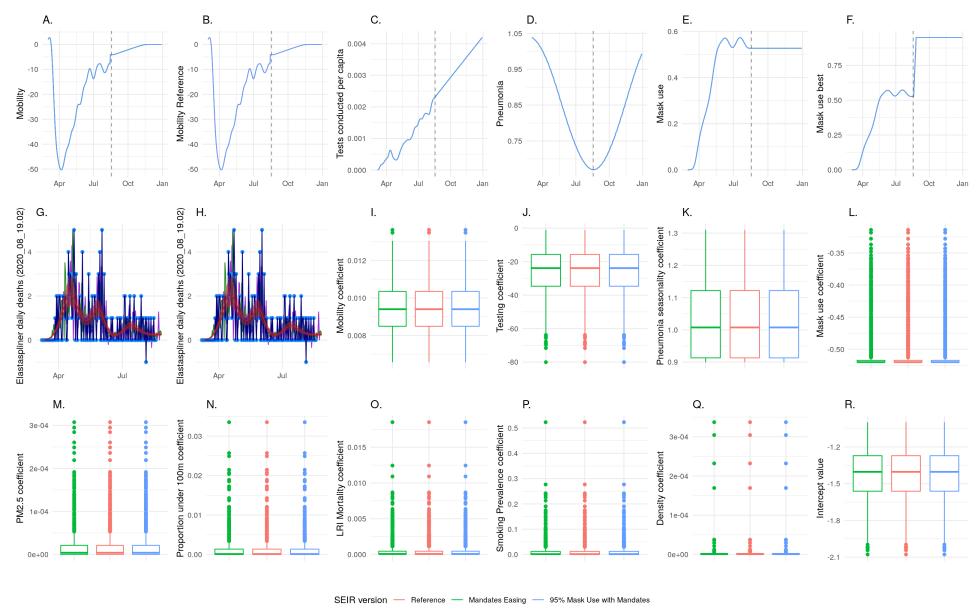
Louisiana: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 41 Maine: SEIR fit comparison



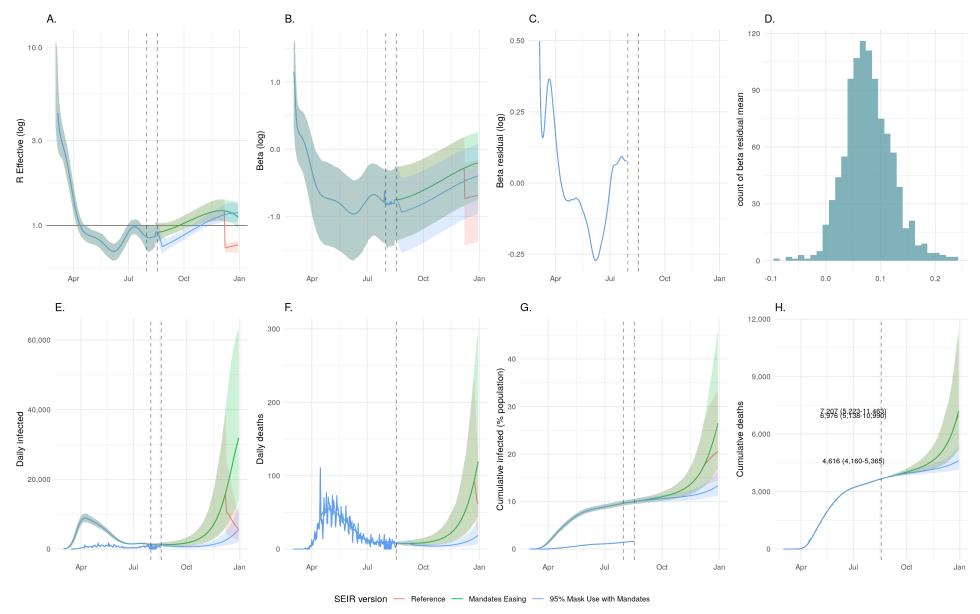
Maine: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 42 Maine: Covariate fits and regression coefficients



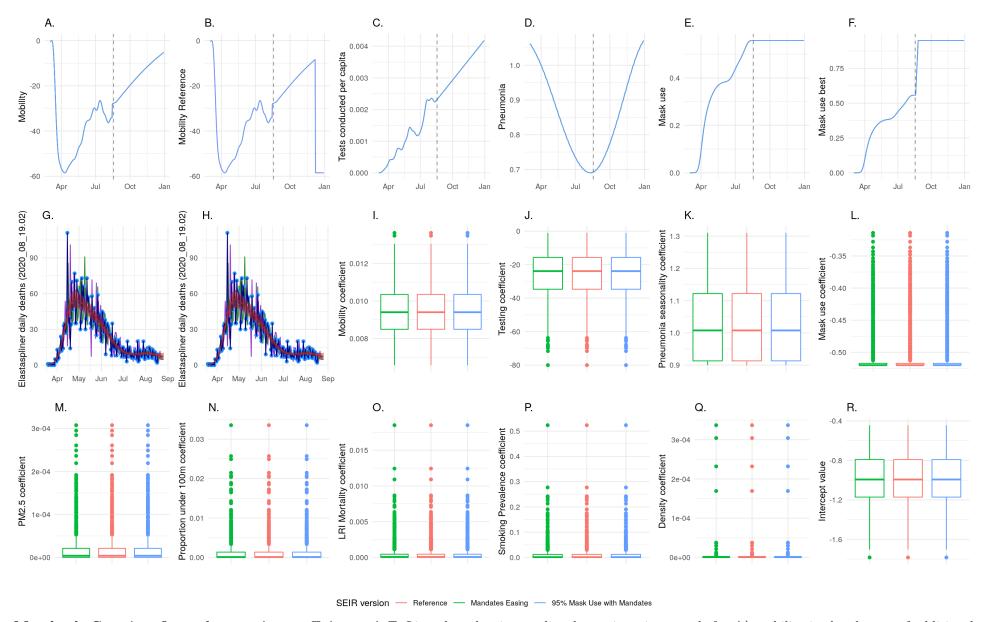
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# 43 Maryland: SEIR fit comparison



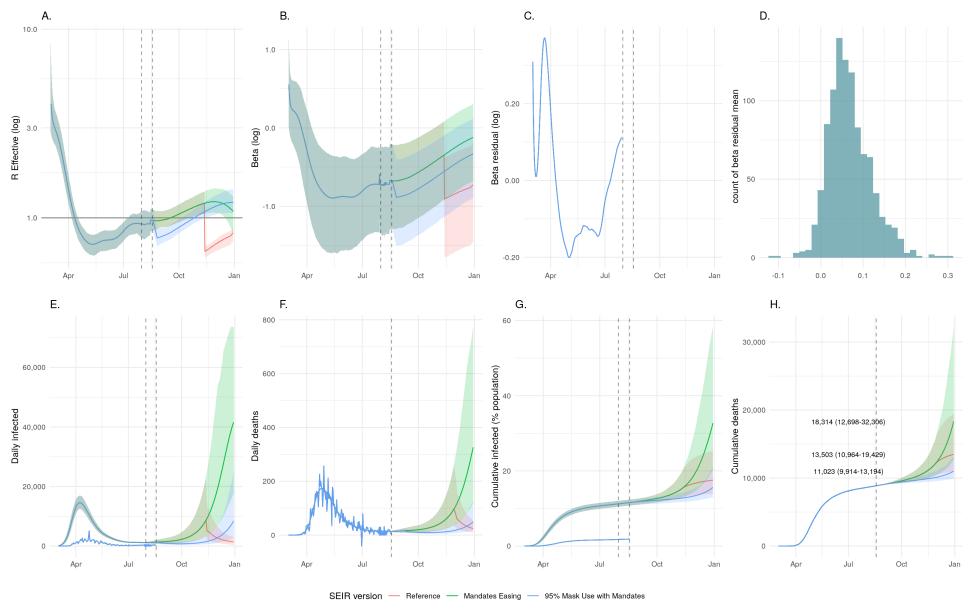
Maryland: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 44 Maryland: Covariate fits and regression coefficients



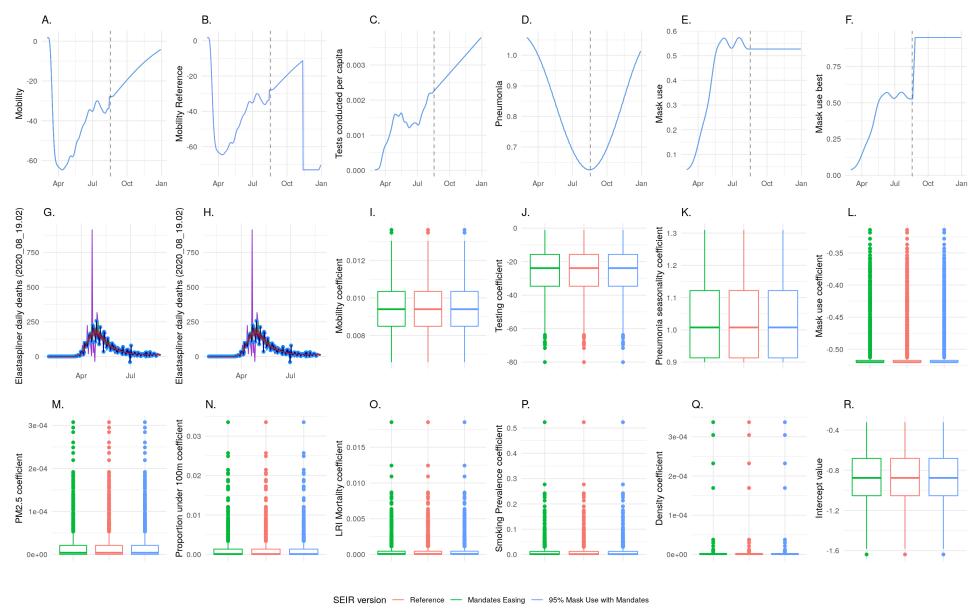
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### 45 Massachusetts: SEIR fit comparison



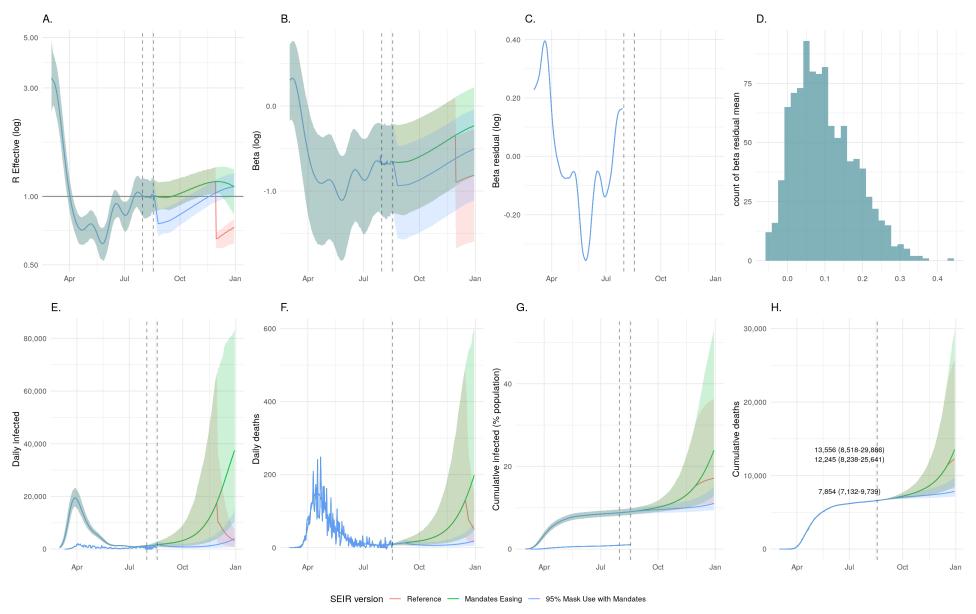
Massachusetts: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 46 Massachusetts: Covariate fits and regression coefficients



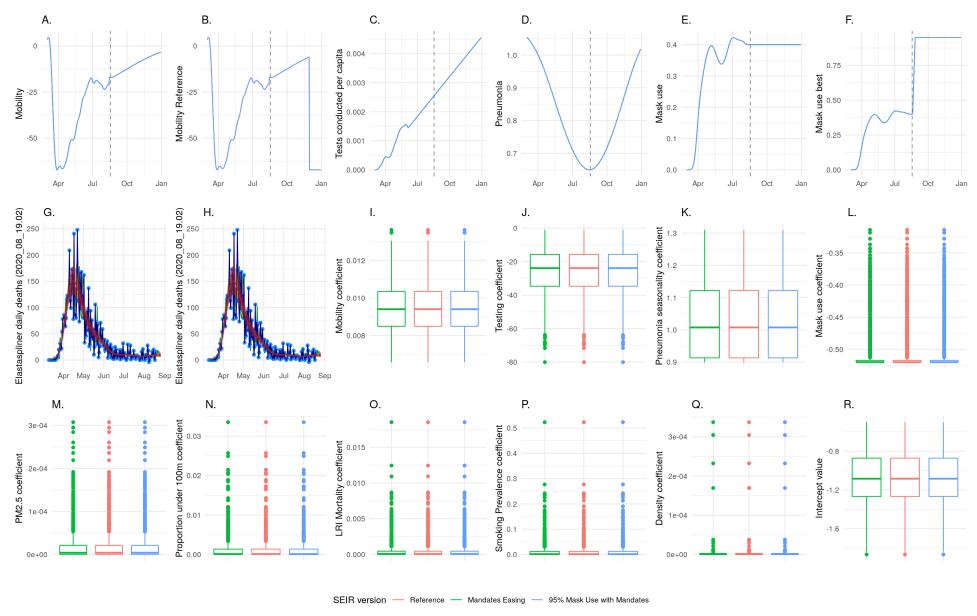
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# 47 Michigan: SEIR fit comparison



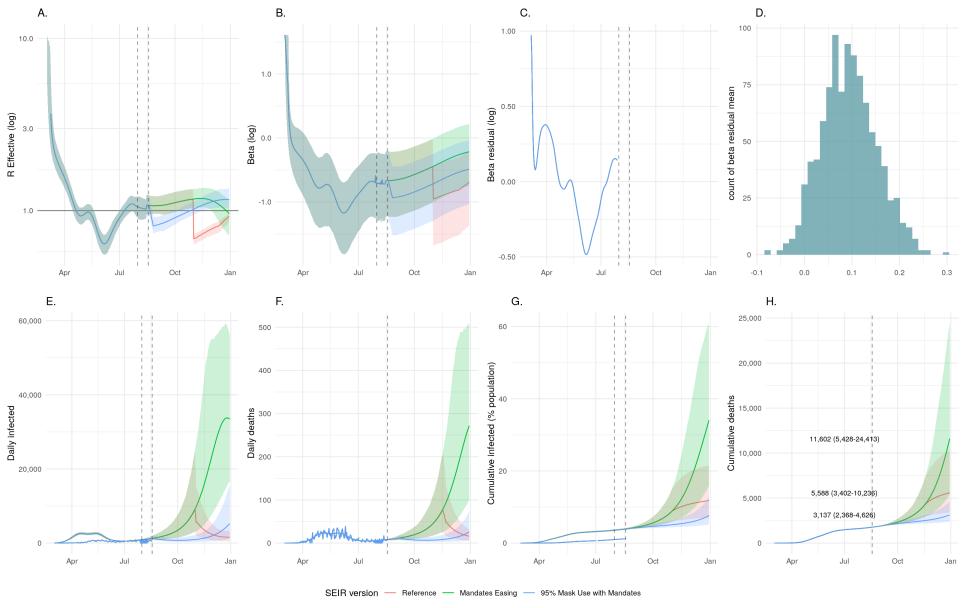
Michigan: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 48 Michigan: Covariate fits and regression coefficients



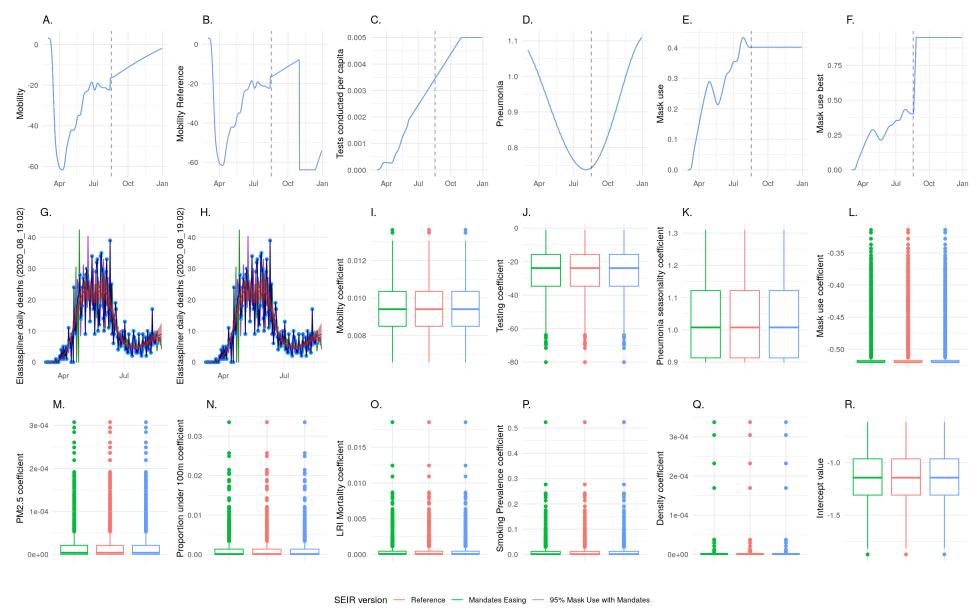
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### 49 Minnesota: SEIR fit comparison



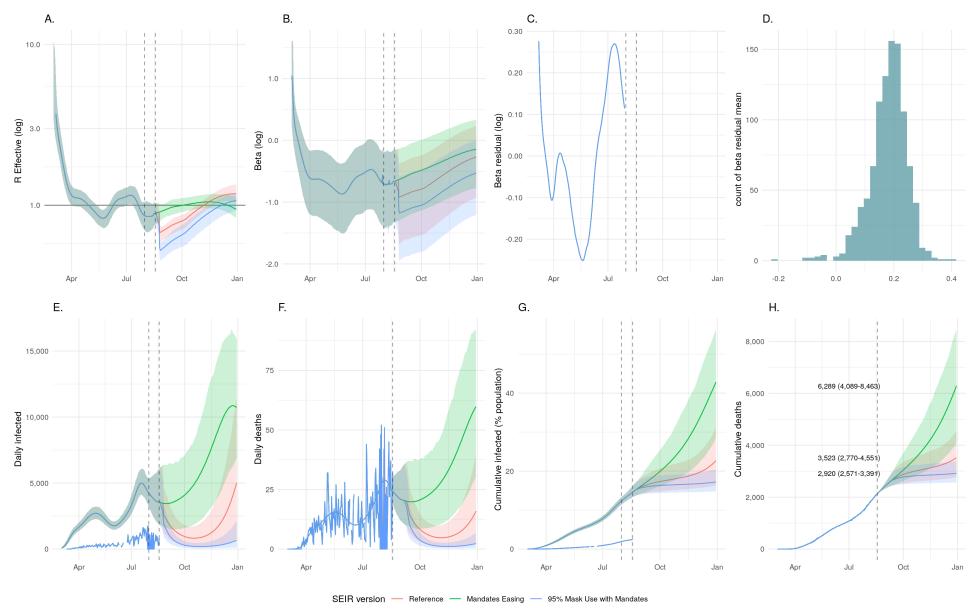
Minnesota: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 50 Minnesota: Covariate fits and regression coefficients



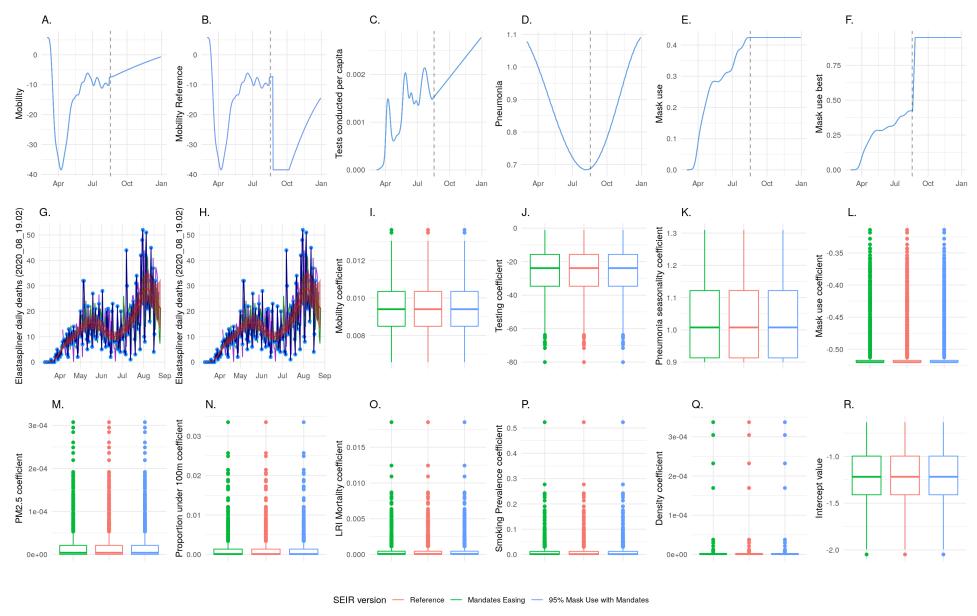
Minnesota: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 51 Mississippi: SEIR fit comparison



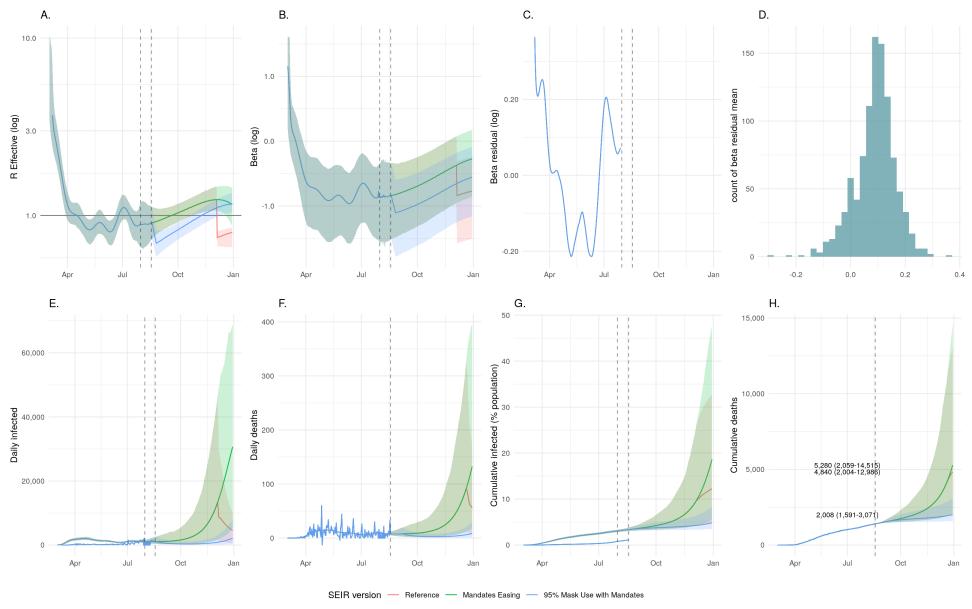
Mississippi: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 52 Mississippi: Covariate fits and regression coefficients



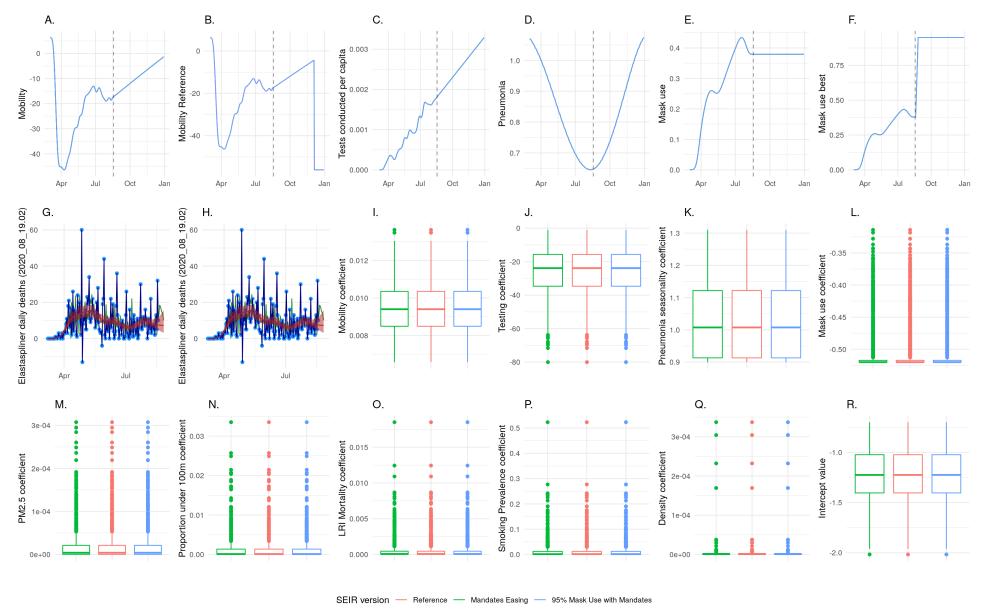
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### 53 Missouri: SEIR fit comparison



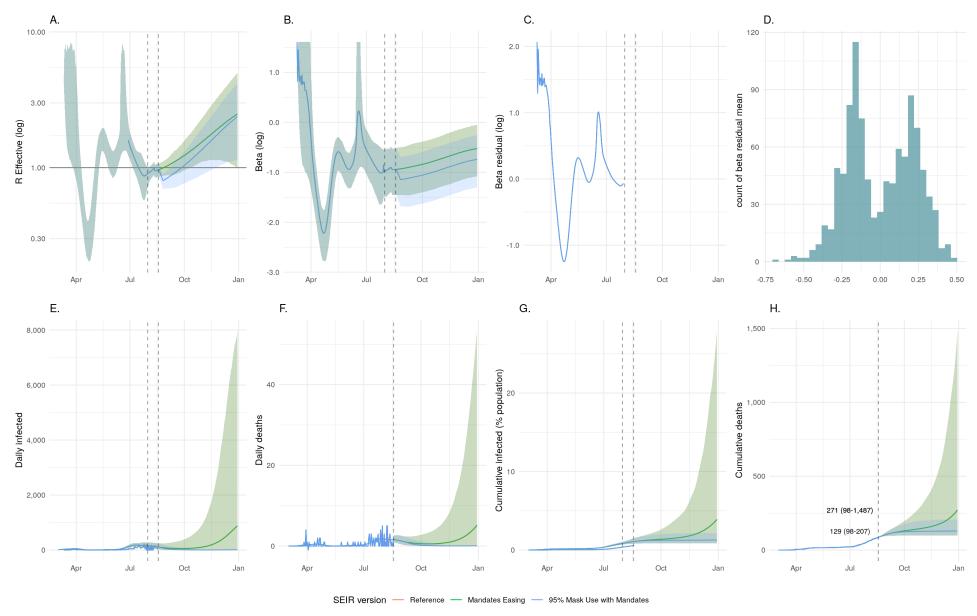
Missouri: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 54 Missouri: Covariate fits and regression coefficients



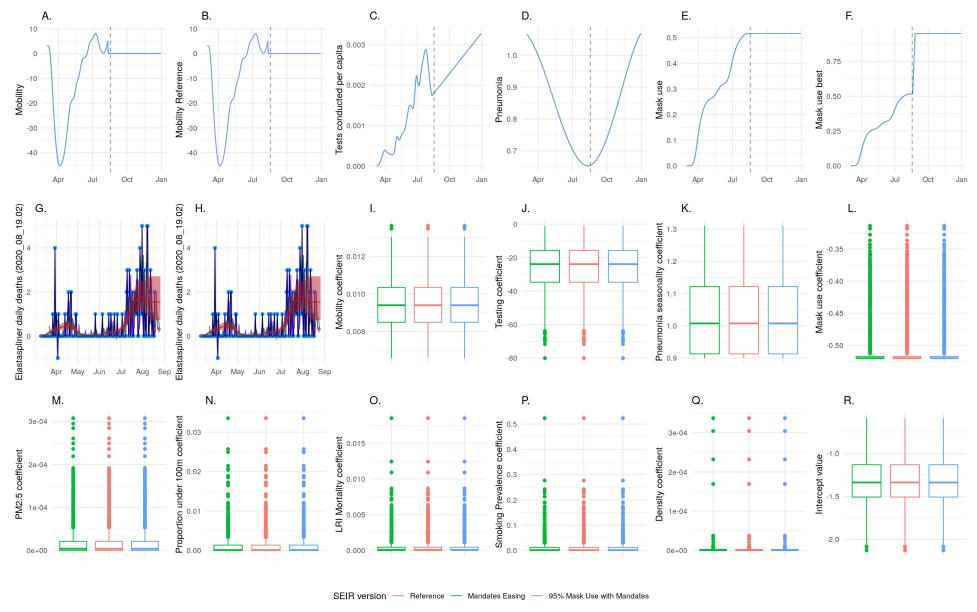
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### 55 Montana: SEIR fit comparison



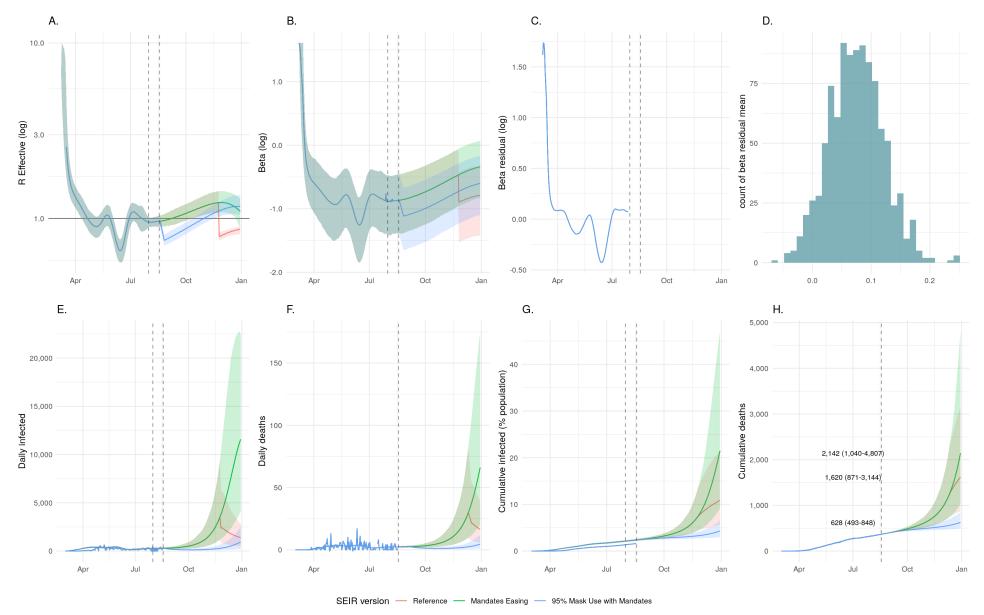
Montana: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 56 Montana: Covariate fits and regression coefficients



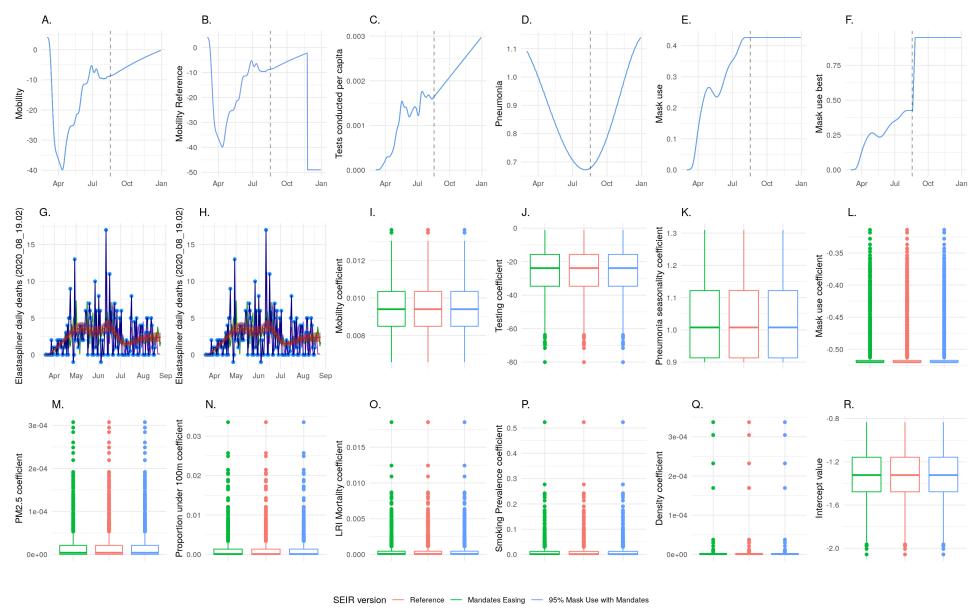
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### 57 Nebraska: SEIR fit comparison



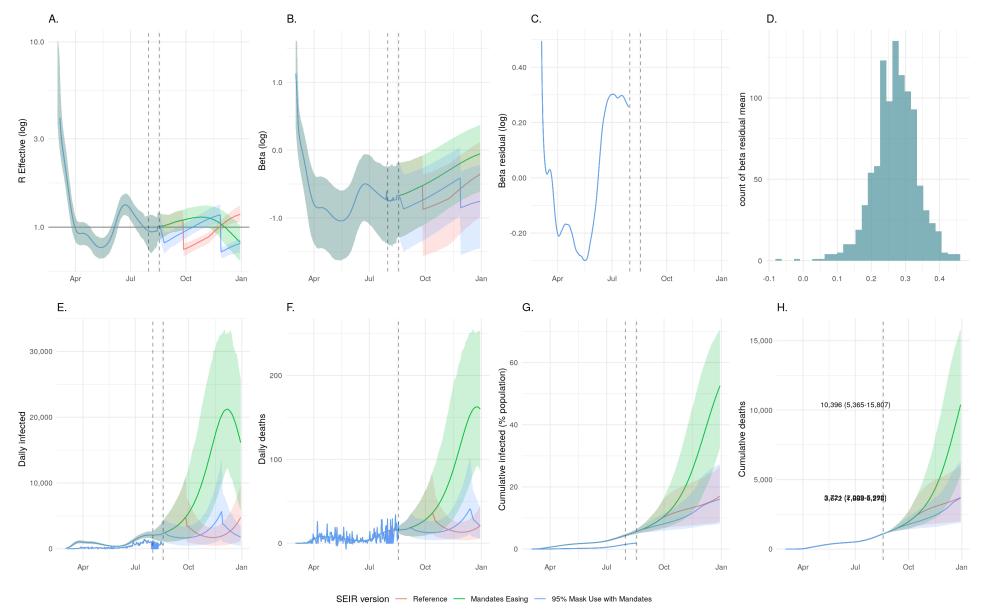
Nebraska: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 58 Nebraska: Covariate fits and regression coefficients



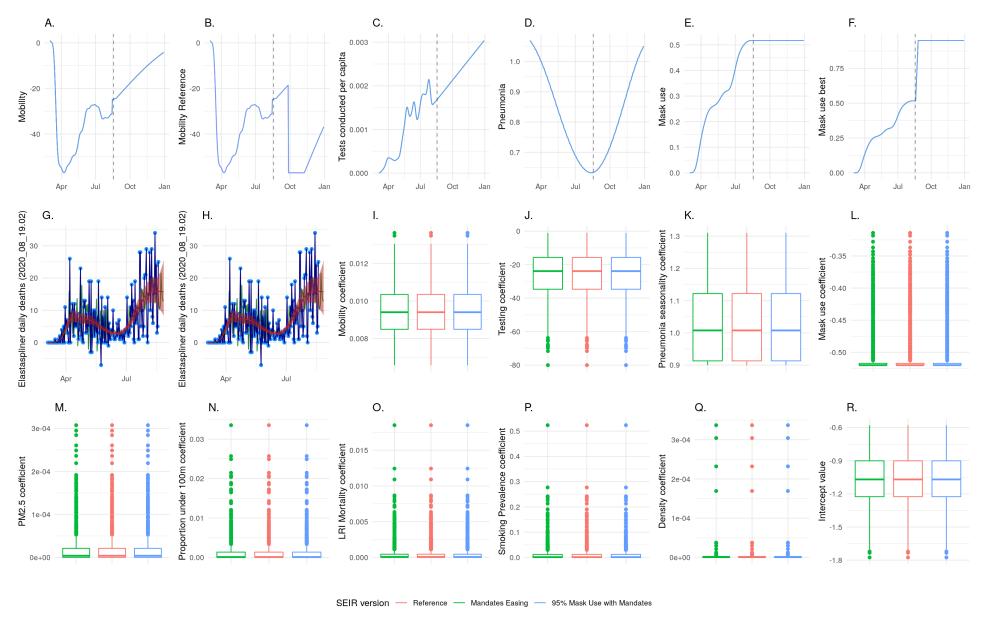
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### 59 Nevada: SEIR fit comparison



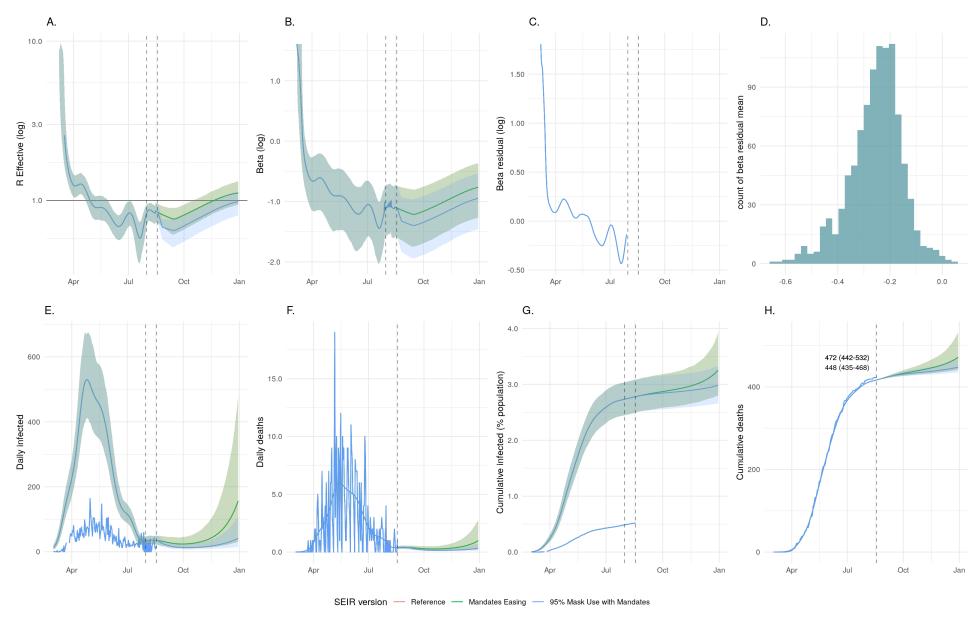
Nevada: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 60 Nevada: Covariate fits and regression coefficients



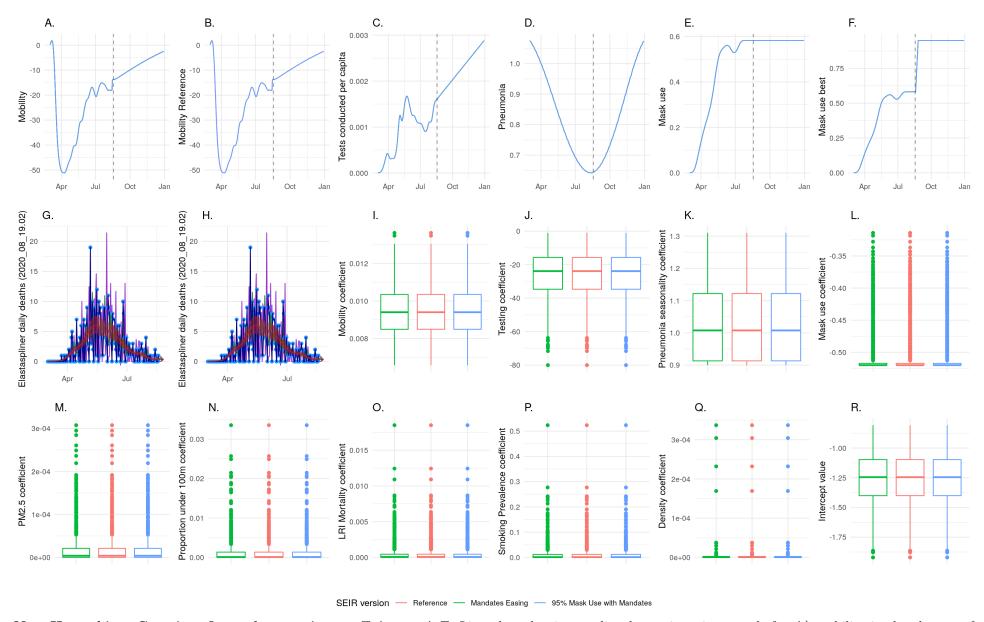
Nevada: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 61 New Hampshire: SEIR fit comparison



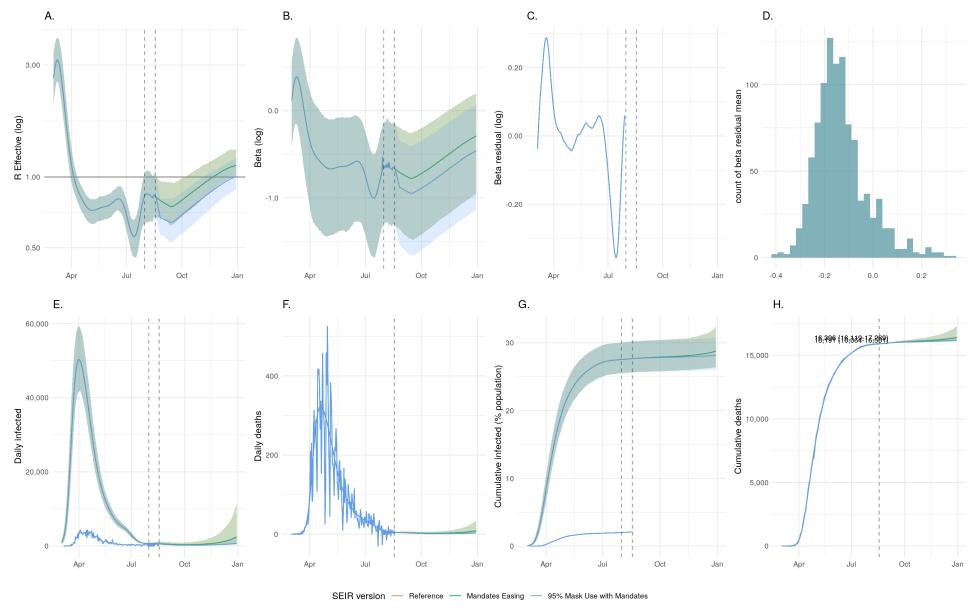
New Hampshire: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

### 62 New Hampshire: Covariate fits and regression coefficients



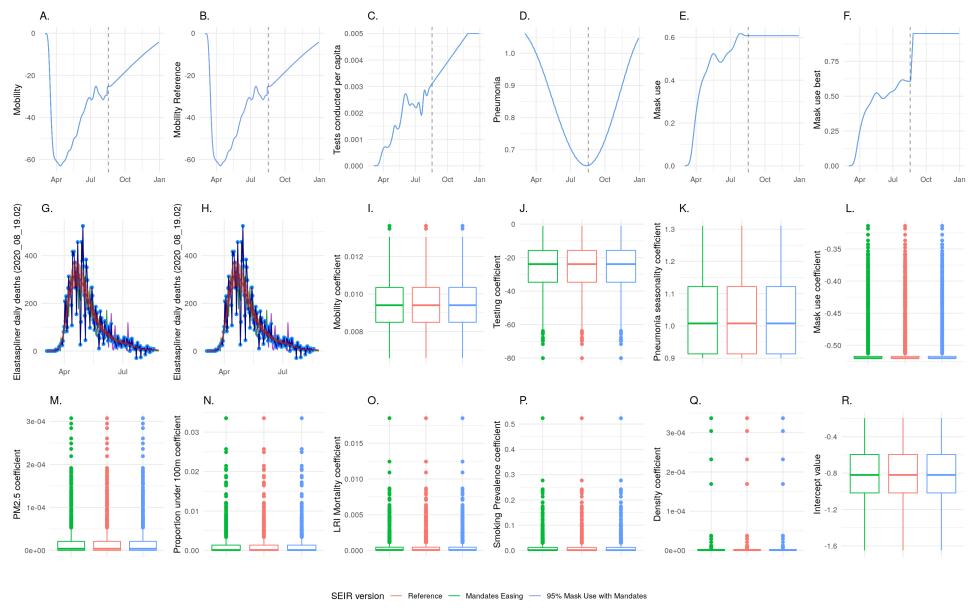
New Hampshire: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 63 New Jersey: SEIR fit comparison



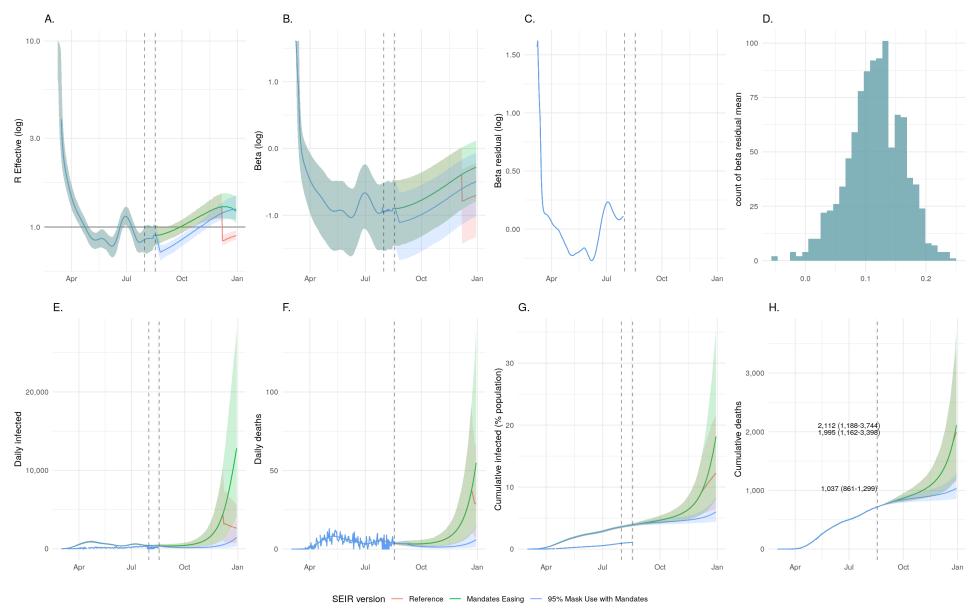
New Jersey: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

# 64 New Jersey: Covariate fits and regression coefficients



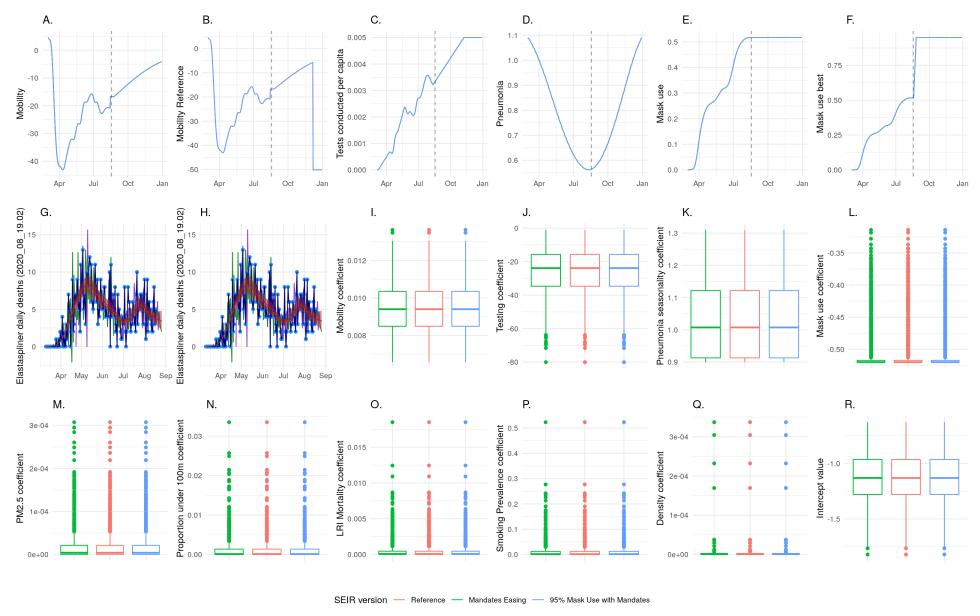
New Jersey: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

### 65 New Mexico: SEIR fit comparison



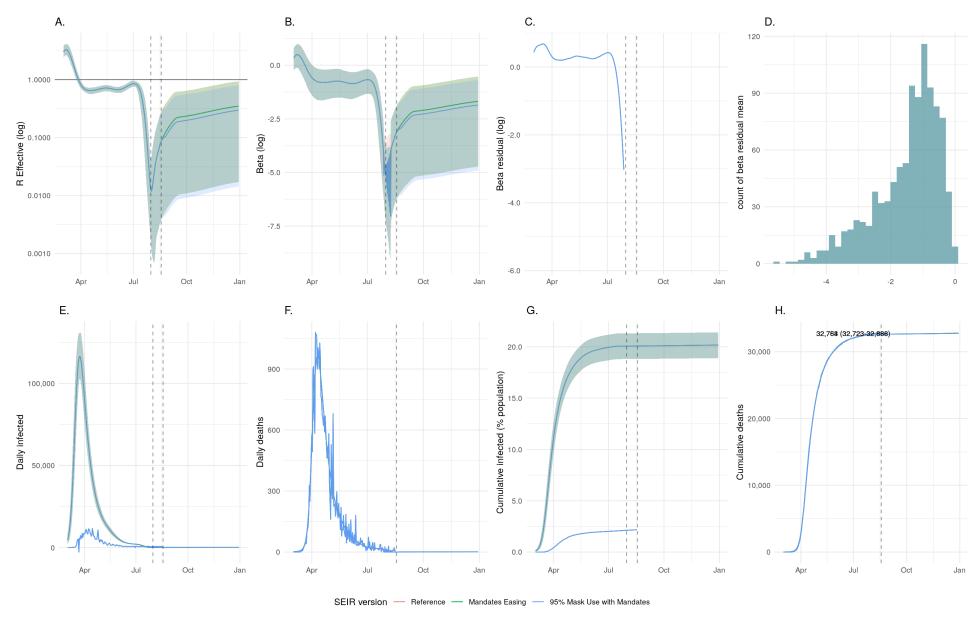
New Mexico: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 66 New Mexico: Covariate fits and regression coefficients



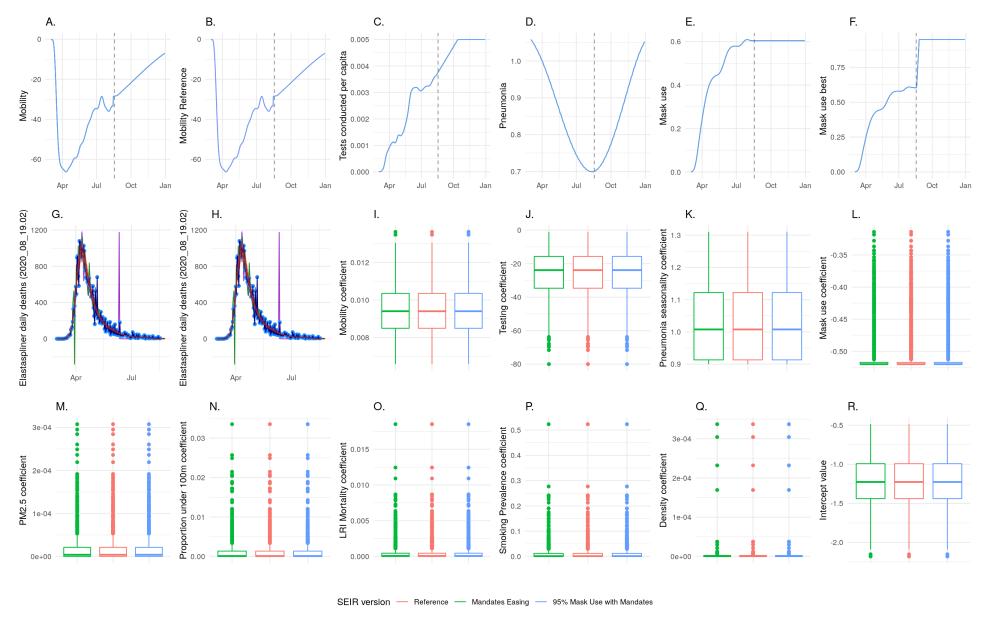
New Mexico: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 67 New York: SEIR fit comparison



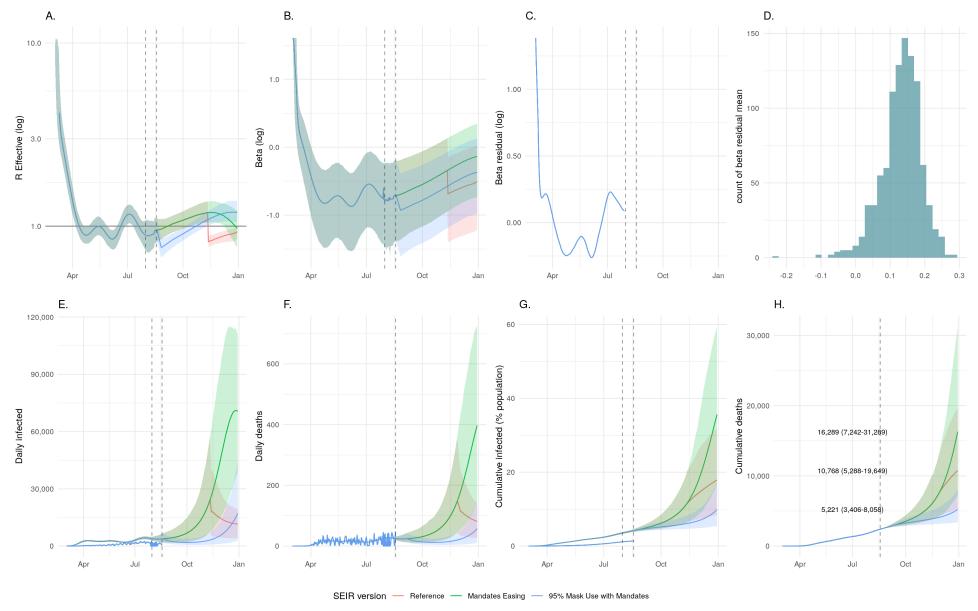
New York: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 68 New York: Covariate fits and regression coefficients



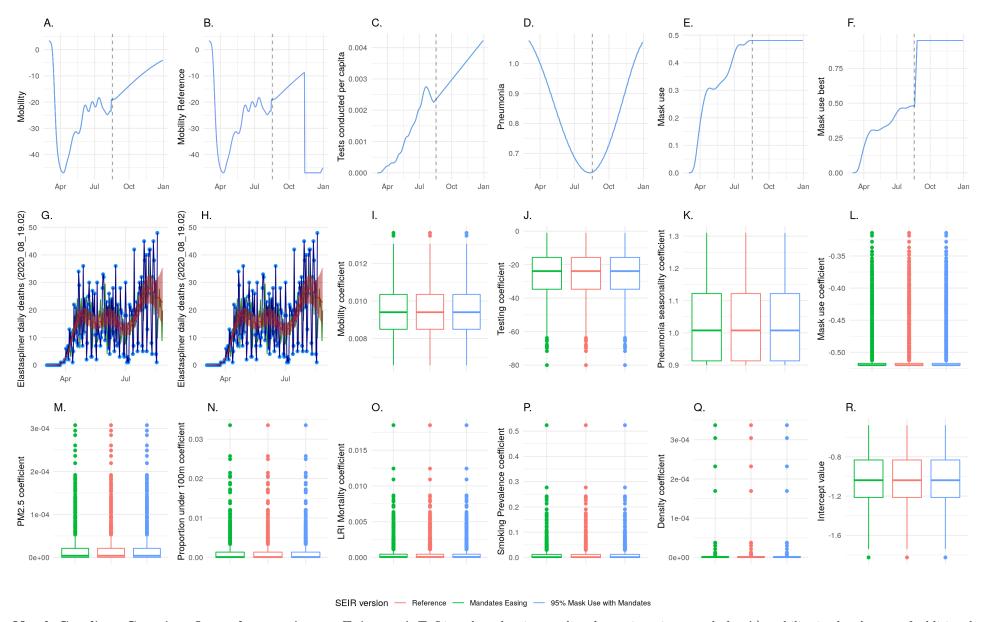
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## 69 North Carolina: SEIR fit comparison



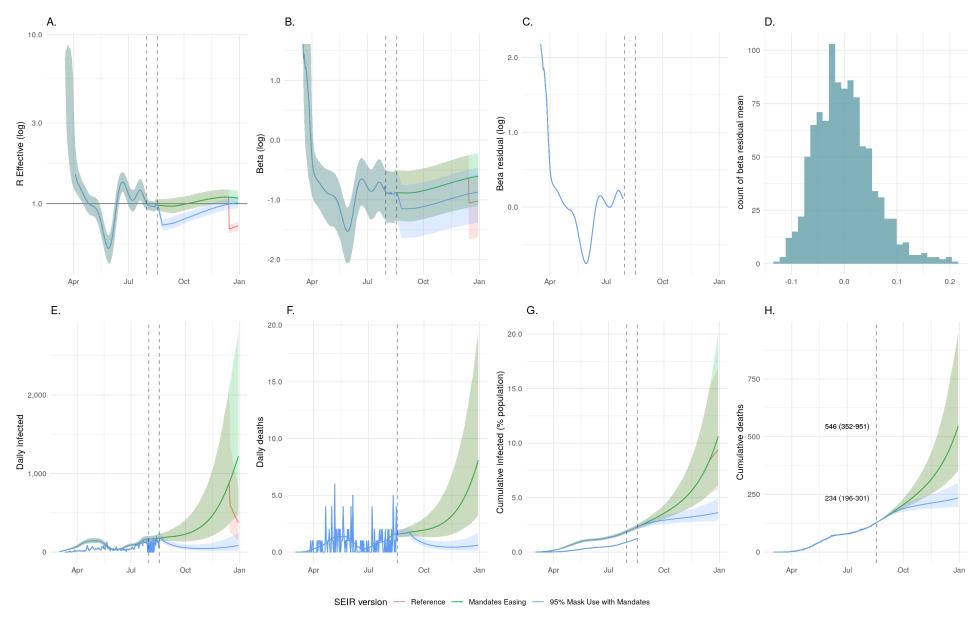
North Carolina: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 70 North Carolina: Covariate fits and regression coefficients



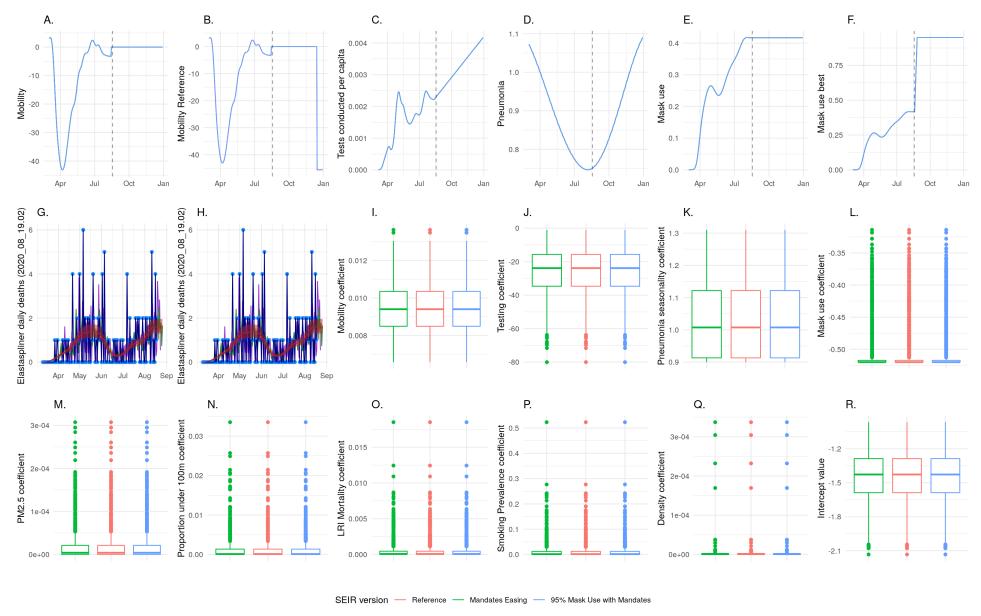
North Carolina: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

## 71 North Dakota: SEIR fit comparison



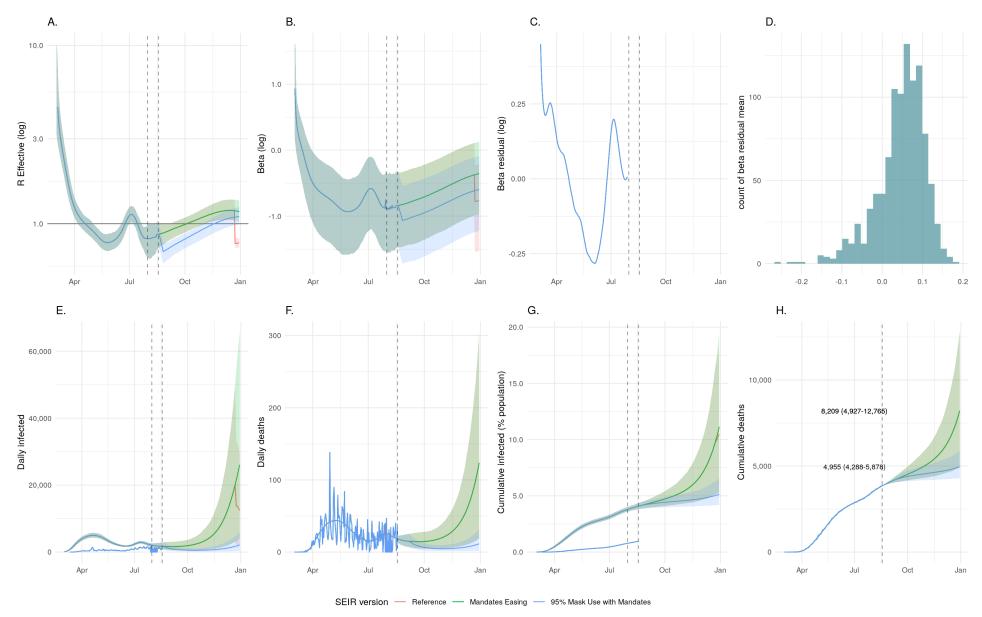
North Dakota: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 72 North Dakota: Covariate fits and regression coefficients



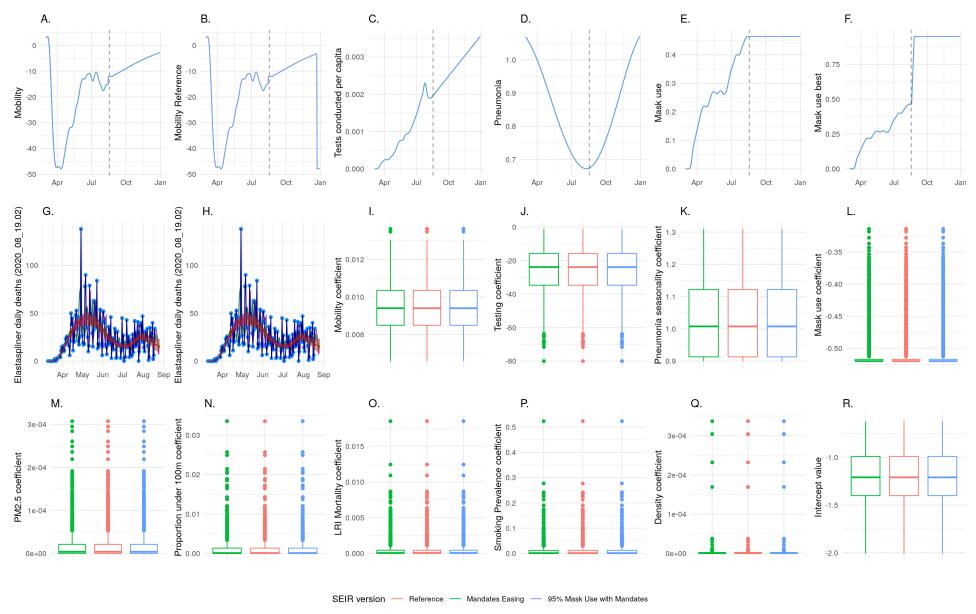
North Dakota: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 73 Ohio: SEIR fit comparison



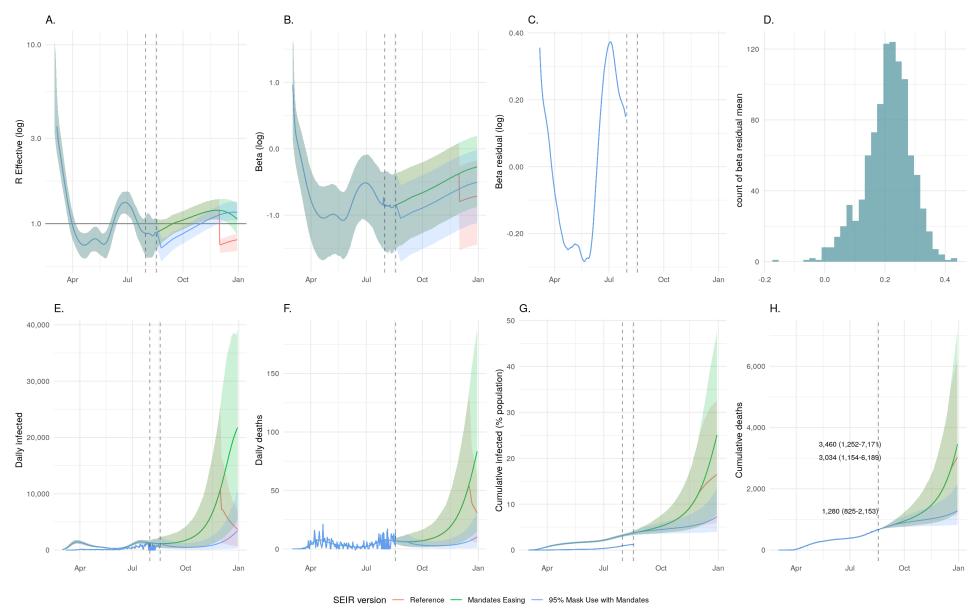
Ohio: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

# 74 Ohio: Covariate fits and regression coefficients



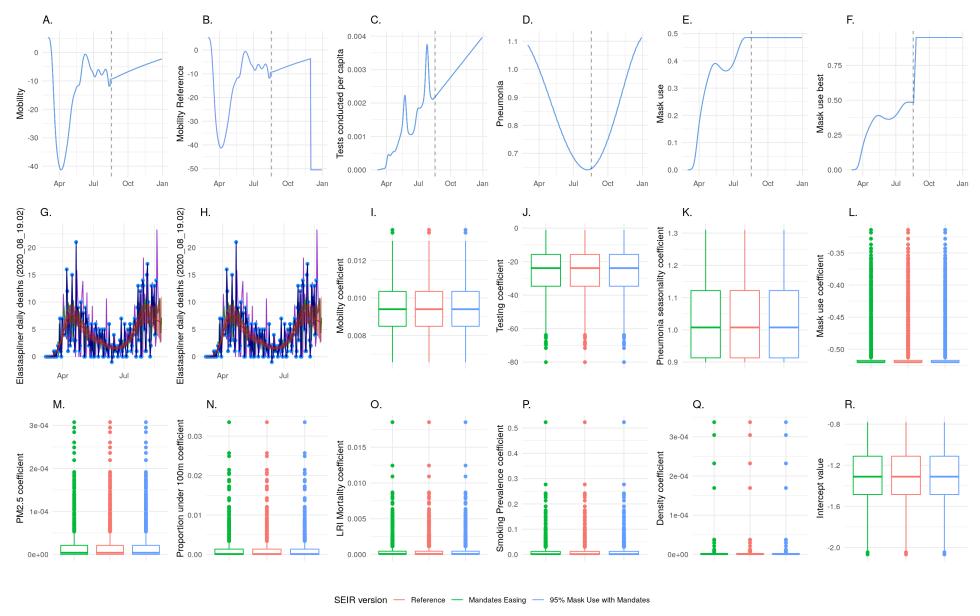
Ohio: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 75 Oklahoma: SEIR fit comparison



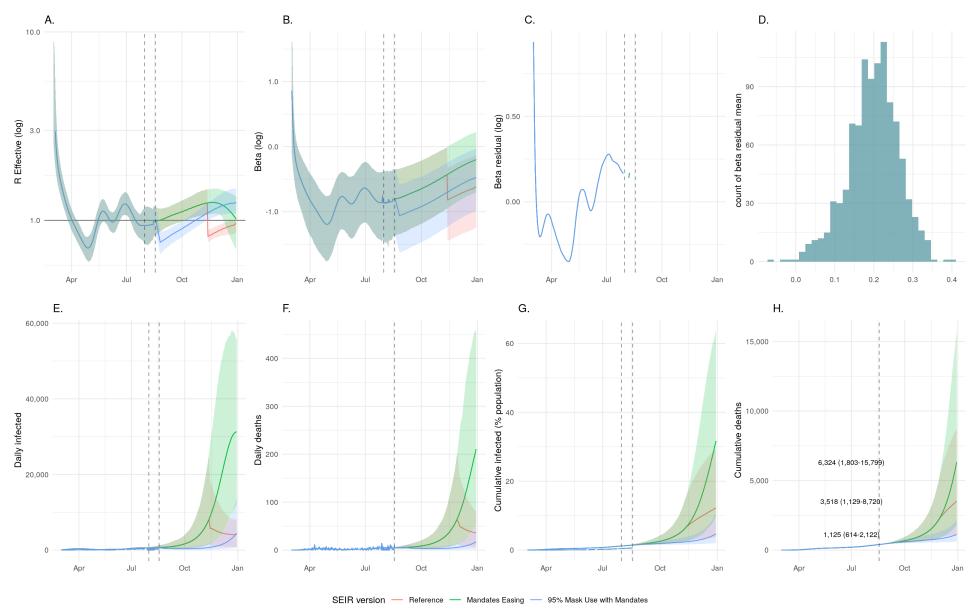
Oklahoma: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 76 Oklahoma: Covariate fits and regression coefficients



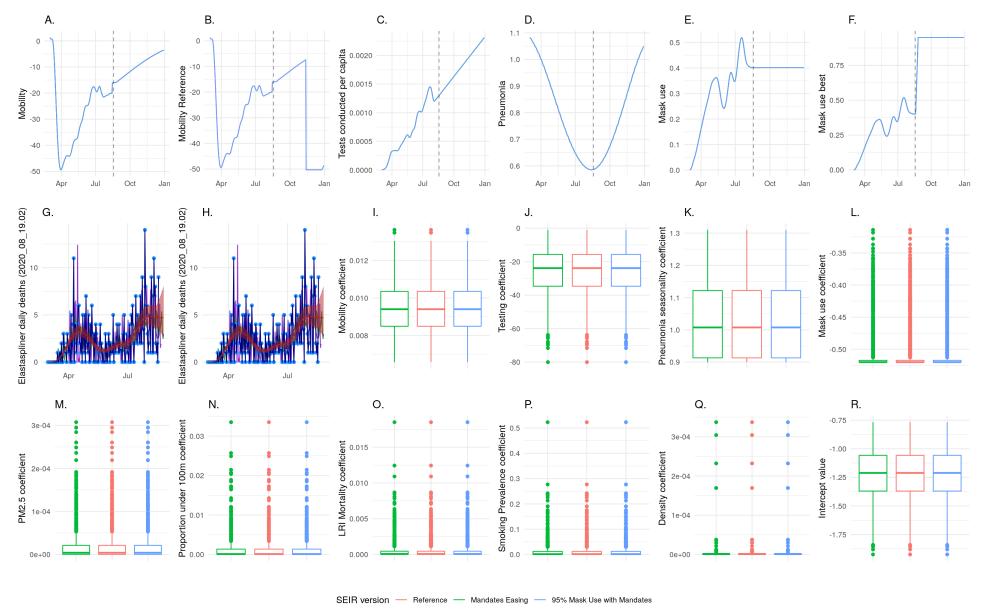
Oklahoma: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 77 Oregon: SEIR fit comparison



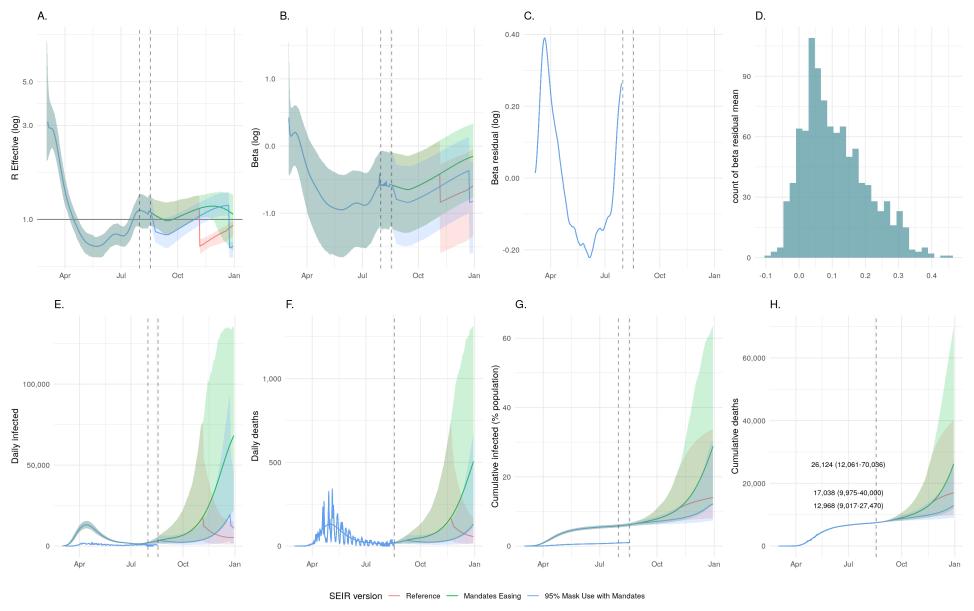
Oregon: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 78 Oregon: Covariate fits and regression coefficients



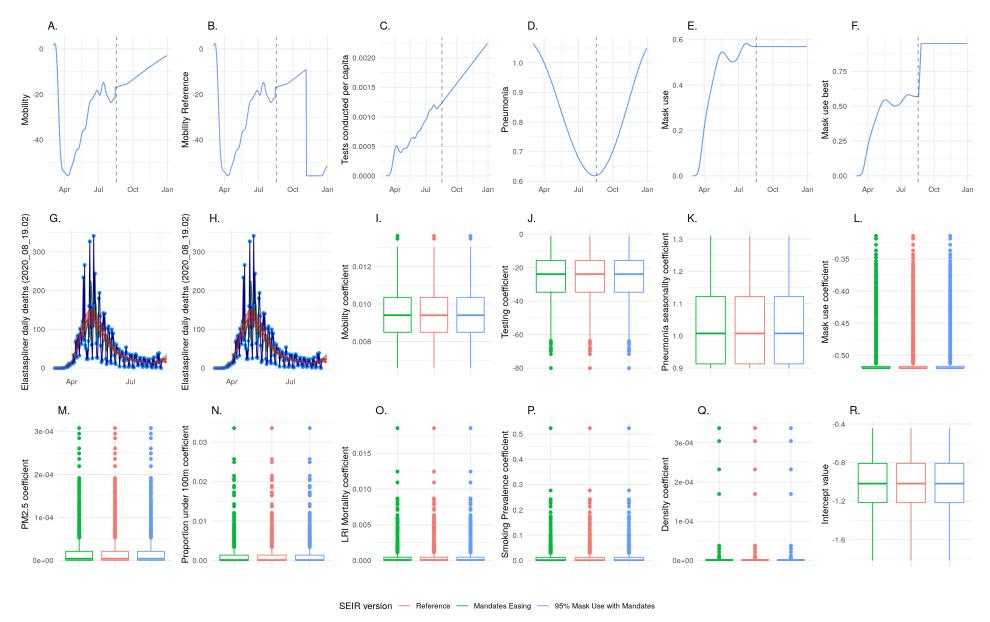
Oregon: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

## 79 Pennsylvania: SEIR fit comparison



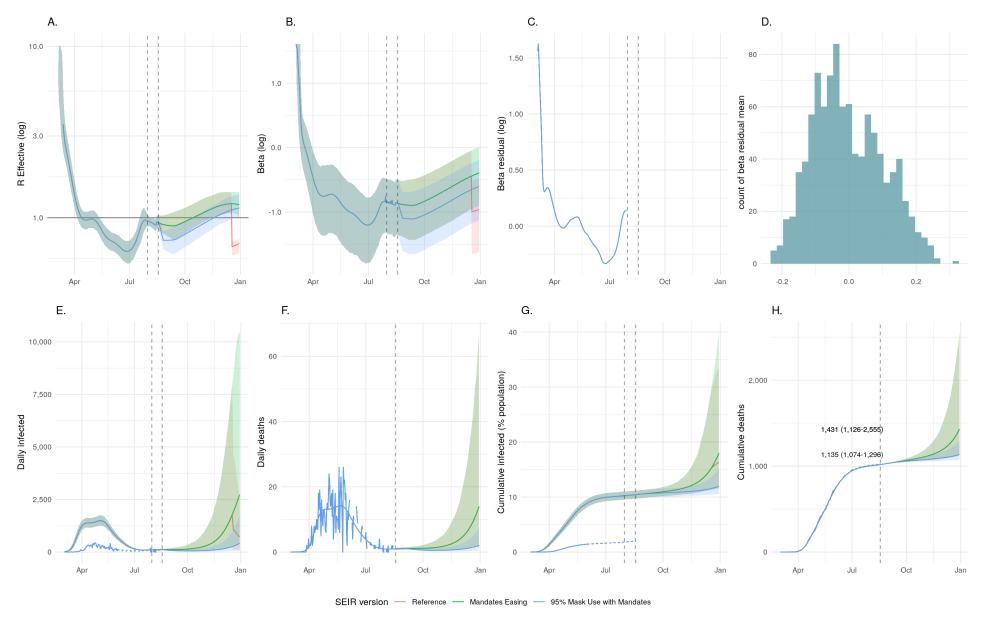
**Pennsylvania: SEIR fit comparison.** A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 80 Pennsylvania: Covariate fits and regression coefficients



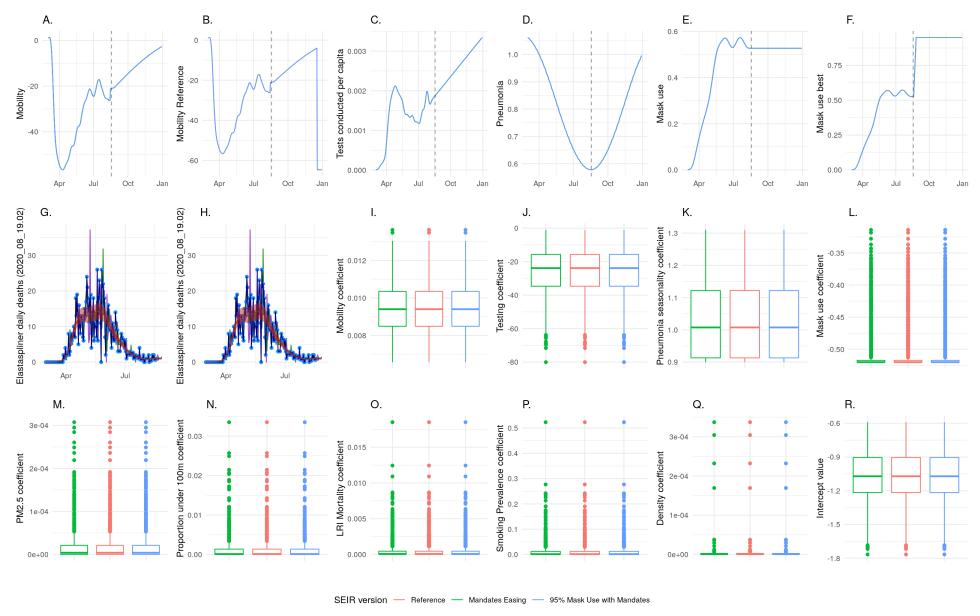
Pennsylvania: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 81 Rhode Island: SEIR fit comparison



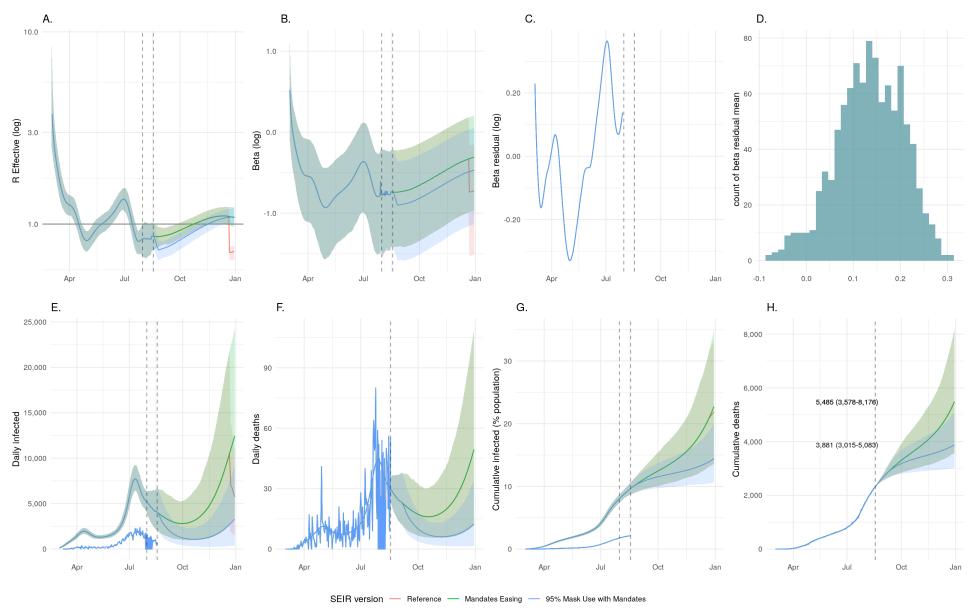
Rhode Island: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 82 Rhode Island: Covariate fits and regression coefficients



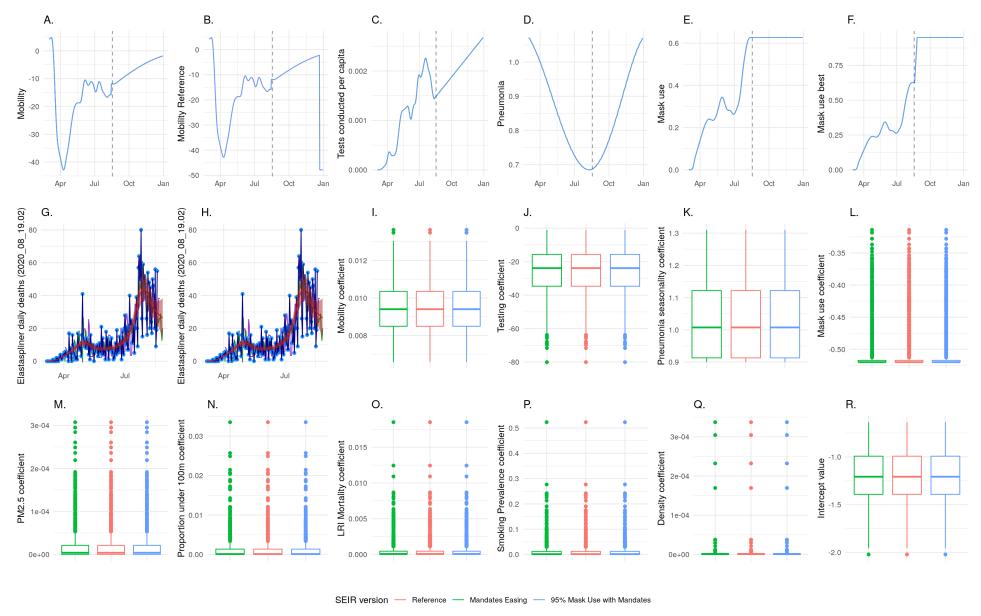
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## 83 South Carolina: SEIR fit comparison



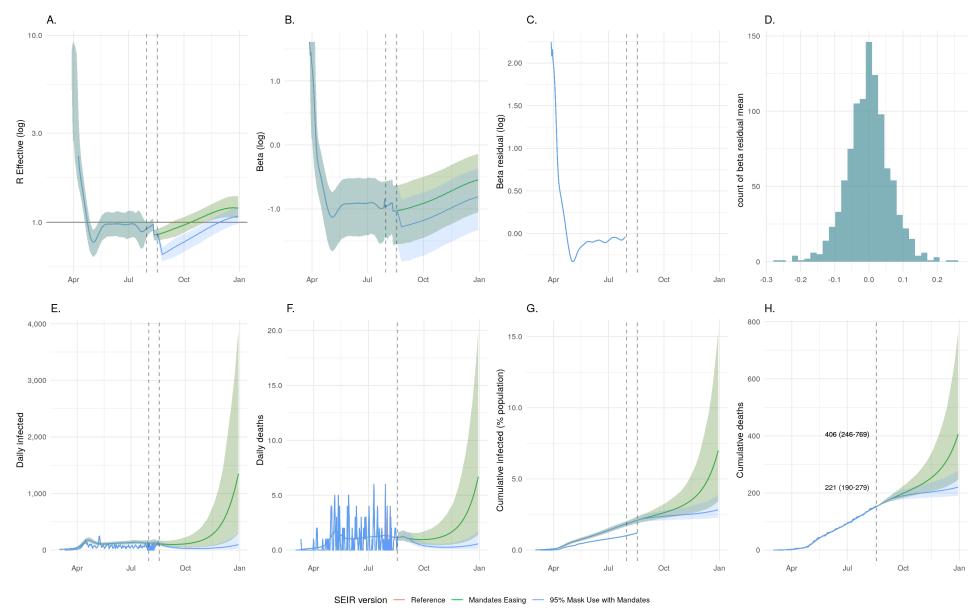
South Carolina: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 84 South Carolina: Covariate fits and regression coefficients



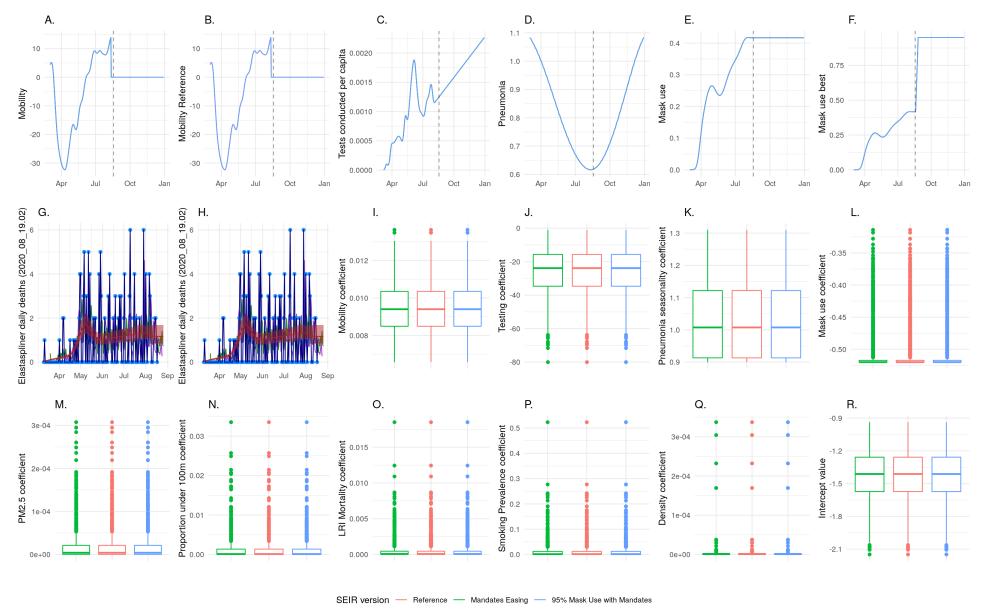
South Carolina: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

## 85 South Dakota: SEIR fit comparison



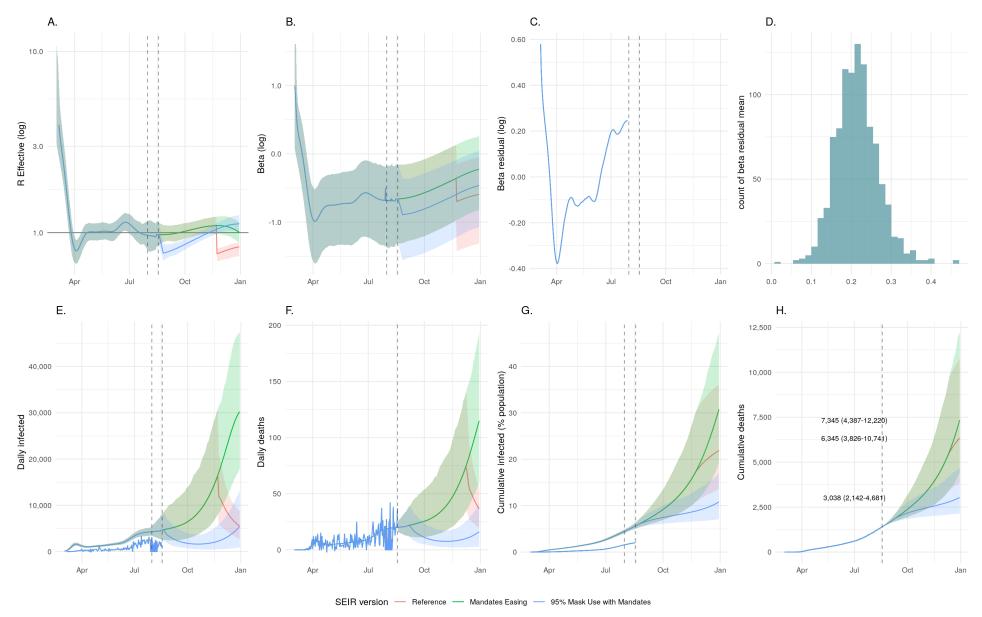
**South Dakota: SEIR fit comparison. A**: predicted R effective for each model through December 31. **B**: predicted SEIR beta parameter. **C**: residual of predicted beta and the observed value calculated directly from infection data over time. **D**: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. **E**: predicted daily infections from each model through December 31. **F**: predicted daily deaths from each model through December 31. **G**: predicted cumulative infections through December 31, as a proportion of the total population. **H**: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 86 South Dakota: Covariate fits and regression coefficients



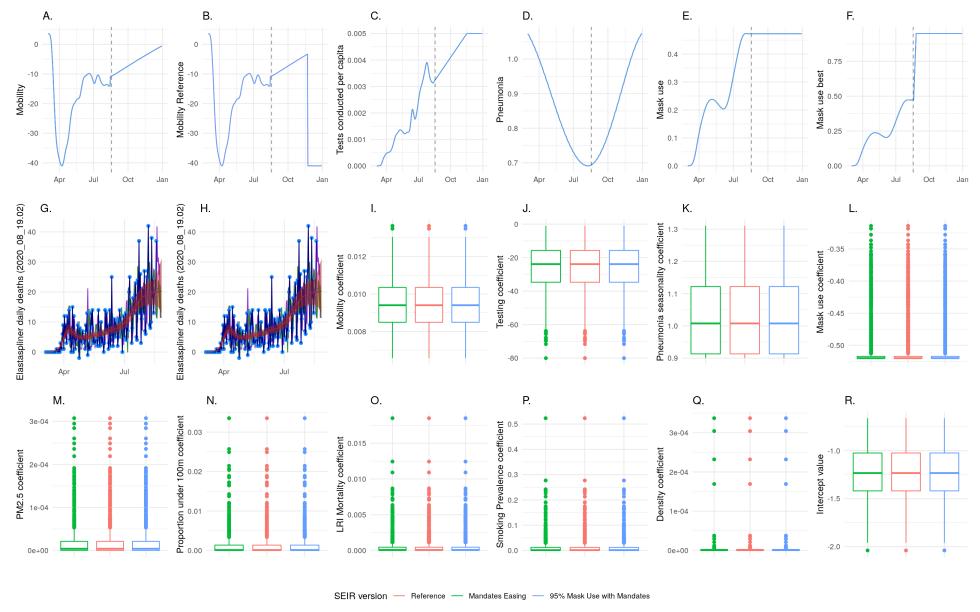
South Dakota: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

## 87 Tennessee: SEIR fit comparison



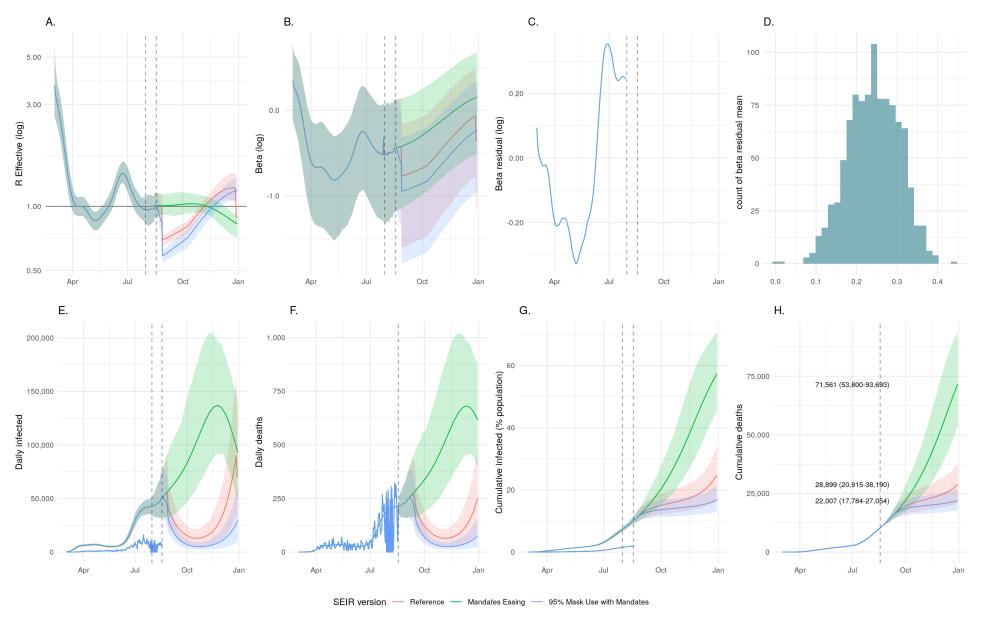
**Tennessee: SEIR fit comparison.** A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 88 Tennessee: Covariate fits and regression coefficients



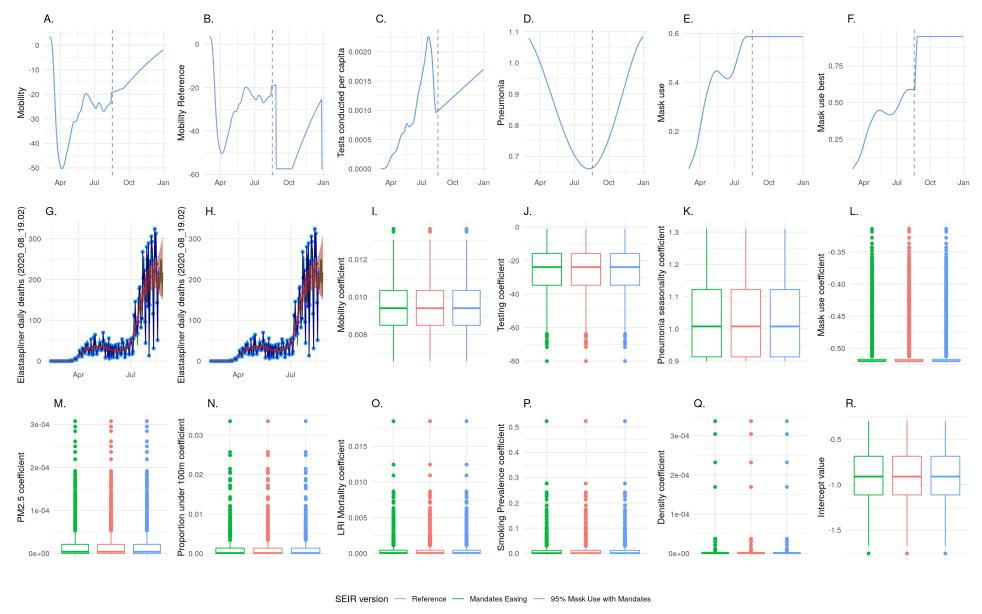
Tennessee: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

## 89 Texas: SEIR fit comparison



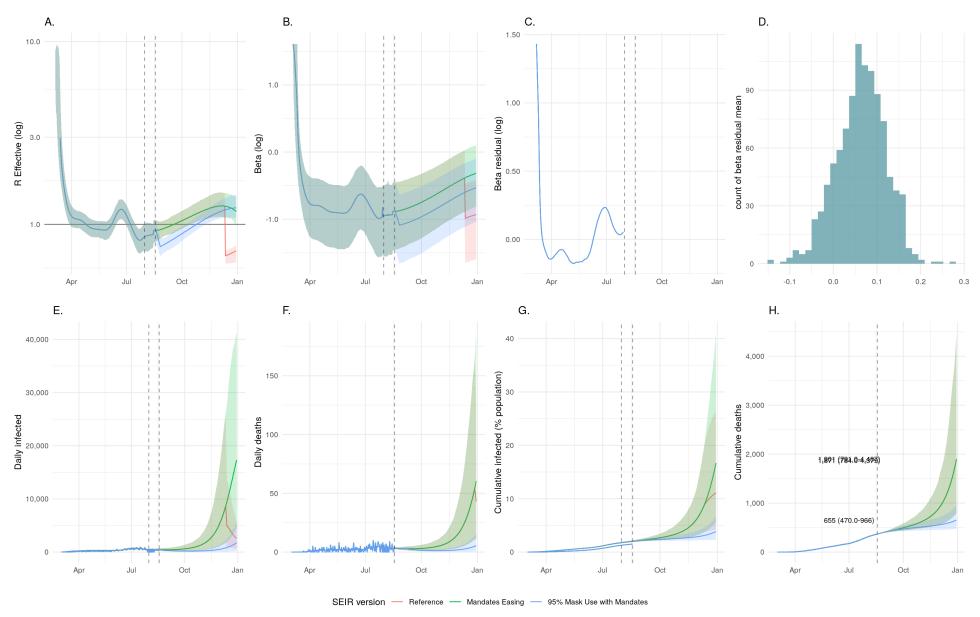
**Texas: SEIR fit comparison. A:** predicted R effective for each model through December 31. **B:** predicted SEIR beta parameter. **C:** residual of predicted beta and the observed value calculated directly from infection data over time. **D:** histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. **E:** predicted daily infections from each model through December 31. **F:** predicted daily deaths from each model through December 31. **G:** predicted cumulative infections through December 31, as a proportion of the total population. **H:** predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

#### 90 Texas: Covariate fits and regression coefficients



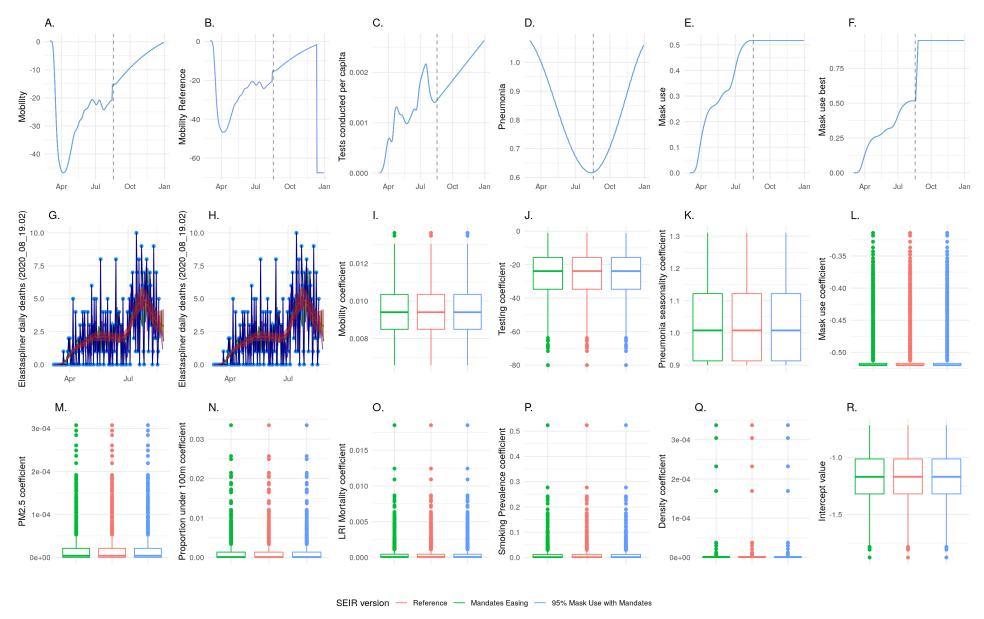
Texas: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 91 Utah: SEIR fit comparison



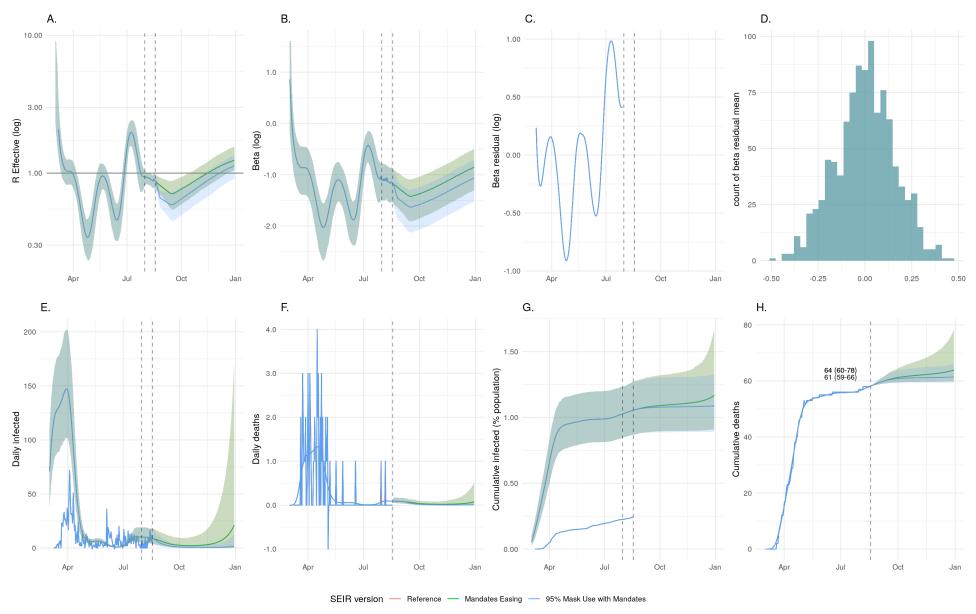
**Utah: SEIR fit comparison. A**: predicted R effective for each model through December 31. **B**: predicted SEIR beta parameter. **C**: residual of predicted beta and the observed value calculated directly from infection data over time. **D**: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. **E**: predicted daily infections from each model through December 31. **F**: predicted daily deaths from each model through December 31. **G**: predicted cumulative infections through December 31, as a proportion of the total population. **H**: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 92 Utah: Covariate fits and regression coefficients



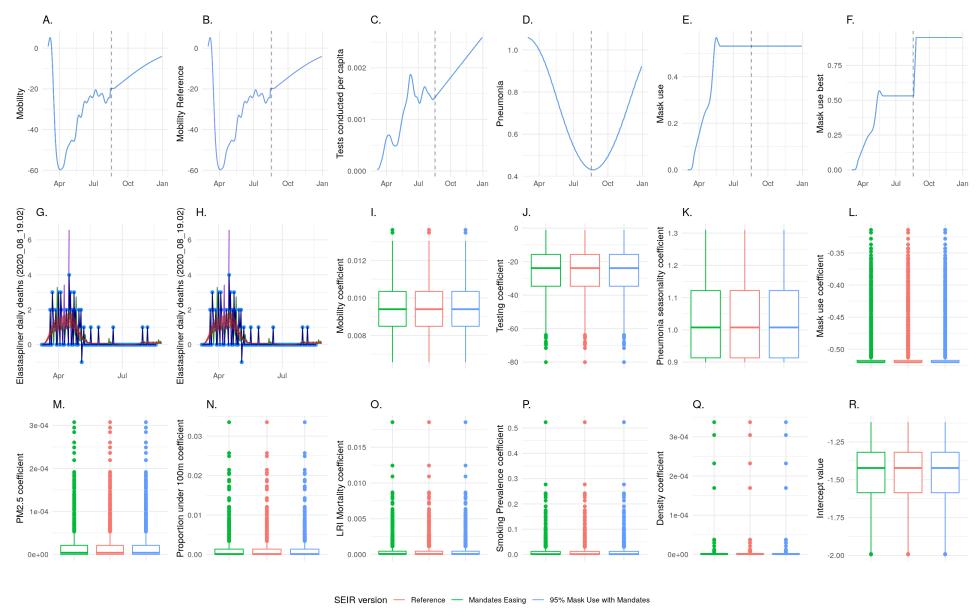
Utah: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

## 93 Vermont: SEIR fit comparison



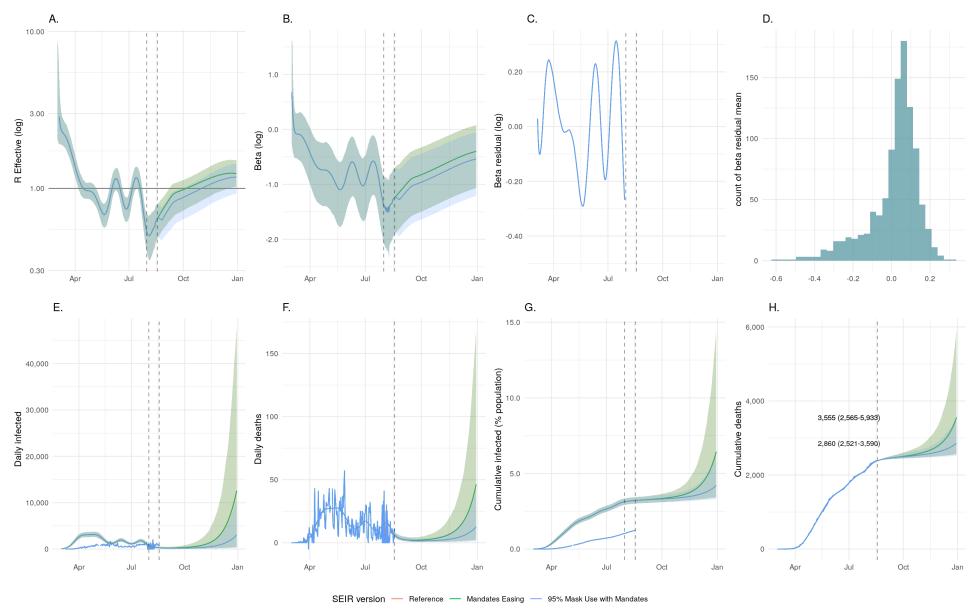
**Vermont: SEIR fit comparison. A**: predicted R effective for each model through December 31. **B**: predicted SEIR beta parameter. **C**: residual of predicted beta and the observed value calculated directly from infection data over time. **D**: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. **E**: predicted daily infections from each model through December 31. **F**: predicted daily deaths from each model through December 31. **G**: predicted cumulative infections through December 31, as a proportion of the total population. **H**: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 94 Vermont: Covariate fits and regression coefficients



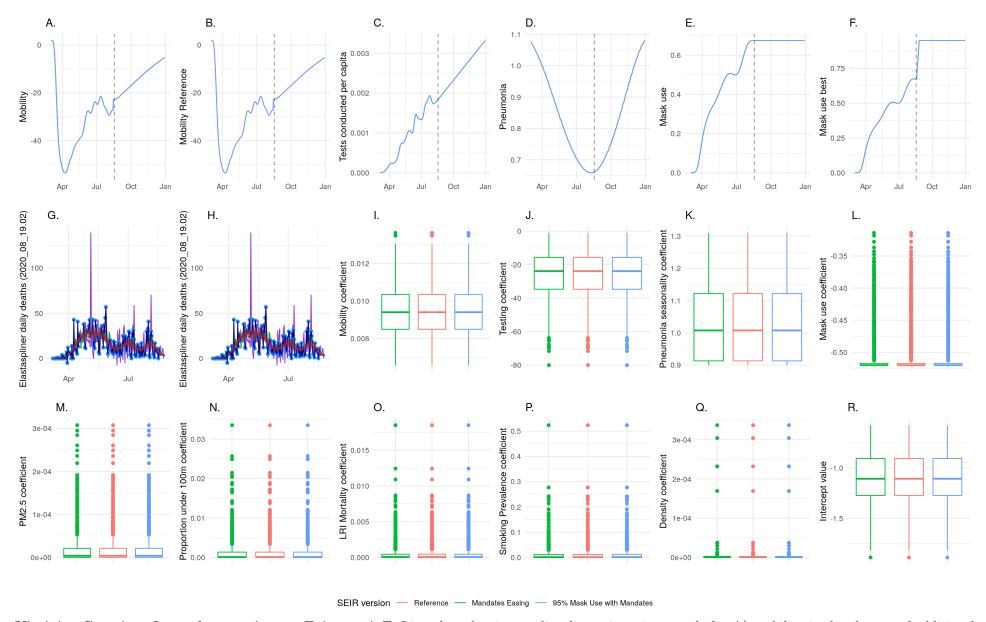
Vermont: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 95 Virginia: SEIR fit comparison



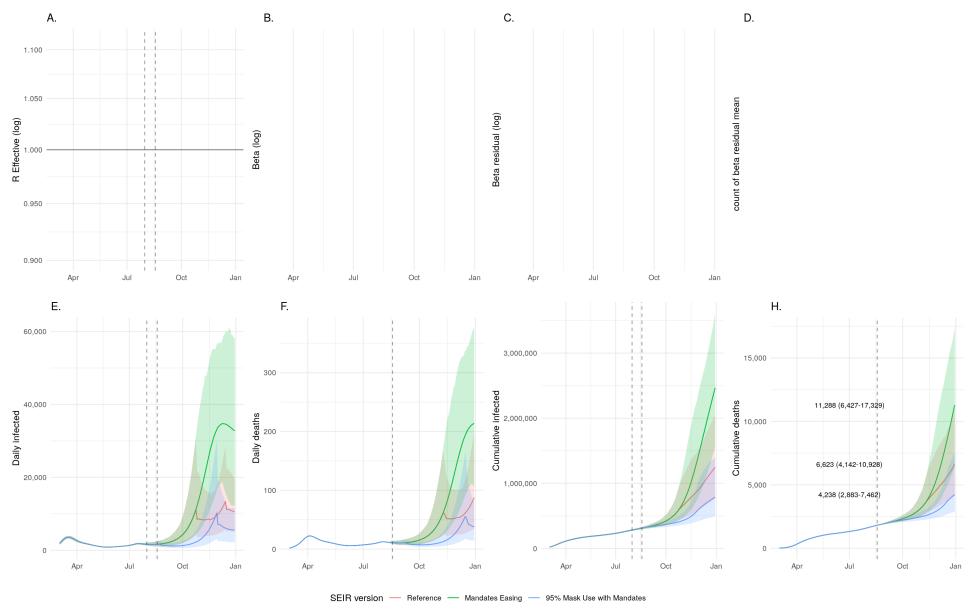
Virginia: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 96 Virginia: Covariate fits and regression coefficients



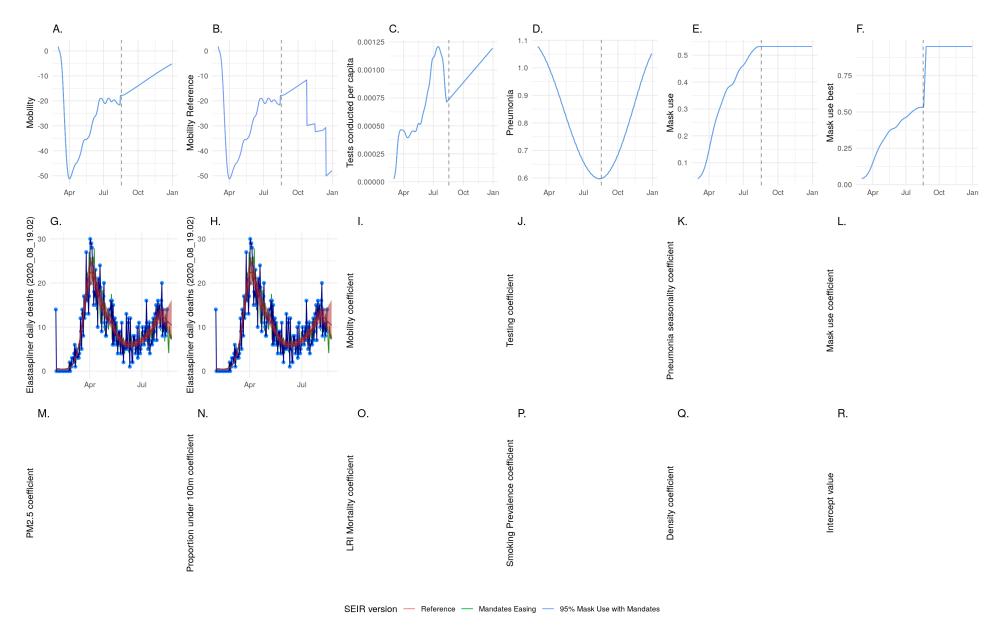
Virginia: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 97 Washington: SEIR fit comparison



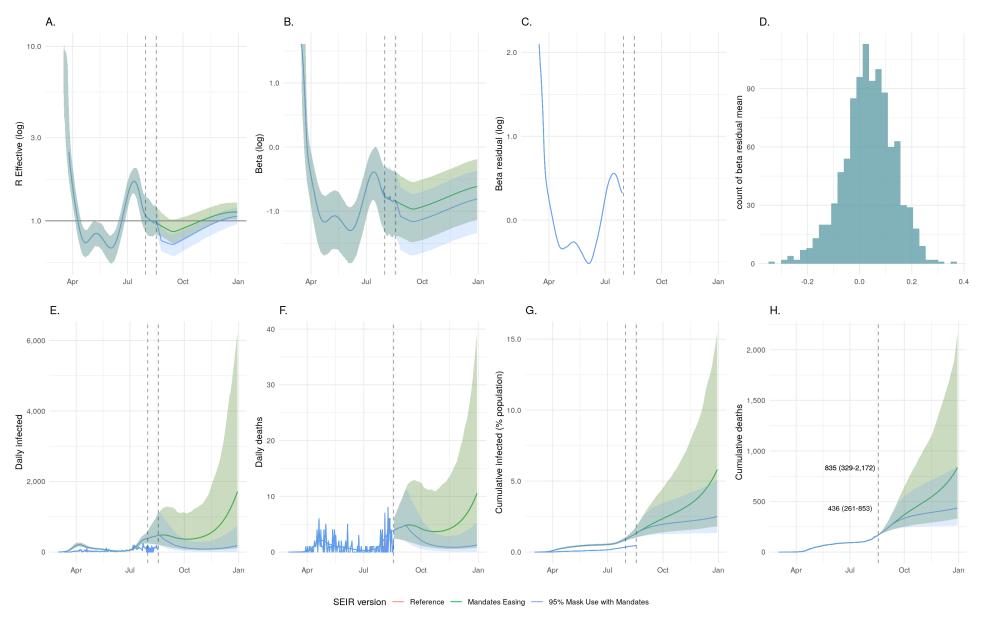
Washington: SEIR fit comparison. Panels A-D display values that are not directly calculated for aggregate locations. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 98 Washington: Covariate fits and regression coefficients



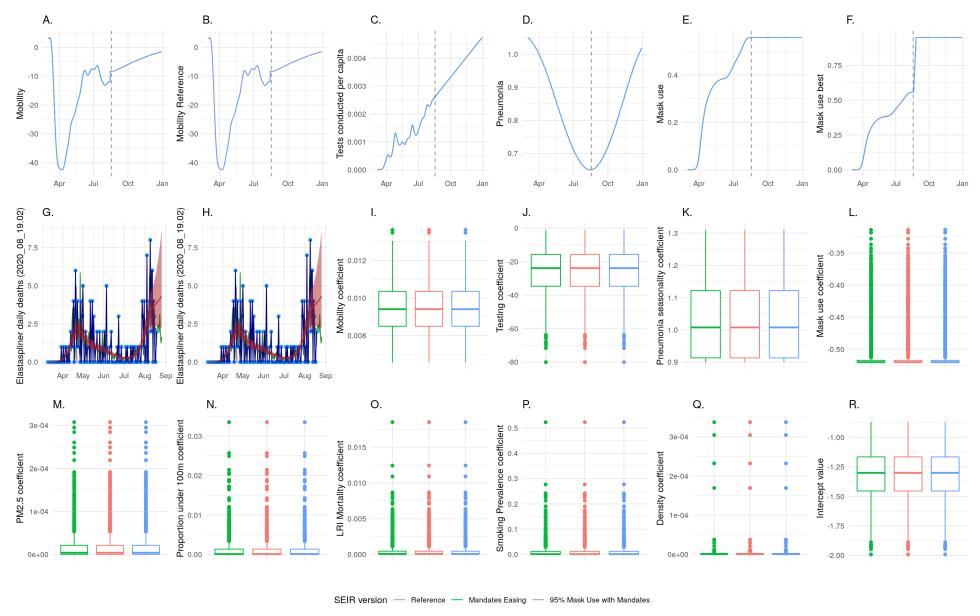
Washington: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). Panels I-R display coefficients for a regression fit to log(beta), which is not applicable to aggregate locations.

## 99 West Virginia: SEIR fit comparison



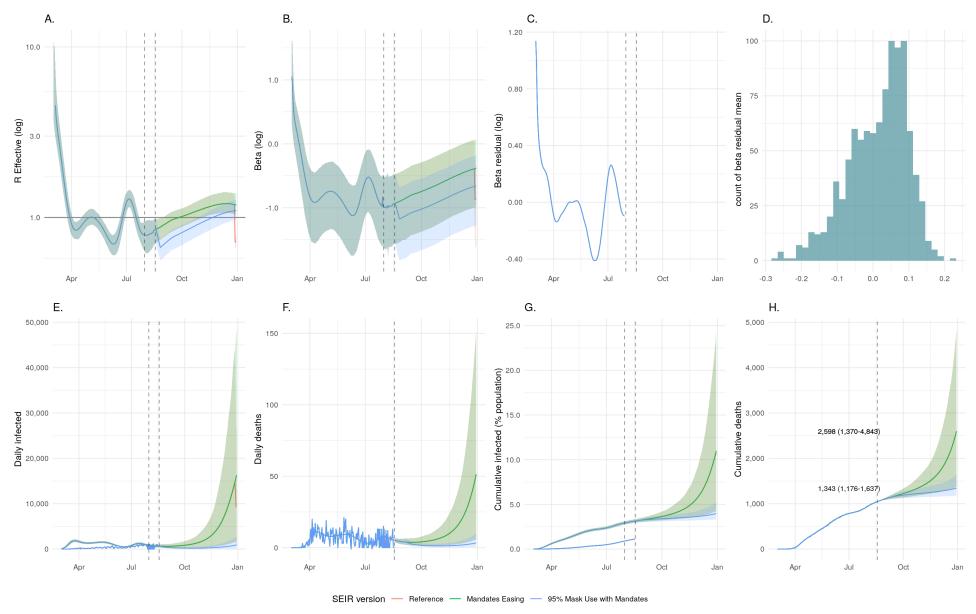
West Virginia: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 100 West Virginia: Covariate fits and regression coefficients



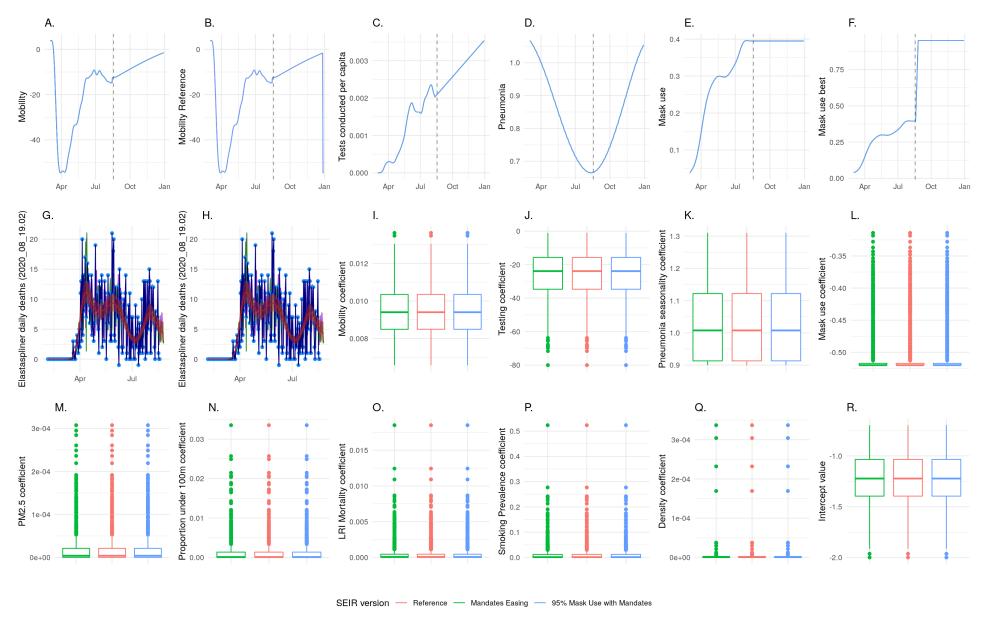
West Virginia: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

## 101 Wisconsin: SEIR fit comparison



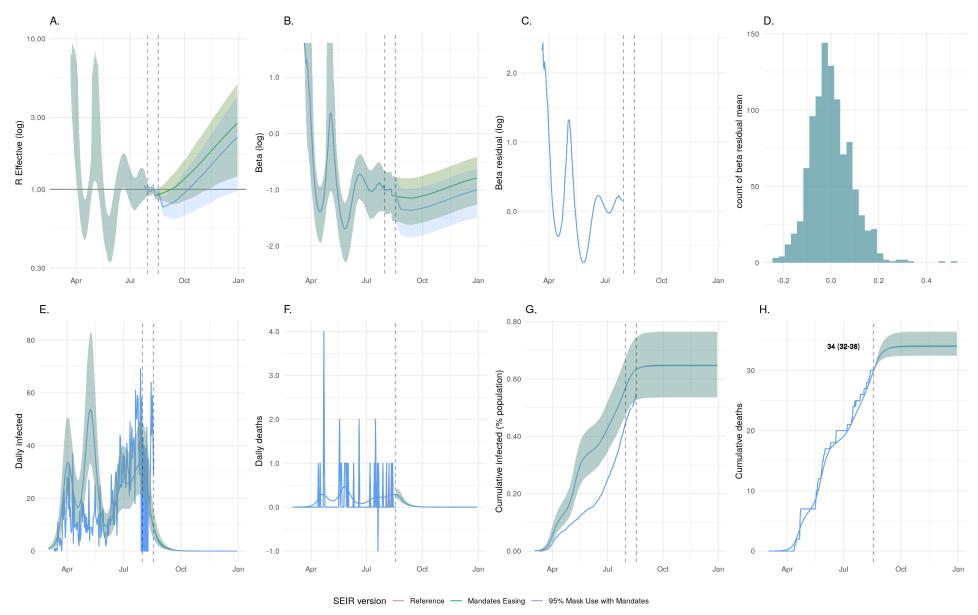
Wisconsin: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 102 Wisconsin: Covariate fits and regression coefficients



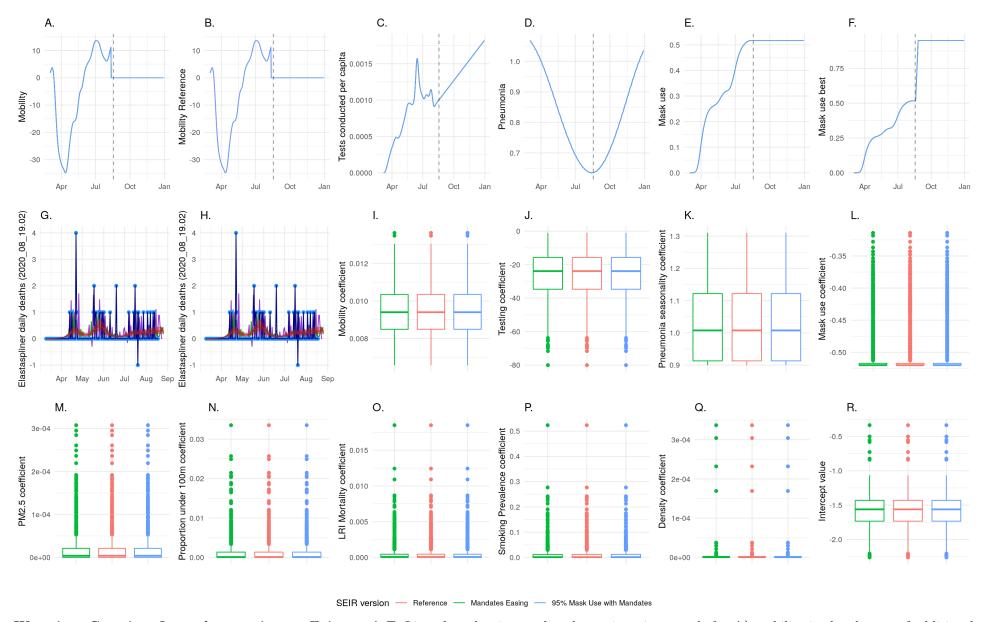
Wisconsin: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 103 Wyoming: SEIR fit comparison



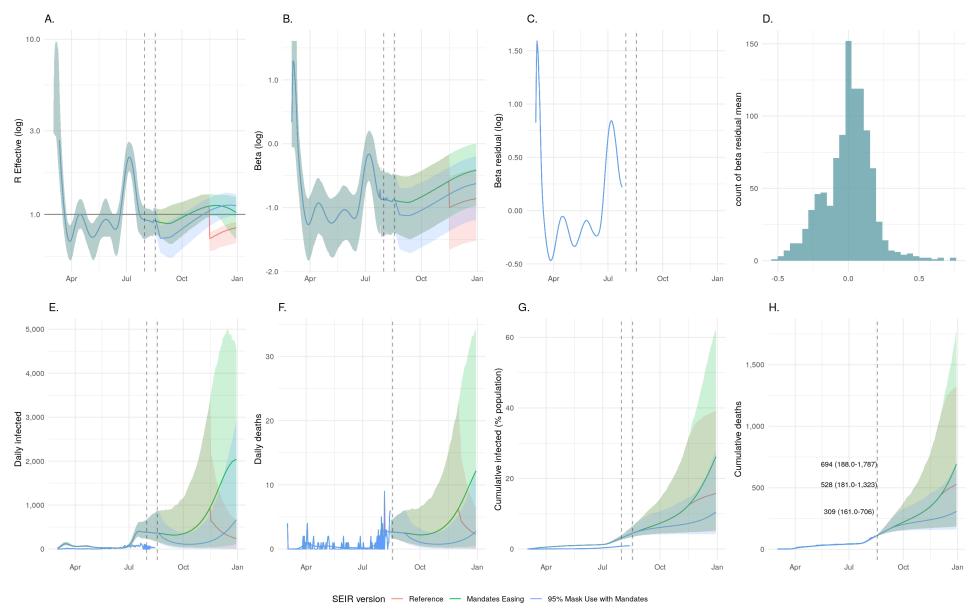
Wyoming: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 104 Wyoming: Covariate fits and regression coefficients



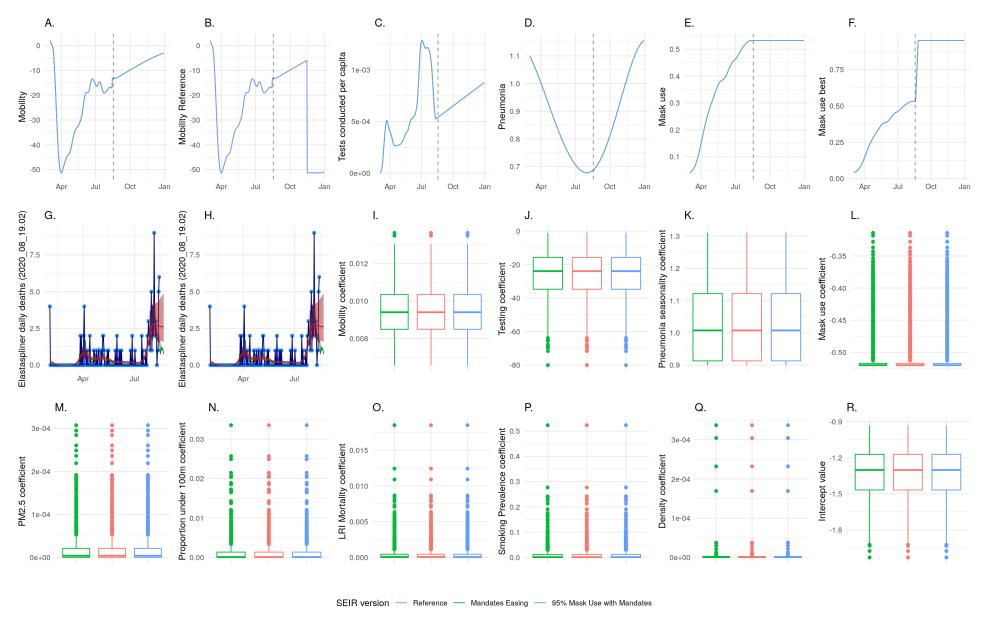
Wyoming: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 105 Spokane County: SEIR fit comparison



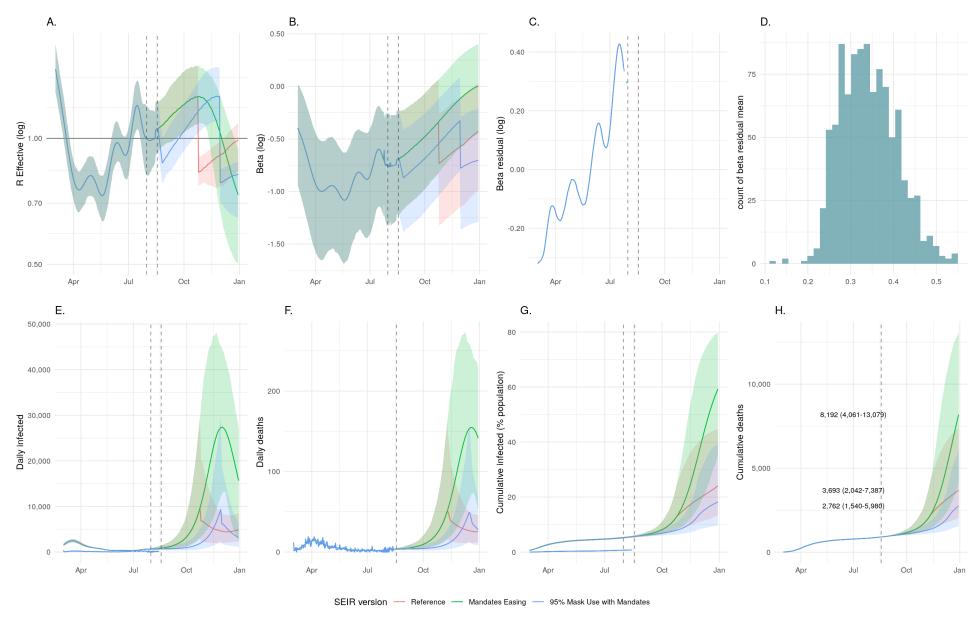
**Spokane County: SEIR fit comparison. A**: predicted R effective for each model through December 31. **B**: predicted SEIR beta parameter. **C**: residual of predicted beta and the observed value calculated directly from infection data over time. **D**: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. **E**: predicted daily infections from each model through December 31. **F**: predicted daily deaths from each model through December 31. **G**: predicted cumulative infections through December 31, as a proportion of the total population. **H**: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 106 Spokane County: Covariate fits and regression coefficients



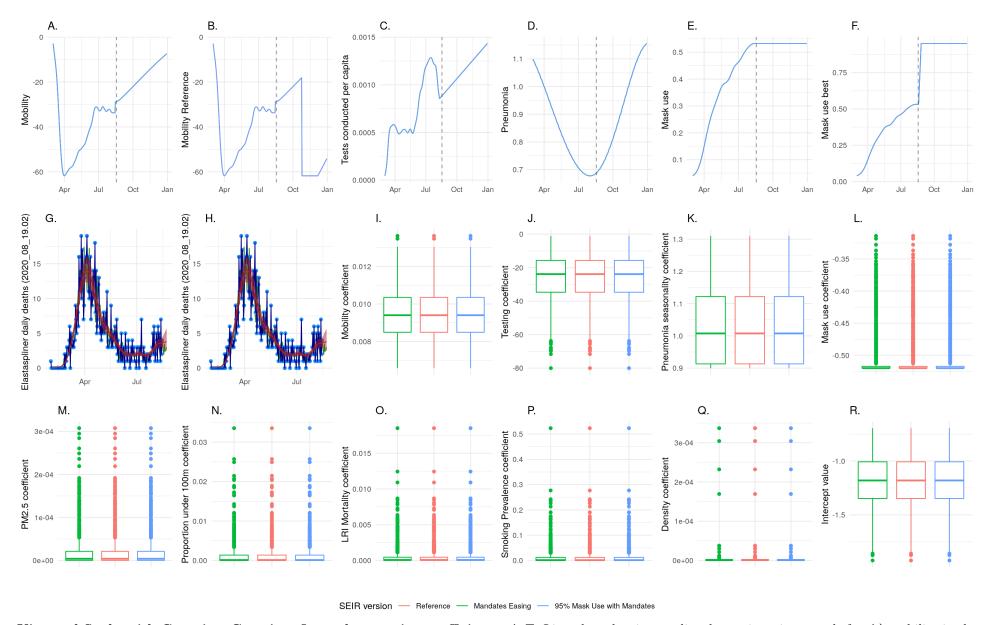
Spokane County: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

# 107 King and Snohomish Counties: SEIR fit comparison



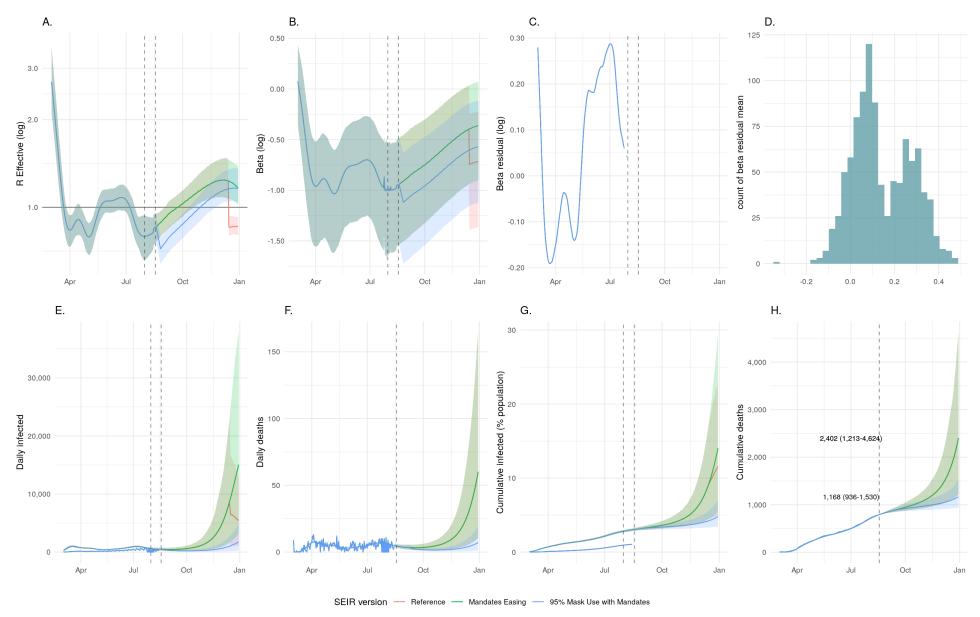
King and Snohomish Counties: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in light blue.

## 108 King and Snohomish Counties: Covariate fits and regression coefficients



King and Snohomish Counties: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).

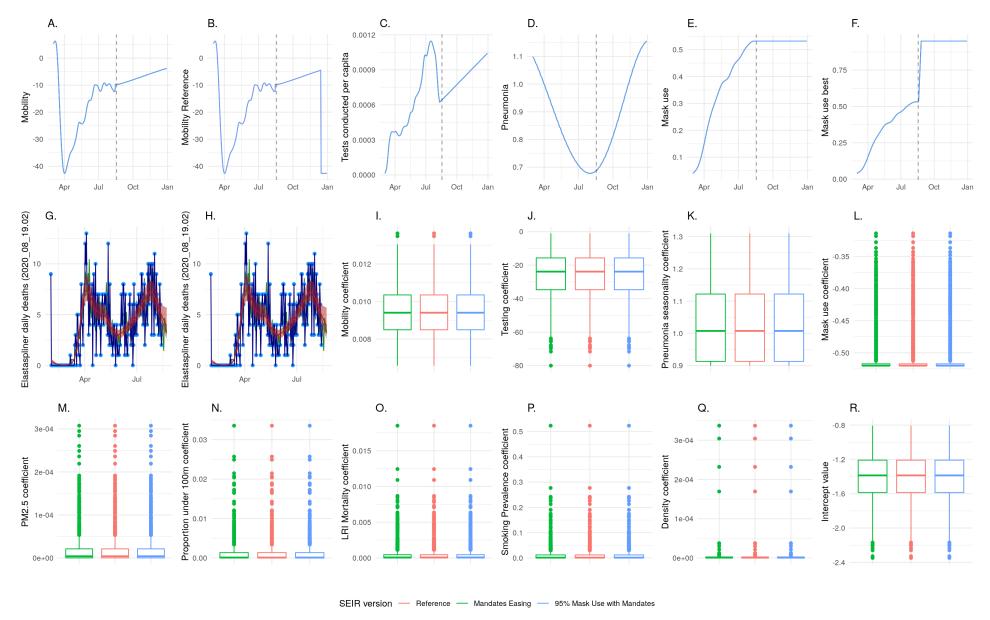
# 109 Washington except for King, Snohomish, and Spokane Counties: SEIR fit comparison



Washington except for King, Snohomish, and Spokane Counties: SEIR fit comparison. A: predicted R effective for each model through December 31. B: predicted SEIR beta parameter. C: residual of predicted beta and the observed value calculated directly from infection data over time. D: histogram of residual values for beta. Panels A, B, C, and D are all displayed in log space, reflecting the space in which the SEIR model is fit. E: predicted daily infections from each model through December 31. F: predicted daily deaths from each model through December 31. G: predicted cumulative infections through December 31, as a proportion of the total population. H: predicted cumulative deaths through December 31. In panels E, F, G, and H, reported death and infections are plotted alongside model predictions in

light blue.

# Washington except for King, Snohomish, and Spokane Counties: Covariate fits and regression coefficients



Washington except for King, Snohomish, and Spokane Counties: Covariate fits and regression coefficients. A-F: Line plots showing predicted covariate time trends for A) mobility in the absence of additional mandates; B) mobility with additional mandates applied; C) diagnostic testing per capita; D) pneumonia seasonality; E) mask use per capita, and; F) mask use in a scenario where adherence increases to 95% of the population. G-H: COVID mortality data generated from

reported daily deaths (blue); estimated based on reported hospitalizations (purple); estimated from reported cases (green); and via a spline fit through all available data types (red, 95% UI in pink). I-R: Box plots showing 1,000 draws of fixed effect coefficients in a multivariate regression fit to log(beta).