

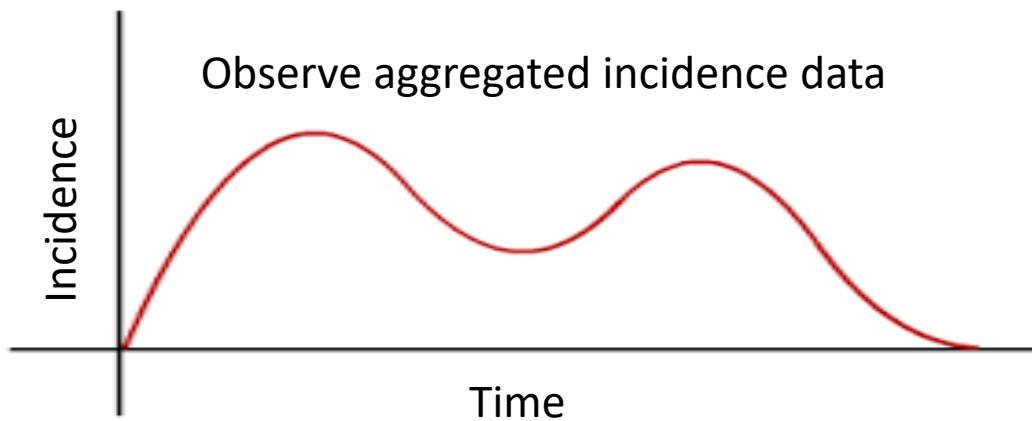
Forecasting epidemiological patterns using multi-scale semi-mechanistic models

Privacy and Ethics in Pandemic Data Collection and Processing

ICERM @ Brown University
19 January 2023

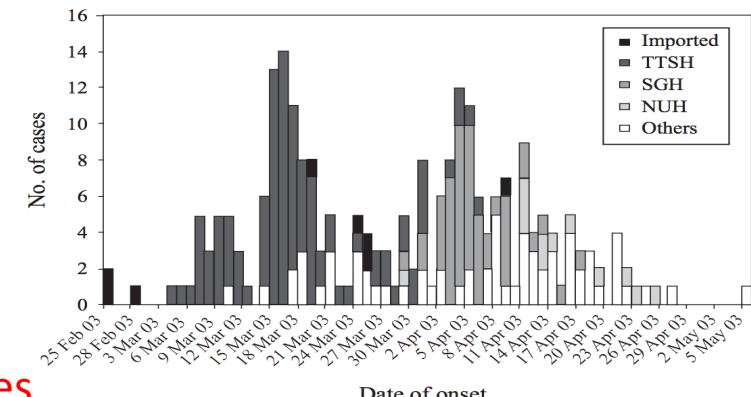
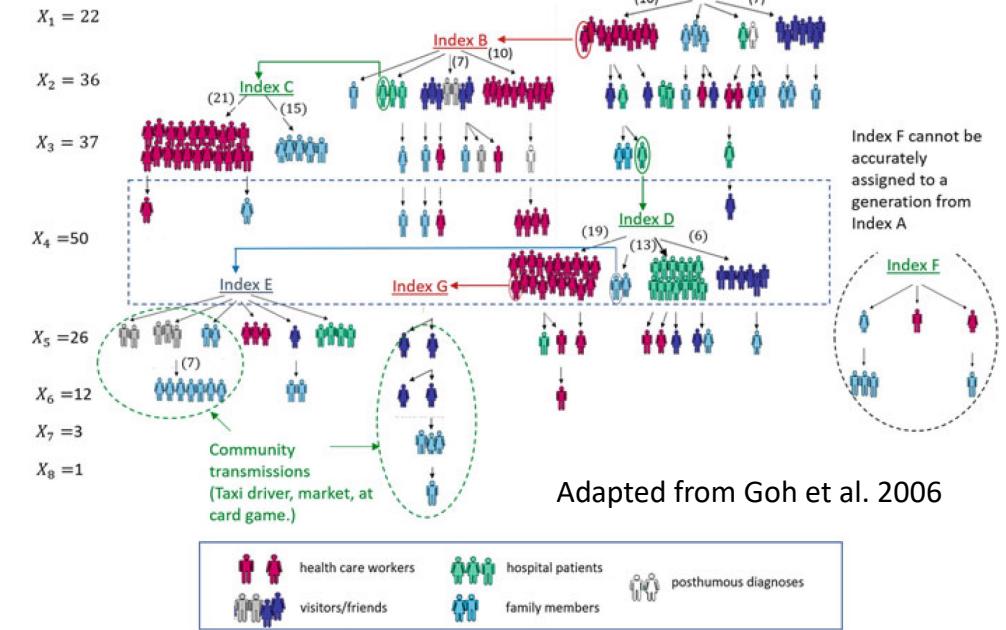
Gerardo Chowell, PhD

Inferring the structure of aggregate epidemiological patterns

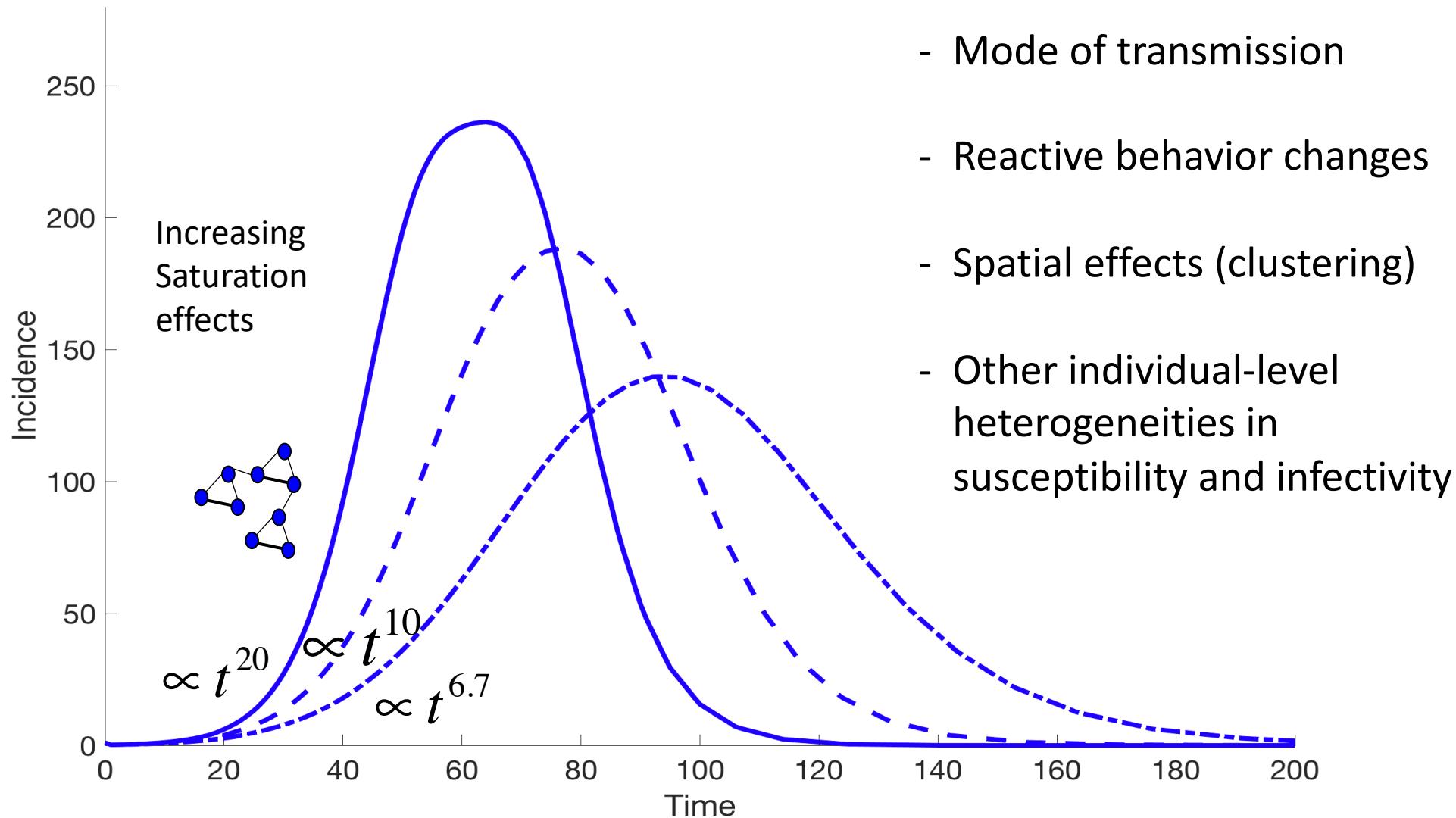


More than one transmission tree could give rise to very similar epidemic curves

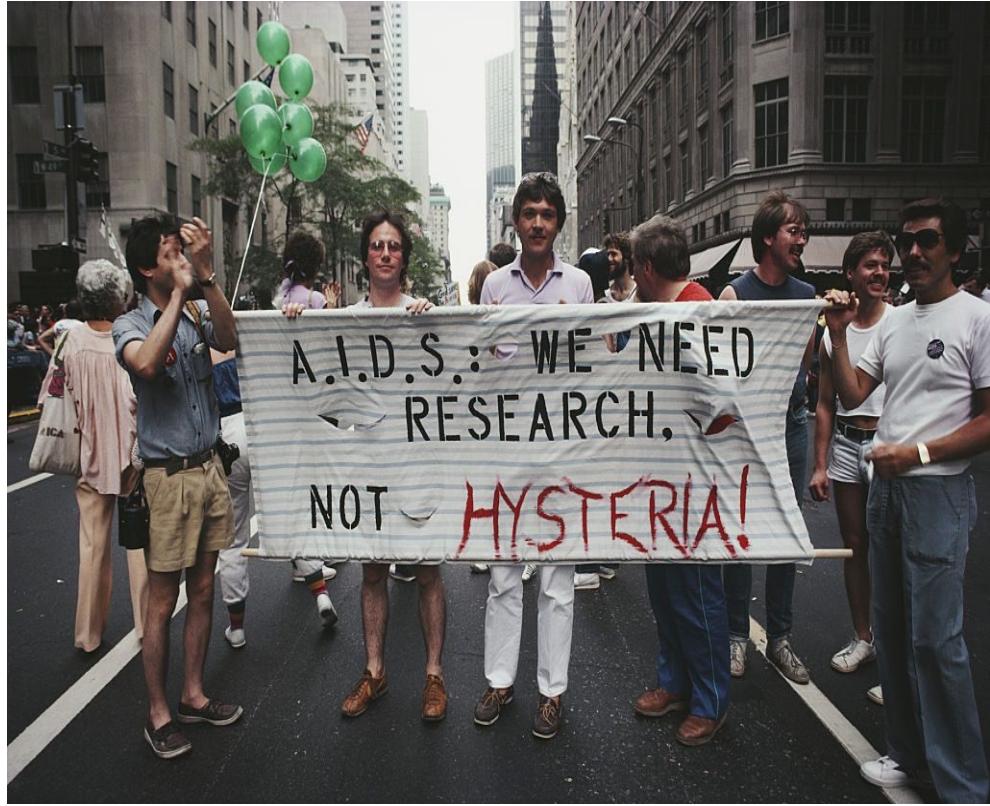
2003 SARS outbreak in Singapore



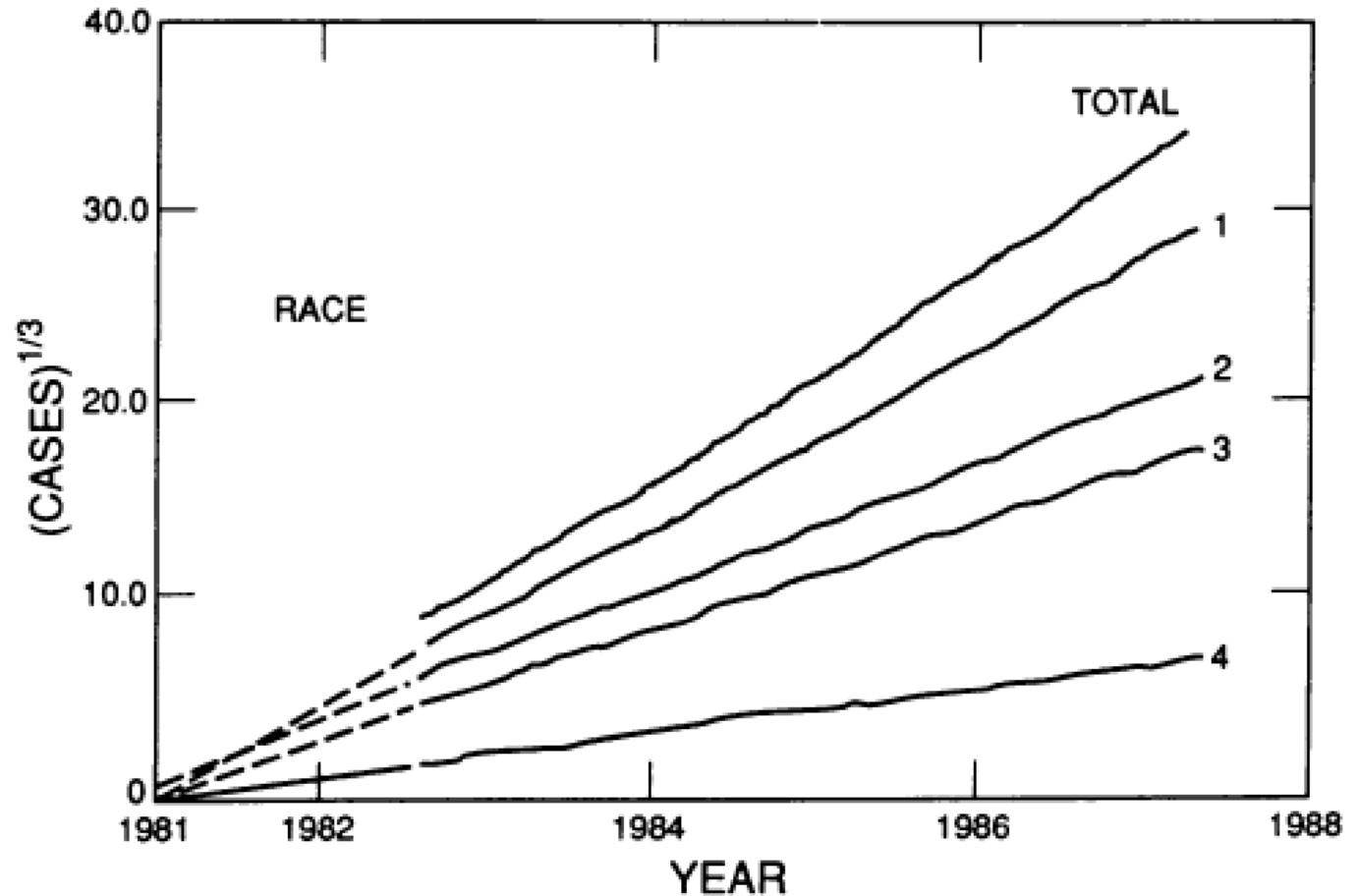
Real epidemics exhibit variable epidemic growth scaling



The initial phase of the HIV/AIDS epidemic in the United States



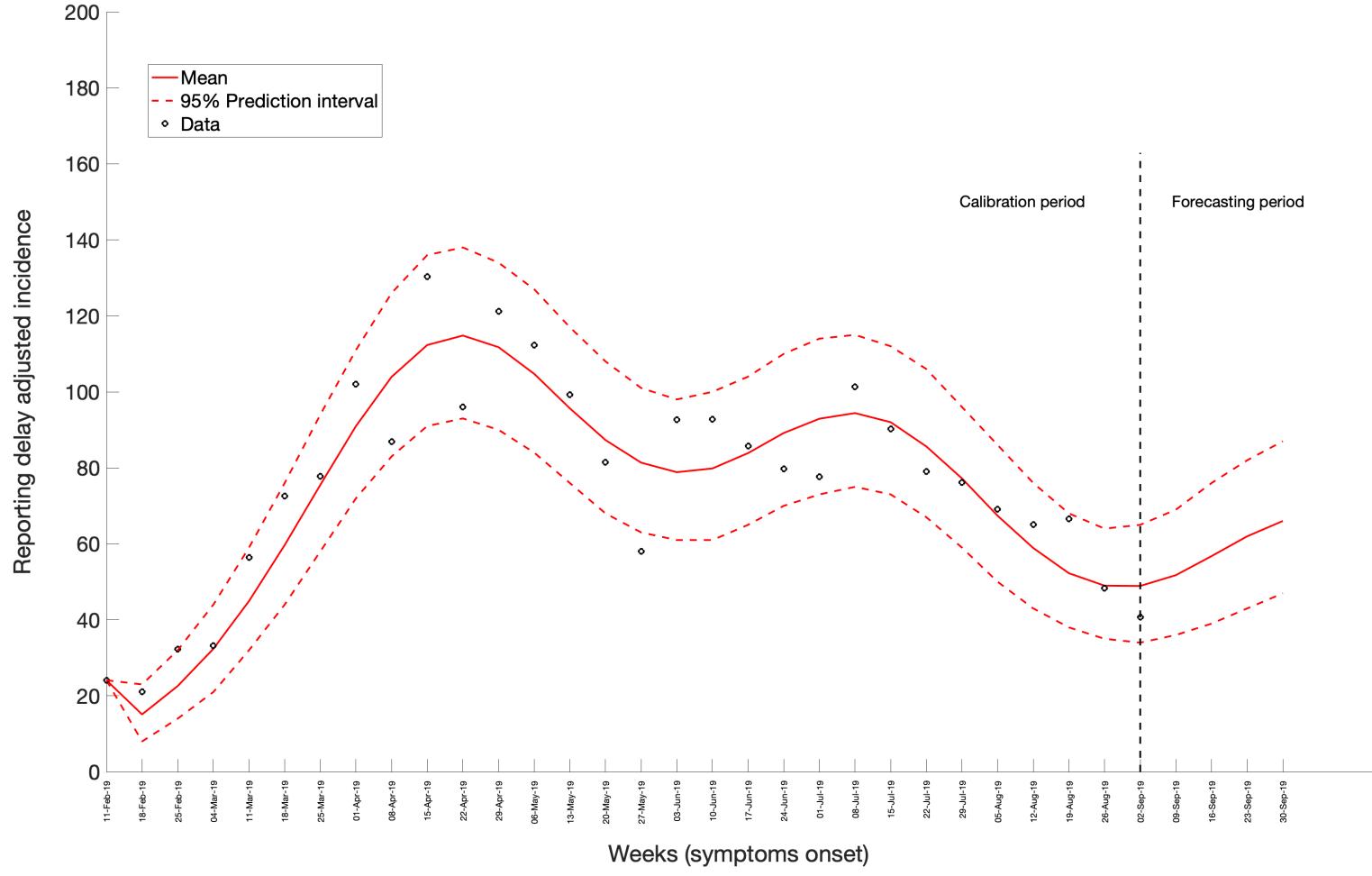
<https://www.pbs.org/newshour/science/america-hiv-outbreak-origins-nyc-gaetan-dugas>



1=White; 2= Black; 3=Hispanic; 4=unknown

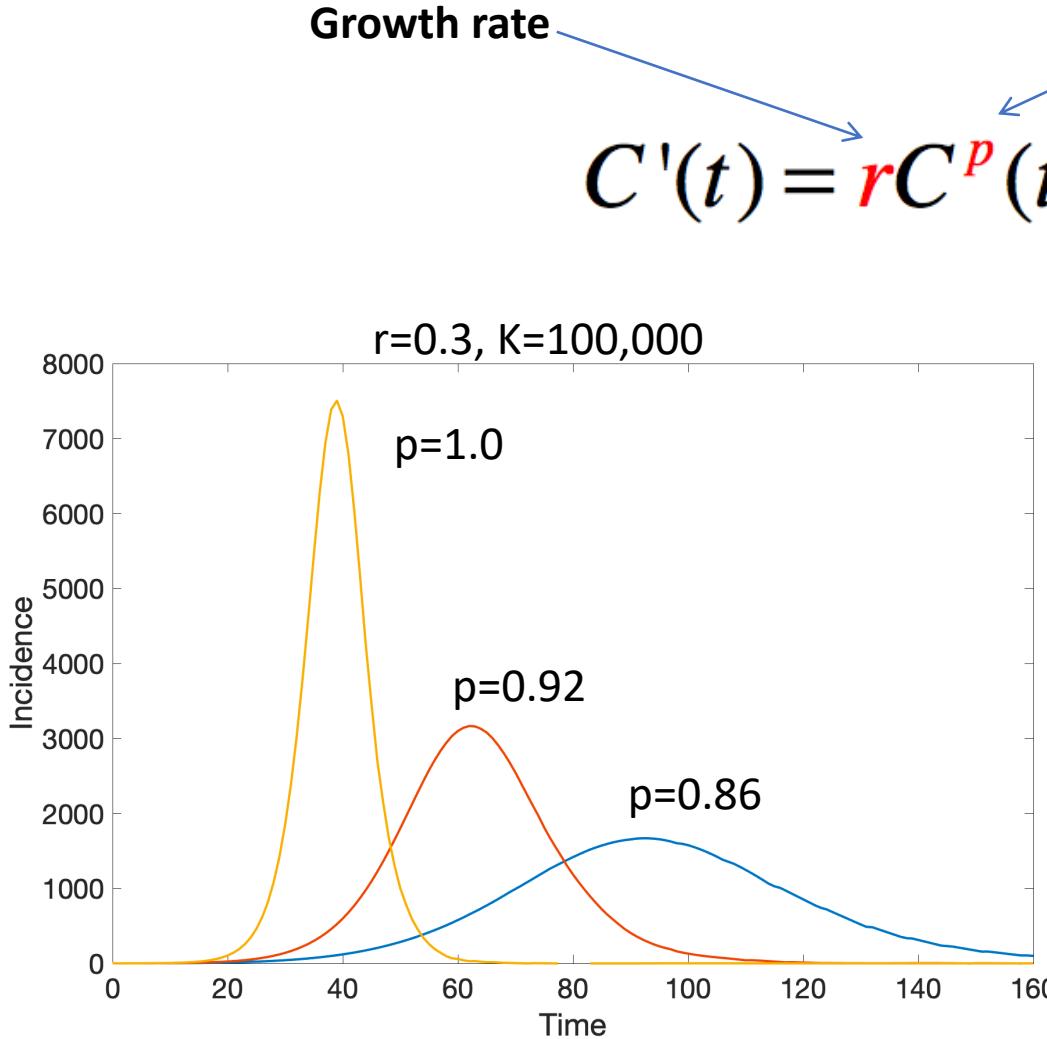
Colgate et al. PNAS 1989

Multimodal epidemics



- Aggregation of multiple underlying spreading mechanisms.
- Transmission occurs in high-risk groups first before affecting others.
- Transmission in different geographic areas occurs at different times.
- Emergence of new variants

Building block: generalized logistic growth model



$$C'(t) = r C^p(t) \left[1 - \left(\frac{C(t)}{K} \right) \right]$$

Growth rate

Deceleration of growth parameter

Epidemic size

Where:

- $C'(t)$ describes the incidence curve over time t
- r is a positive parameter denoting the growth rate
- $p \in [0,1]$ is an “deceleration” growth parameter
- K is the final epidemic size parameter

Overlapping sub-epidemic wave model

- Each sub-epidemic is modelled by a generalized logistic growth model
- An epidemic wave comprising of a set of n overlapping sub-epidemics is modelled using coupled differential equations

$$\frac{dC_i(t)}{dt} = rA_{i-1}(t)C_i(t)^p \left(1 - \frac{C_i(t)}{K_i}\right)$$

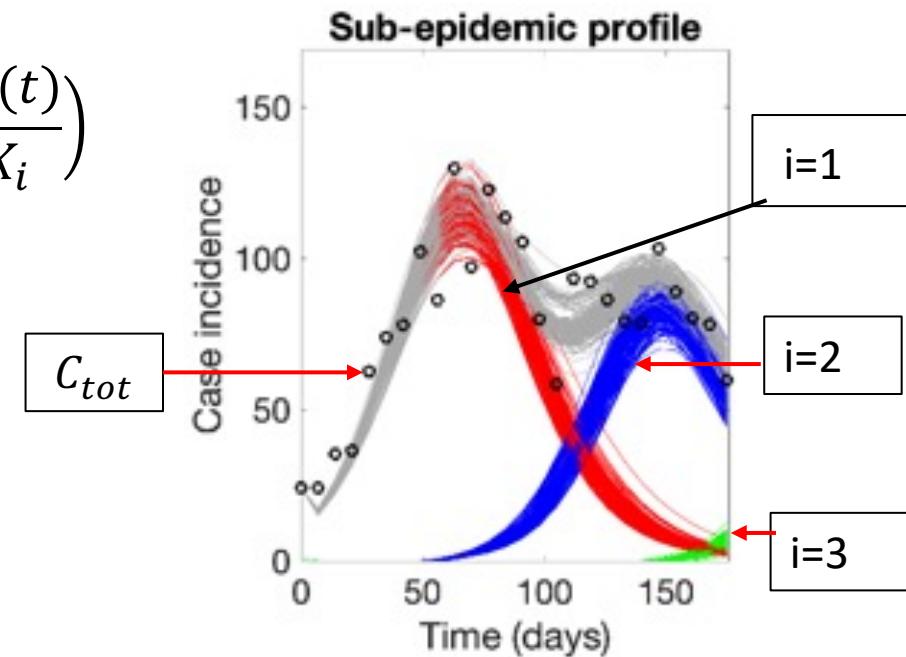
Where,

C_i = cumulative number of infections for sub-epidemic i ,

K_i = size of the i_{th} sub-epidemic

A_i = indicator variable

$$A_i(t)=f(x) = \begin{cases} 1 & C_i(t) > C_{thr} \\ 0 & Otherwise \end{cases} \quad i = 1,2,3 \dots \dots n$$



Modeling sub-epidemic sizes

Assuming the subsequent sub-epidemic sizes decline exponentially we have,

$$K_i = K_0 e^{-q(i-1)}$$

Where,

K_0 is the size of the initial sub-epidemic ($K_i = K_0$)

The total size of the epidemic wave composed of n overlapping sub-epidemics is given by:

$$K_{tot} = \sum_{i=1}^n K_0 e^{-q(i-1)} = \frac{K_0(1 - e^{-qn})}{1 - e^{-q}}$$

Where,

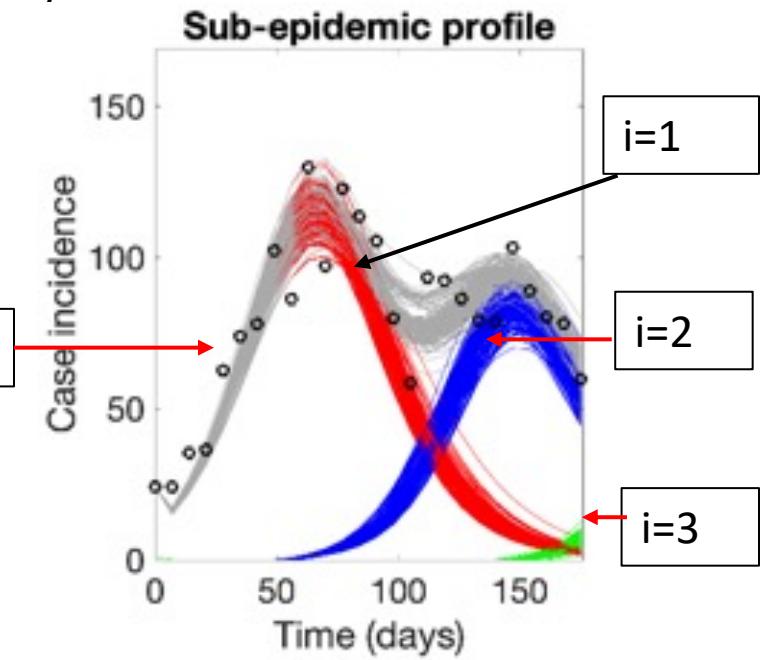
q = interventions or behavior change, $q > 0$ and n = number of sub-epidemics

$$K_{tot} = nK_0 , \quad \text{when } q = 0$$

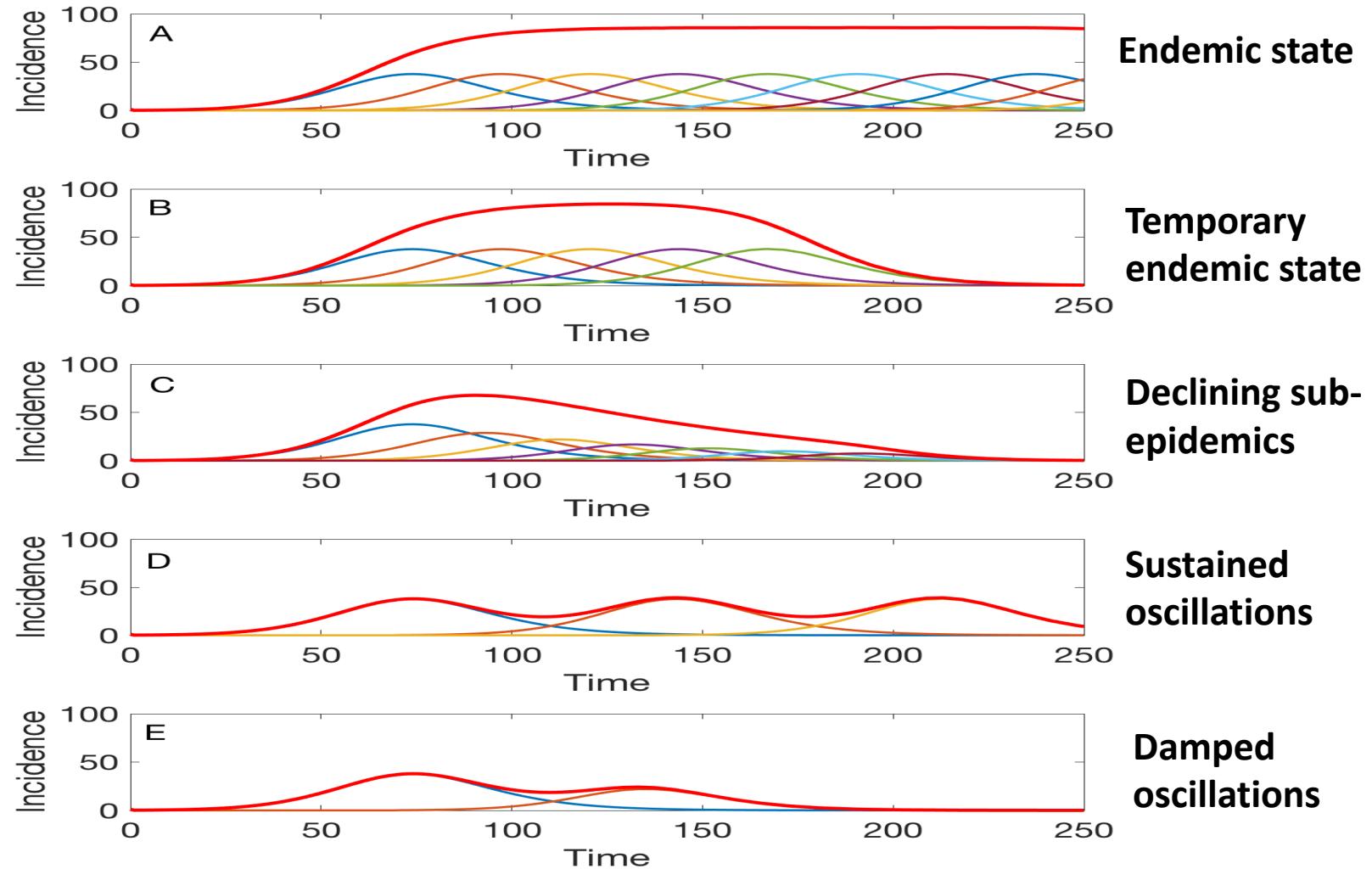
Where K_{tot} = total epidemic size

$$C_{tot}(t) = \sum_{i=1}^n C_i(t)$$

Where $C_{tot}(t)$ = cumulative curve of epidemic wave



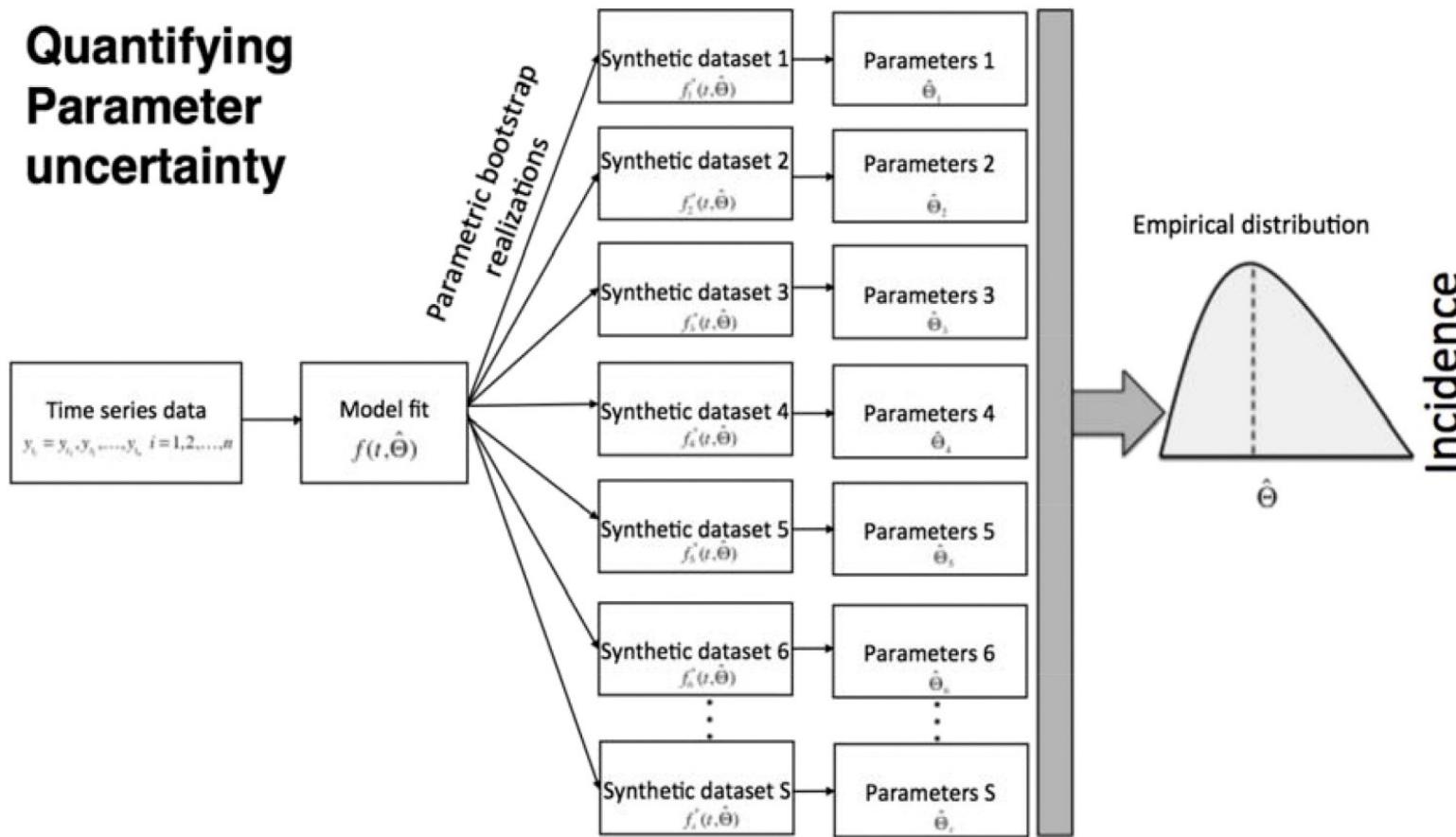
Representative epidemic waves composed of overlapping sub-epidemics



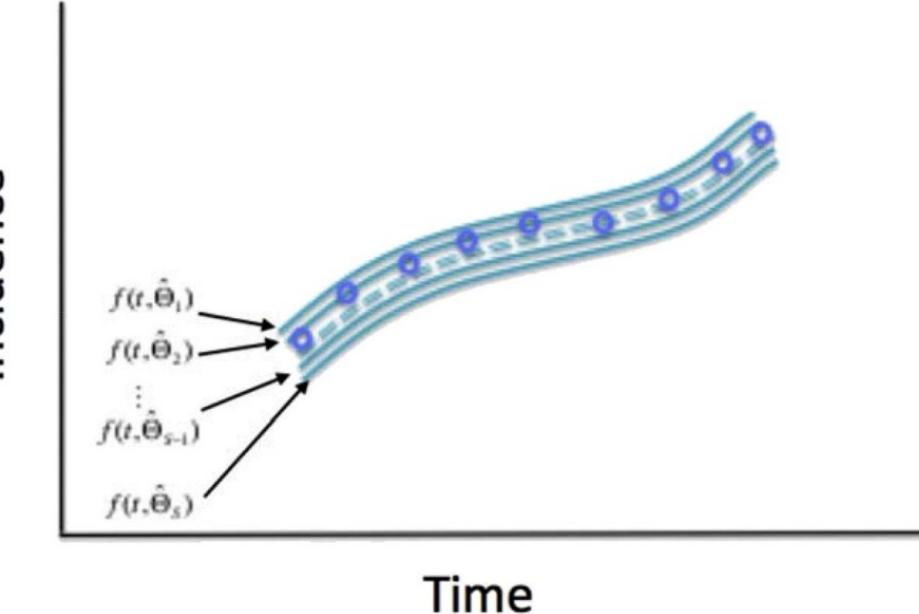
Number of model parameters is $n > 1$

Parameter estimation

Quantifying Parameter uncertainty

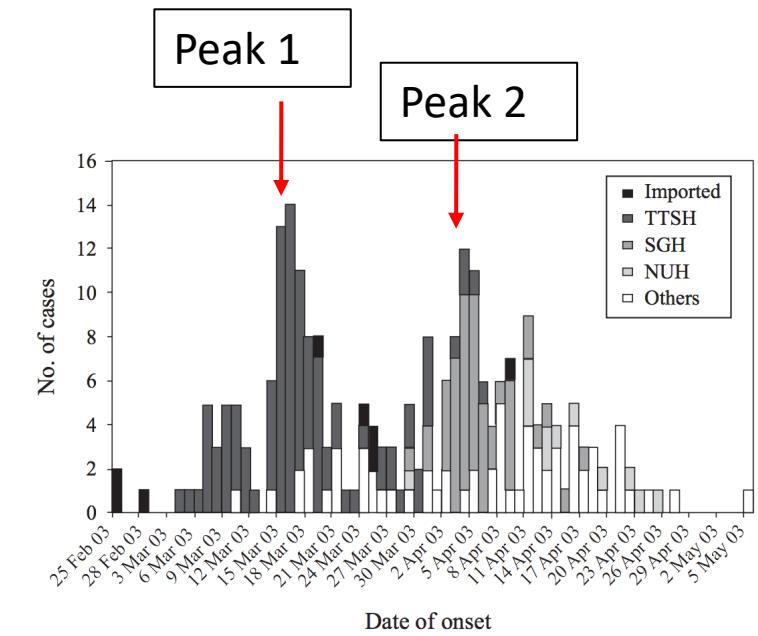
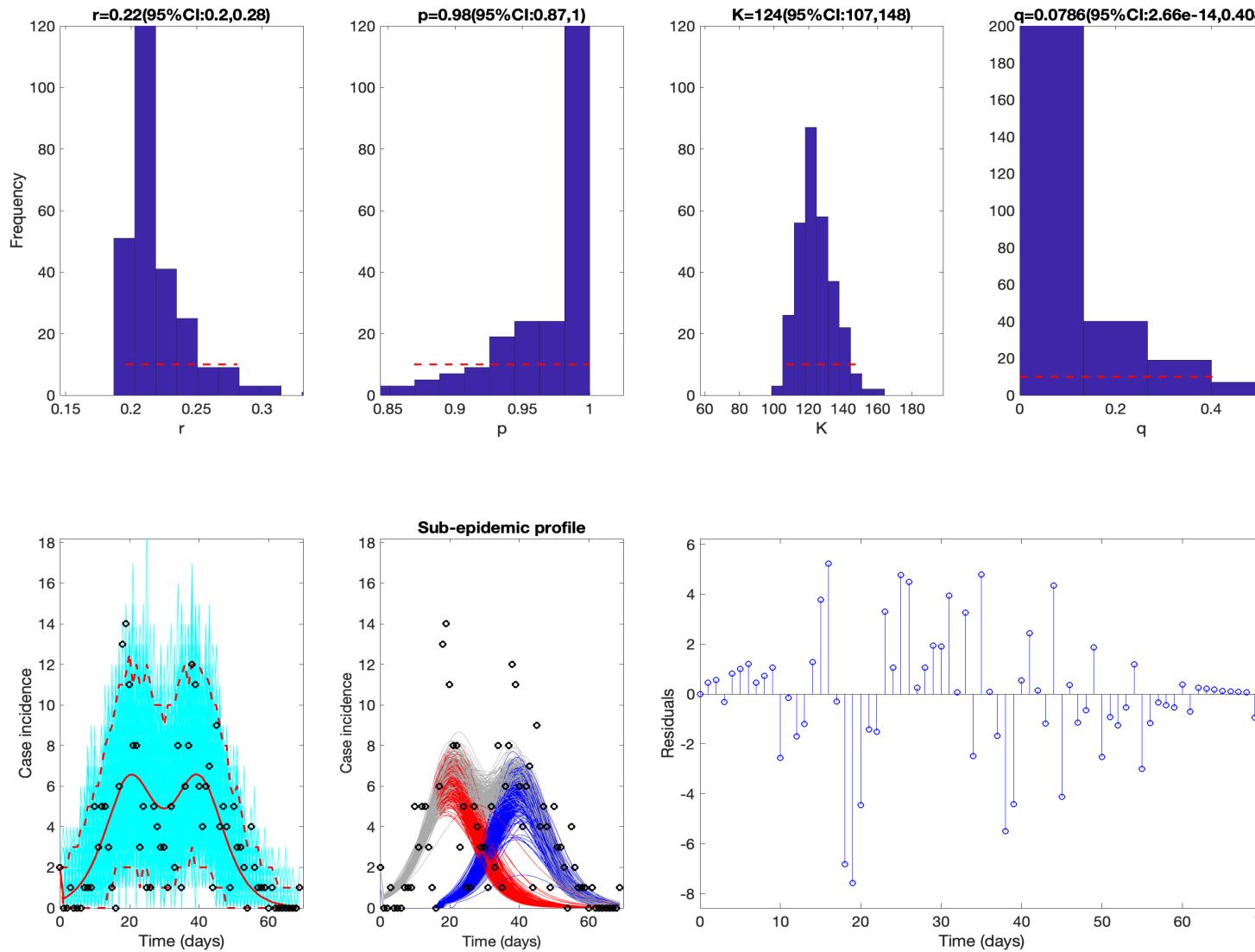


Model parameters are estimated by fitting the model to the aggregated incidence curve

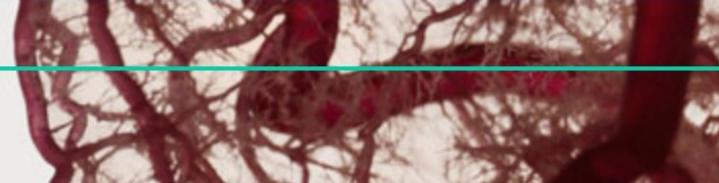


Nonlinear least squares fitting or maximum likelihood estimation with parametric bootstrapping.

SARS outbreak in Singapore, 2003



Hence two sub-epidemics in the sub-epidemic profile



Research article | [Open Access](#) | [Published: 22 August 2019](#)

A novel sub-epidemic modeling framework for short-term forecasting epidemic waves

[Gerardo Chowell](#)  [Amna Tariq](#) & [James M. Hyman](#)

[BMC Medicine](#) **17**, Article number: 164 (2019) | [Cite this article](#)

10k Accesses | **72** Citations | **11** Altmetric | [Metrics](#)

Spatial wave sub-epidemic framework - MATLAB Toolbox

The screenshot shows a GitHub repository page for the 'spatial_wave_subepidemic_framework' project. The repository is public and owned by 'gchowell'. The main page displays the repository's structure, recent commits, and a detailed README.md file.

Repository Structure:

- Code: main branch, 1 branch, 0 tags
- Issues: 0
- Pull requests: 0
- Actions: 0
- Projects: 0
- Wiki: 0
- Security: 0
- Insights: 0
- Settings: 0

Commits:

Author	Commit Message	Date	Commits
gchowell	Add files via upload	e4c6cad last week	68 commits
	spatialWave_subepidemicFramewo...	Add files via upload	last week
	LICENSE	Initial commit	2 months ago
	README.md	Update README.md	2 months ago

README.md Content:

spatial_wave_subepidemic_framework

A Matlab toolbox for fitting and forecasting epidemic trajectories using the spatial wave sub-epidemic framework

It carries out the following tasks:

- fitting models to time series data,
- estimation of parameters with quantified uncertainty,
- plotting the fits of the top-ranked models,
- plotting the AICc values of the top-ranked models,
- generates forecasts of the top-ranked models
- generates ensemble forecasts based on the top-ranked models.

Additional features include:

- fitting models using different parameter estimation approaches (least-squares, maximum likelihood estimation),

About:

A Matlab toolbox for fitting and forecasting epidemic trajectories using the spatial wave sub-epidemic framework

Readme:

GPL-3.0 license:

Stars: 0

Watching: 1

Forks: 0

Releases:

No releases published

[Create a new release](#)

Packages:

No packages published

[Publish your first package](#)

Languages:

MATLAB 99.8% Objective-C 0.2%

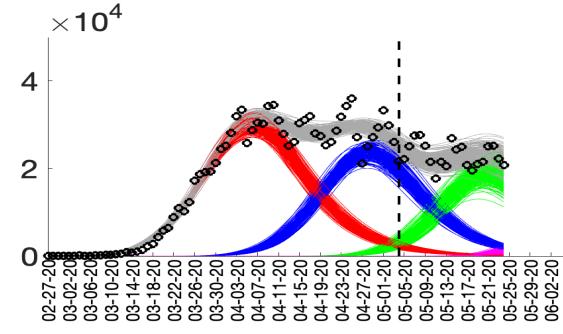
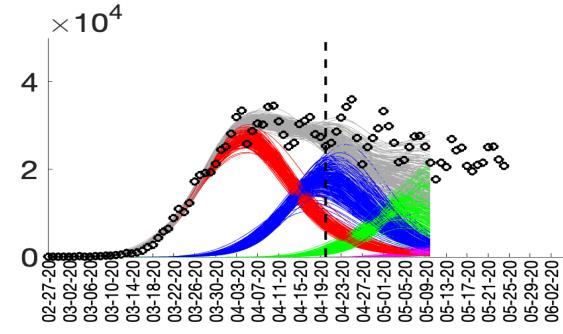
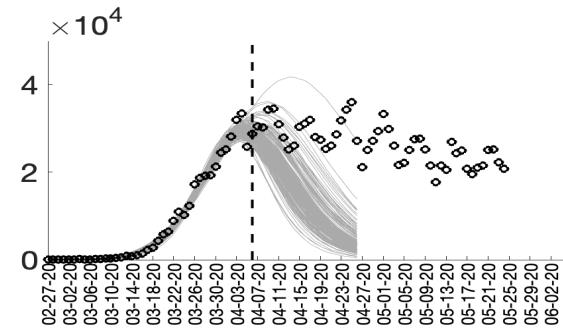
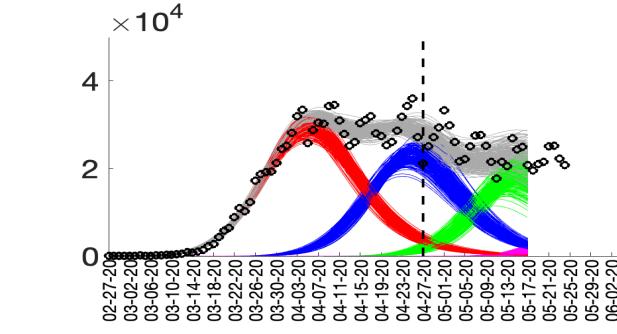
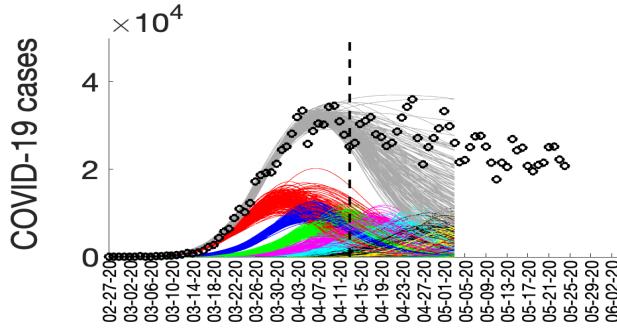
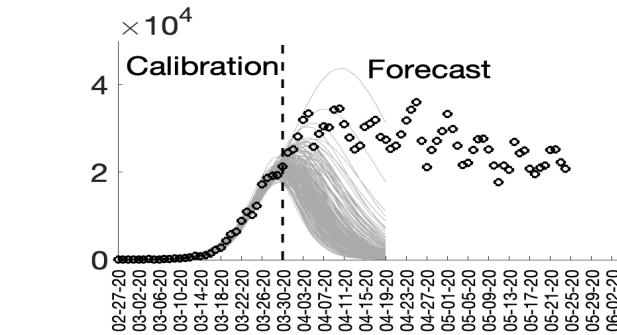
https://github.com/gchowell/spatial_wave_subepidemic_framework

Real-time forecasts of the ongoing COVID-19 pandemic



<https://publichealth.gsu.edu/research/coronavirus/>

Sub-epidemic wave model forecasts: USA



Performance metrics

Mean Absolute Error

$$\text{MAE} = \frac{1}{n} \sum_{i=1}^n |f(t_i, \hat{\Theta}) - y_{t_i}|$$

y_{t_i} = time series of incident cases
describing epidemic wave

t_i = time points of time series data
 $f(t_i, \hat{\Theta})$ = model fit

Mean Squared Error

$$\text{MSE} = \frac{1}{n} \sum_{i=1}^n (f(t_i, \hat{\Theta}) - y_{t_i})^2$$

Coverage of the 95% Prediction Interval

$$\frac{1}{n} \sum_{t=1}^n \mathbb{1}\{y_t > L_t \cap y_t < U_t\}$$

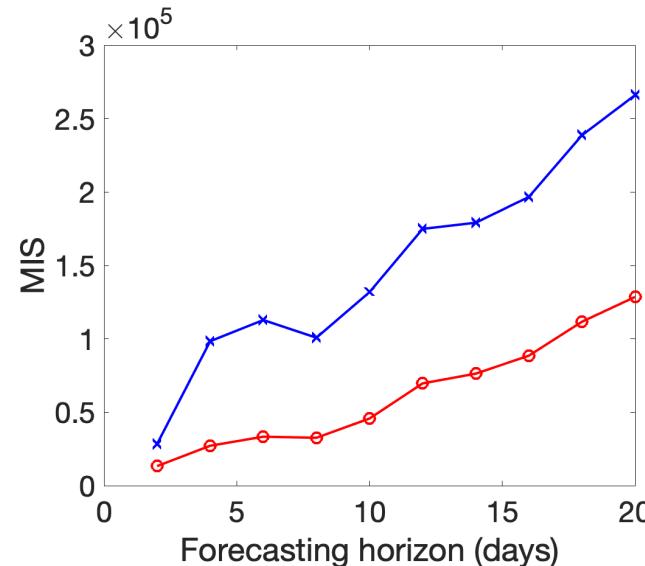
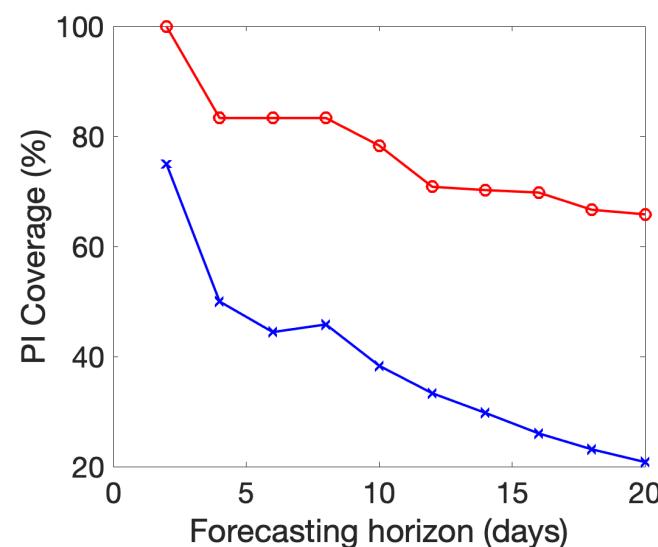
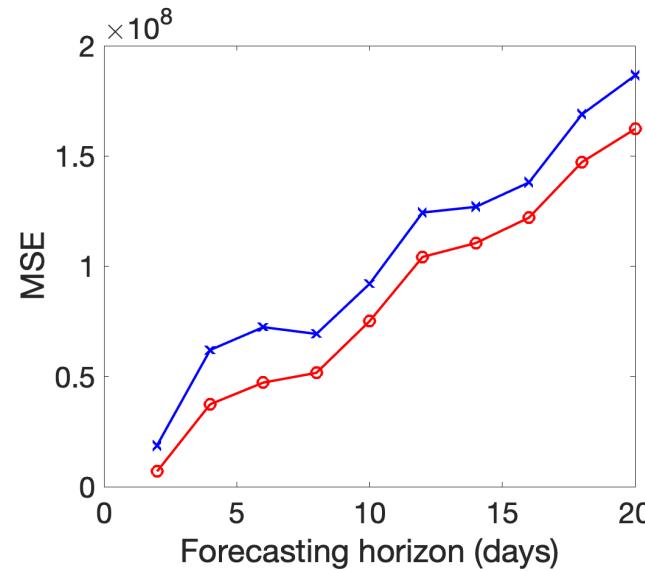
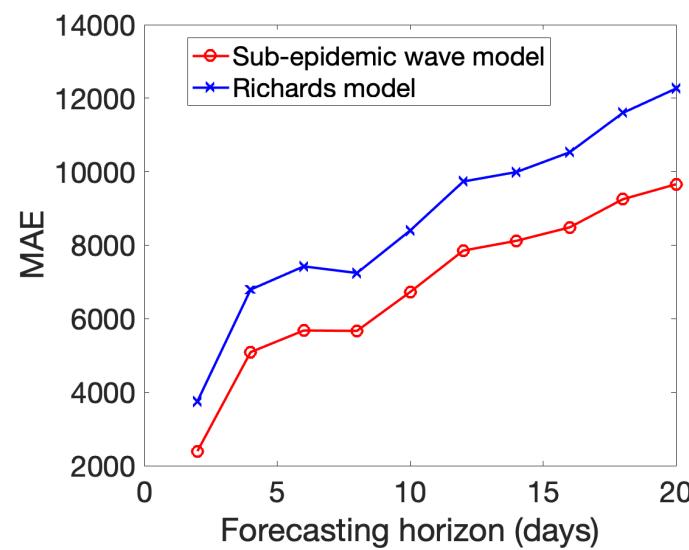
Weighted Interval Score

$$WIS_{\alpha_0:K}(F, y) = \frac{1}{K + \frac{1}{2}} \cdot (w_0 \cdot |y - m| + \sum_{k=1}^K w_k \cdot IS_{\alpha_k}(F, y))$$

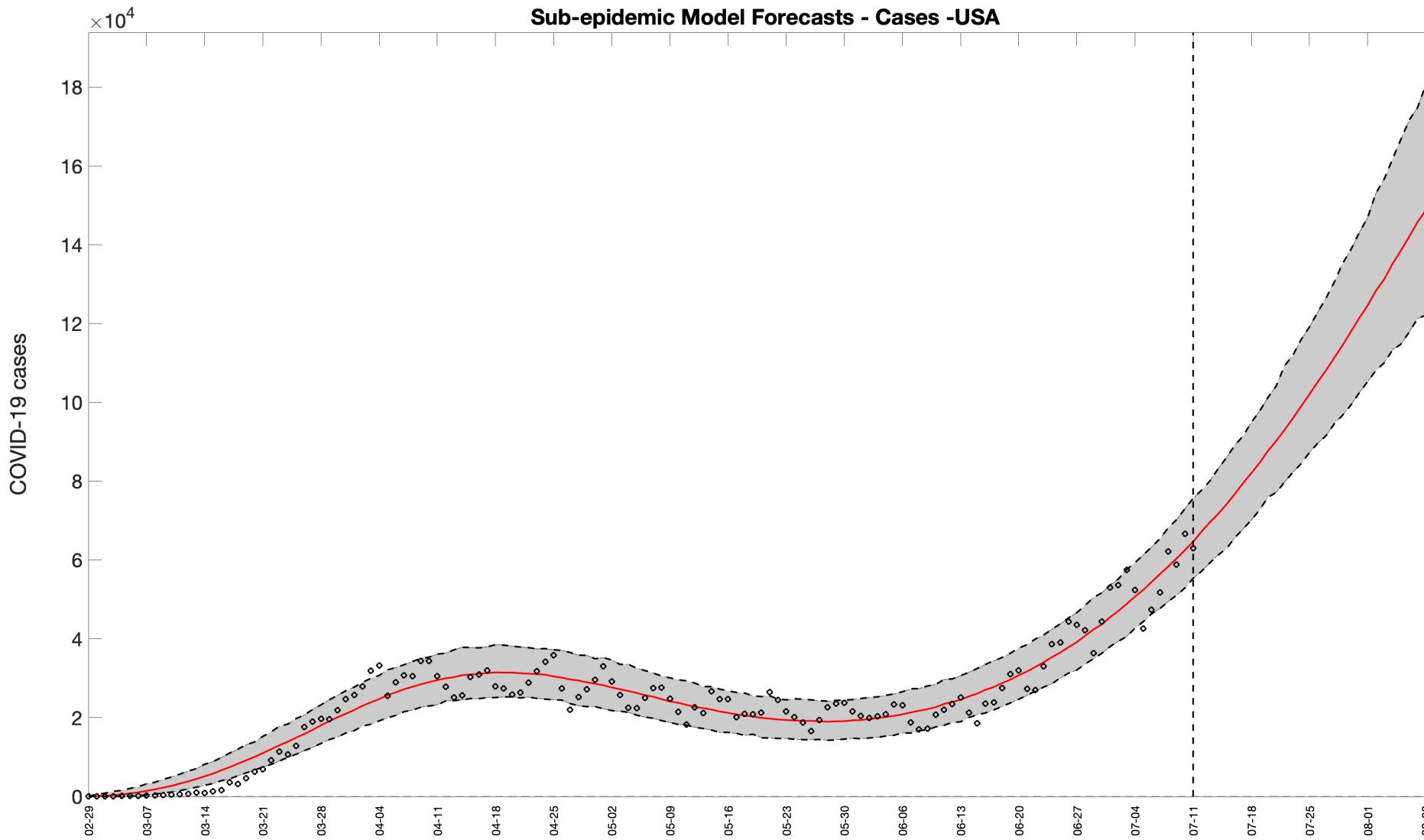
L_t = lower bound of 95% prediction interval

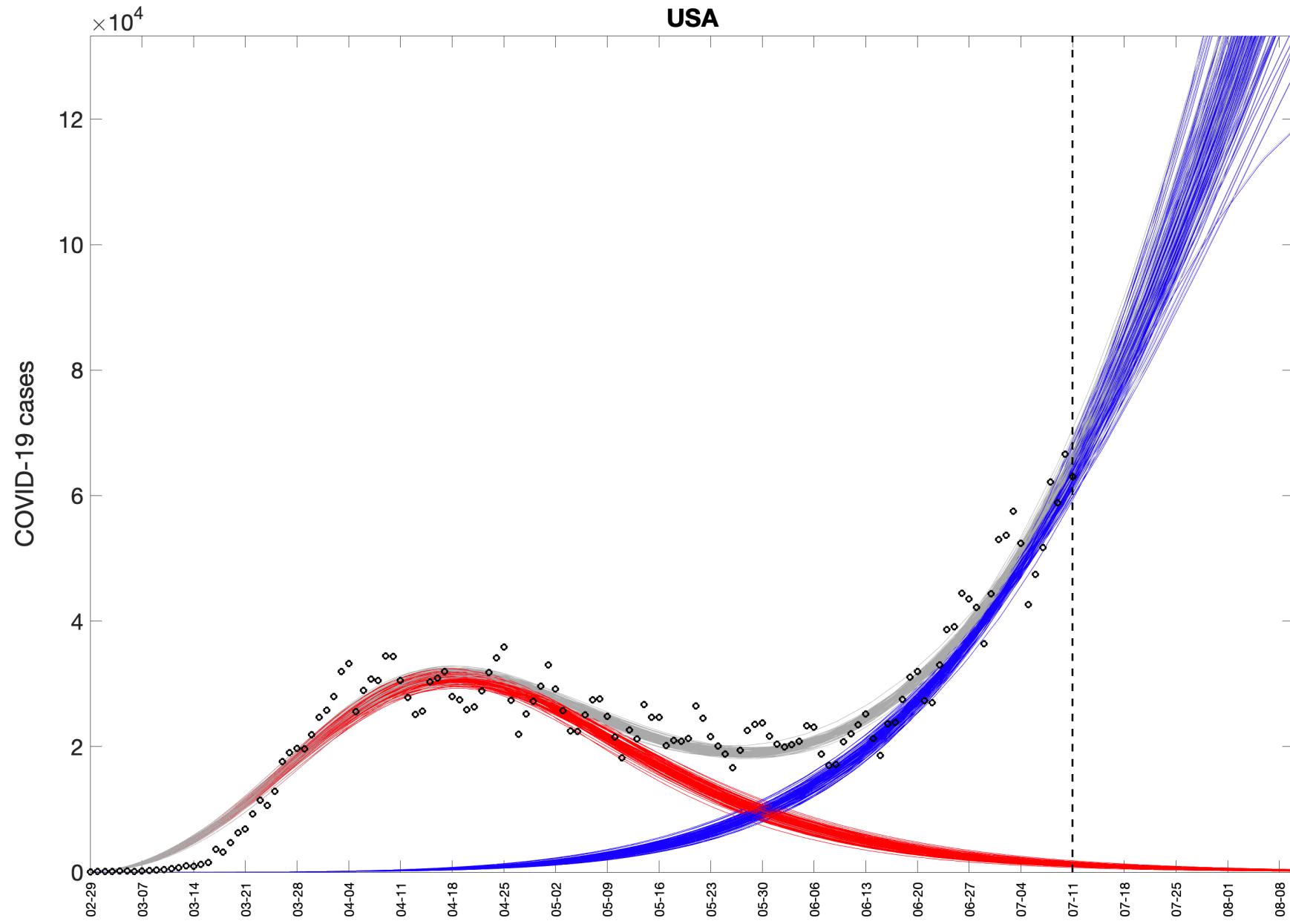
U_t = Upper bound of 95% prediction interval

Sub-epidemic wave model performance - USA



COVID-19 Resurgence after relaxation of social distancing - USA





n-subepidemic modeling framework

- Each sub-epidemic is modelled by a generalized logistic growth model
- An epidemic wave comprising of a set of n overlapping sub-epidemics is modelled using coupled differential equations
- $$\frac{dC_i(t)}{dt} = C_i'(t) = A_i(t)r_i C_i^{p_i}(t) \left(1 - \frac{C_i(t)}{K_{0i}}\right)$$

Where,

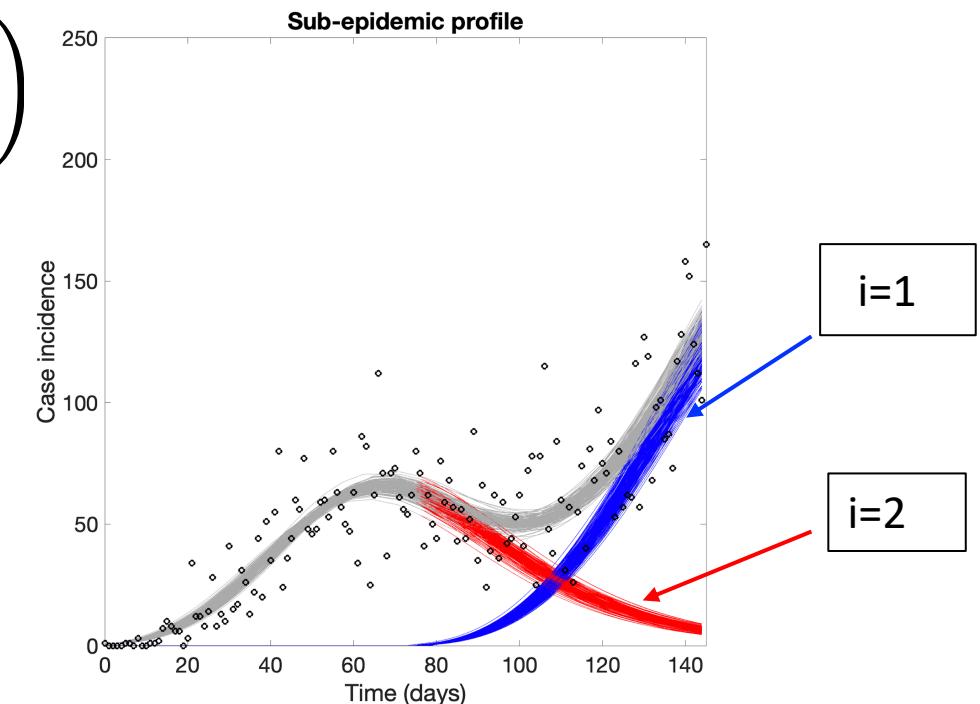
C_i = cumulative number of infections for sub-epidemic i ,

K_i = size of the i_{th} sub-epidemic

A_i = indicator variable

$$A_i(t)=f(x) = \begin{cases} 1 & C_i(t) > C_{thr} \\ 0 & Otherwise \end{cases} \quad i = 1, 2, 3, \dots, n$$

Number of model parameters is $3n+1$



Top-ranking K sub-epidemic models via model selection

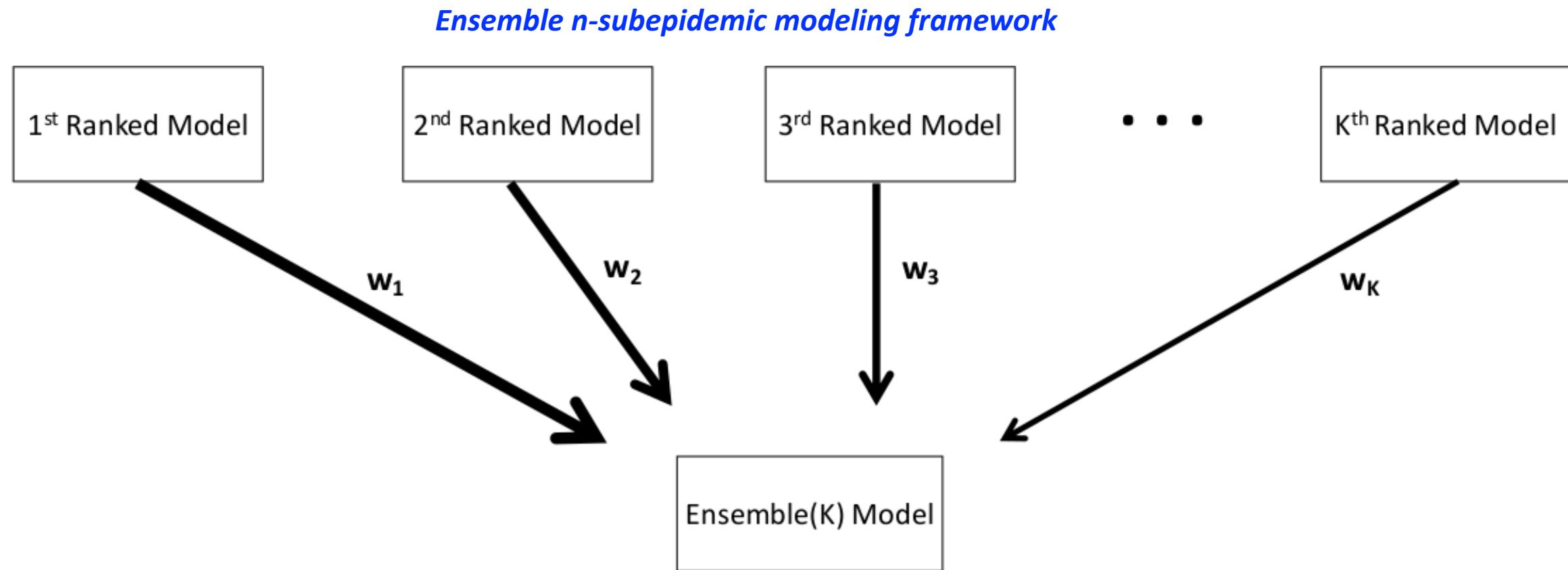
We considered a set of n -subepidemic models with $1 \leq n \leq 2$ and different values of C_{thr} . Then we ranked them from best to worst according to the AIC_c which is given by:

$$AIC_c = n_d \log(SSE) + 2m + \frac{2m(m+1)}{n_d - m - 1}$$

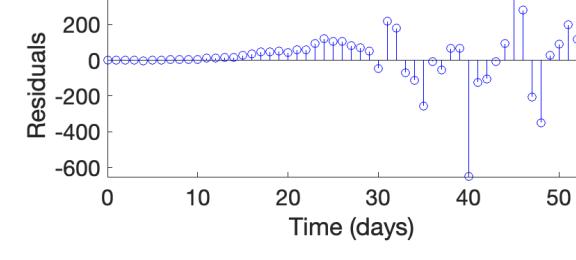
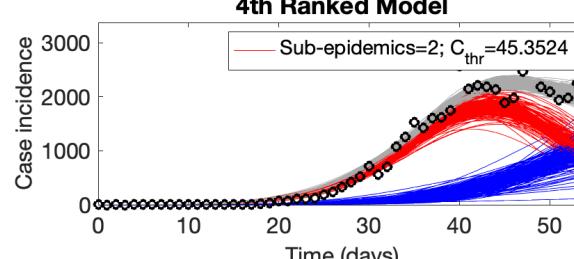
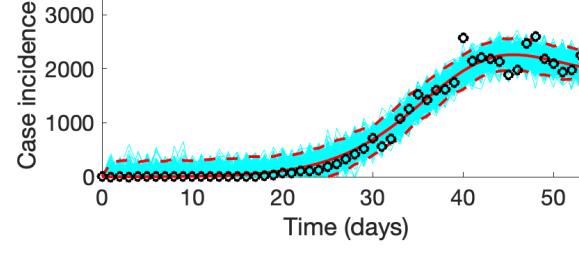
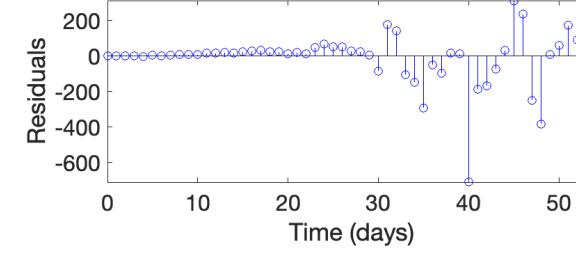
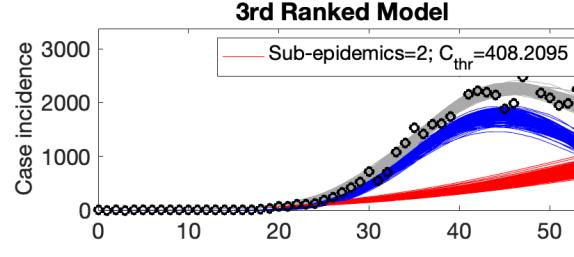
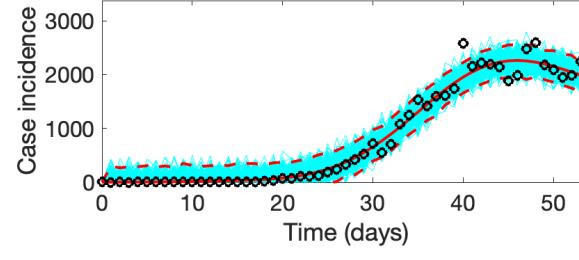
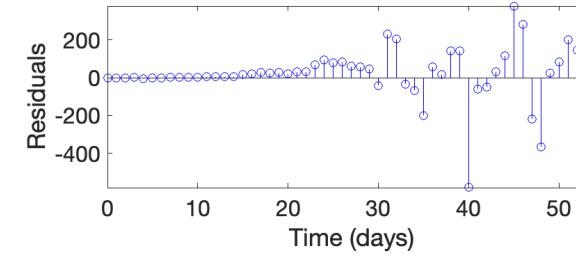
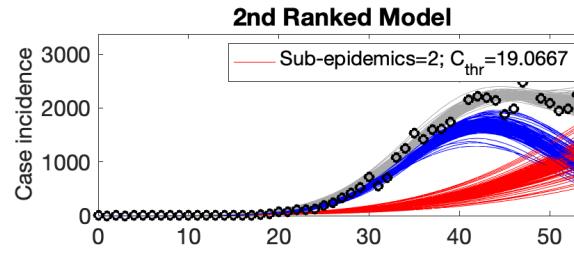
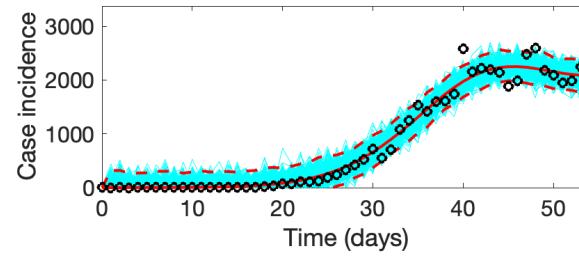
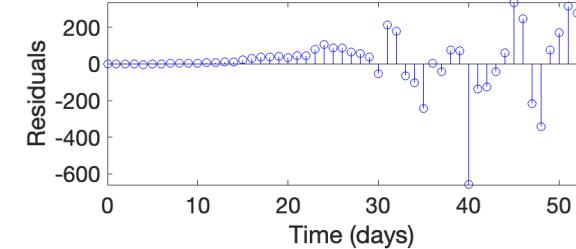
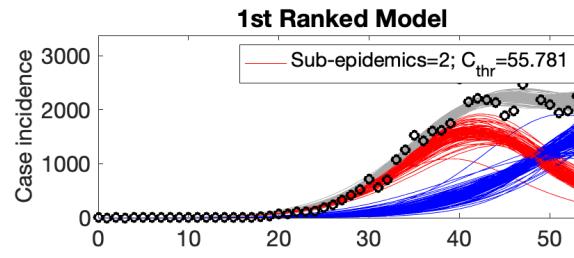
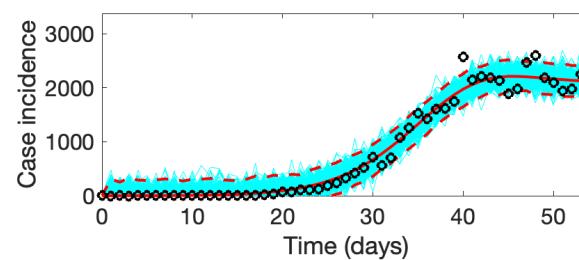
where $SSE = \sum_{j=1}^{n_d} (f(t_j, \hat{\Theta}) - y_{t_j})^2$, $m = 3n + 1$ is the number of model parameters and n_d is the number of data points.

Ensemble(K) models from top-ranking K sub-epidemic models

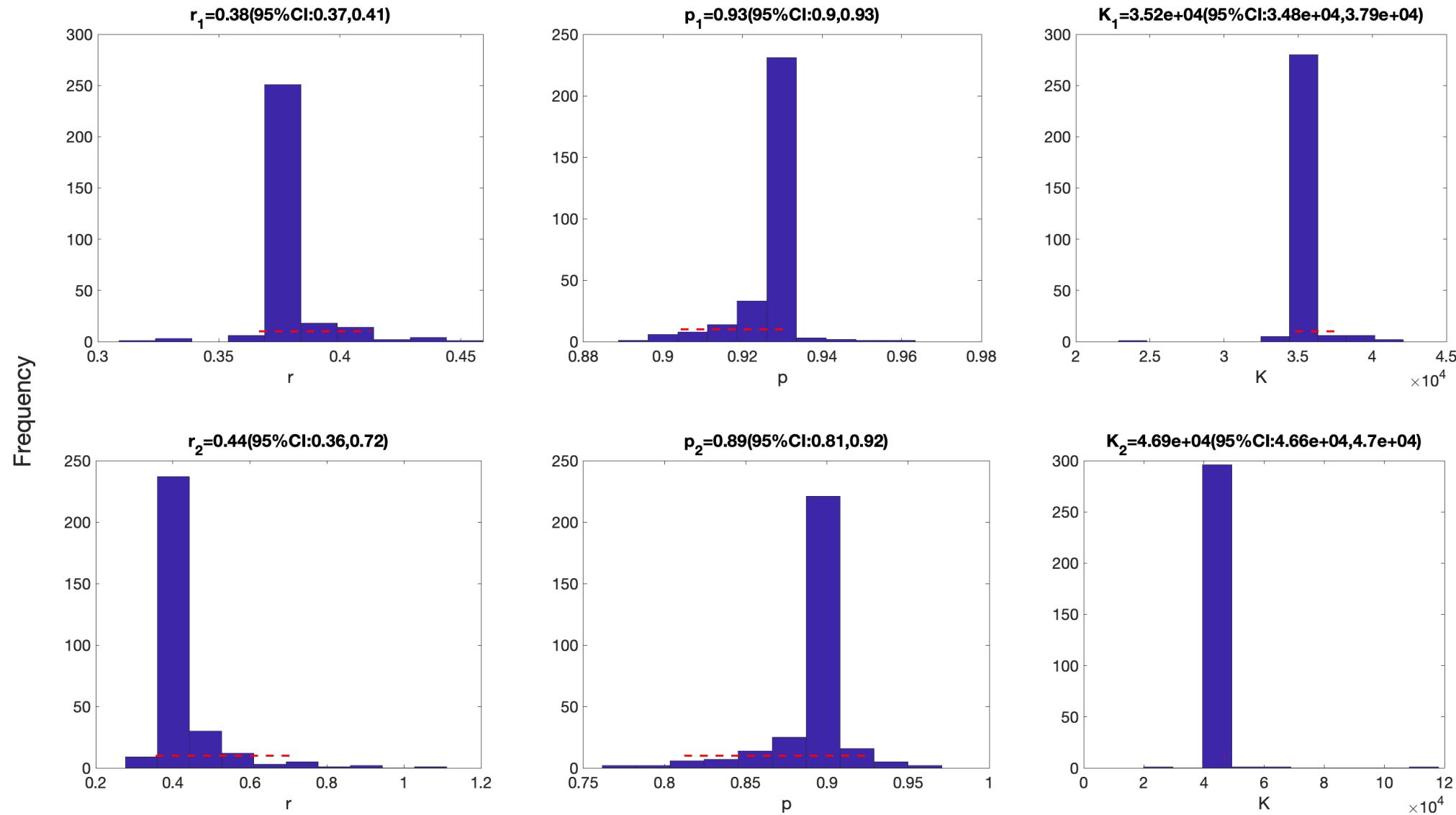
We generate ensemble models from the weighted combination of the highest-ranking sub-epidemic models as deemed by the AIC_C. An ensemble derived from the top-ranking "K" models is denoted by Ensemble(K).



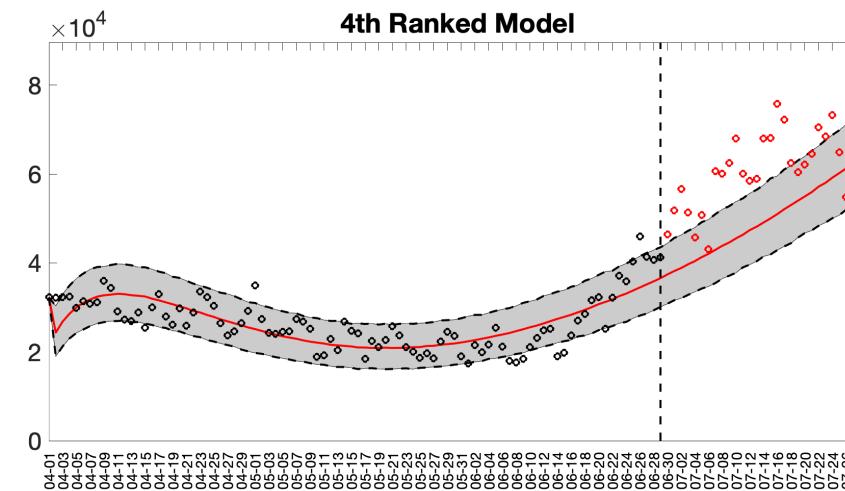
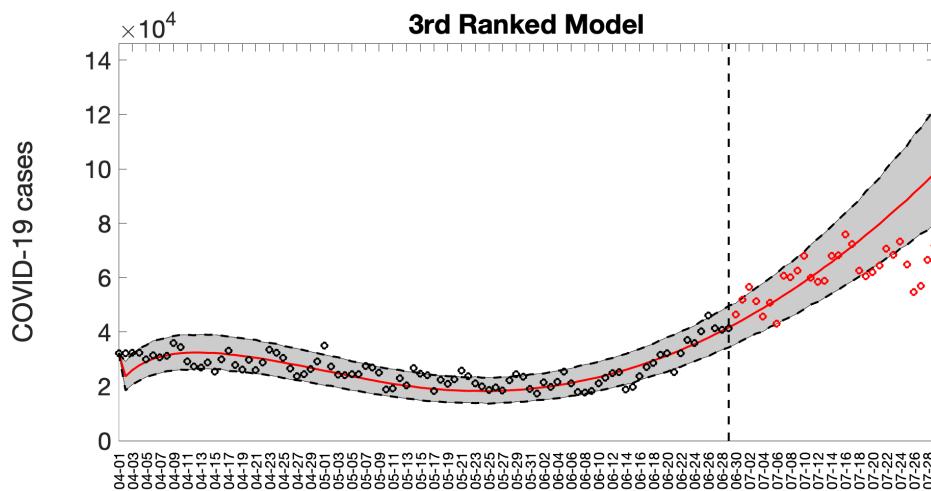
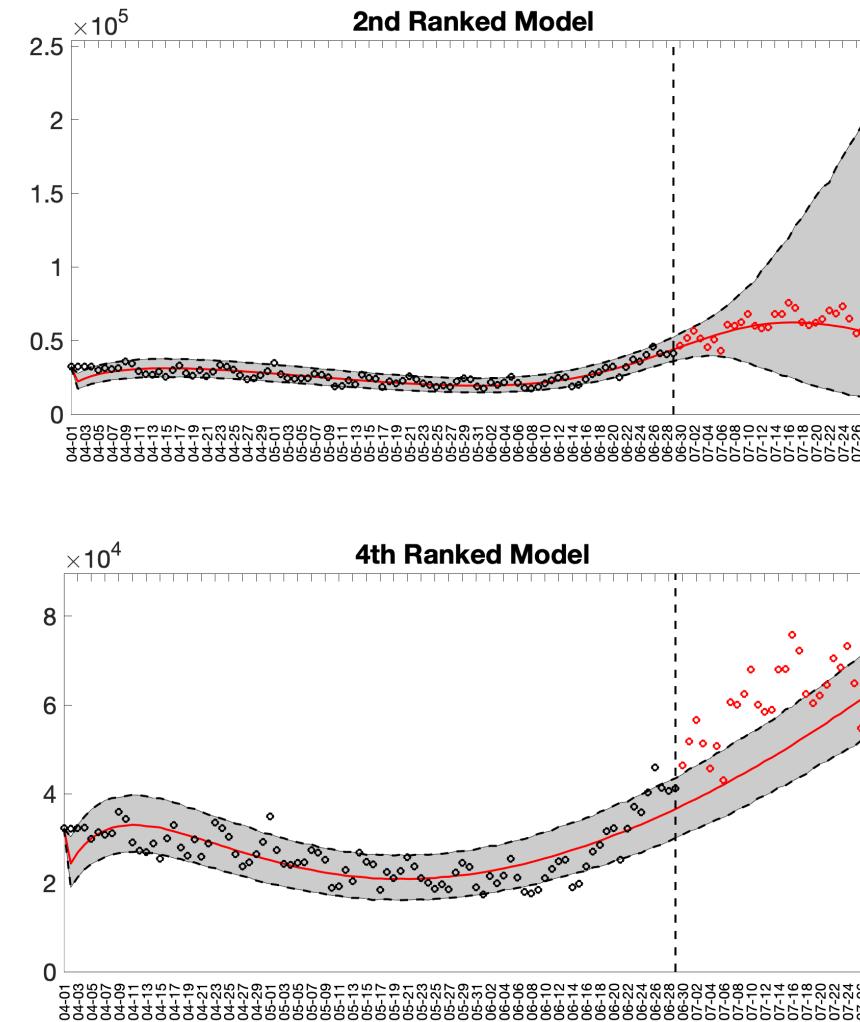
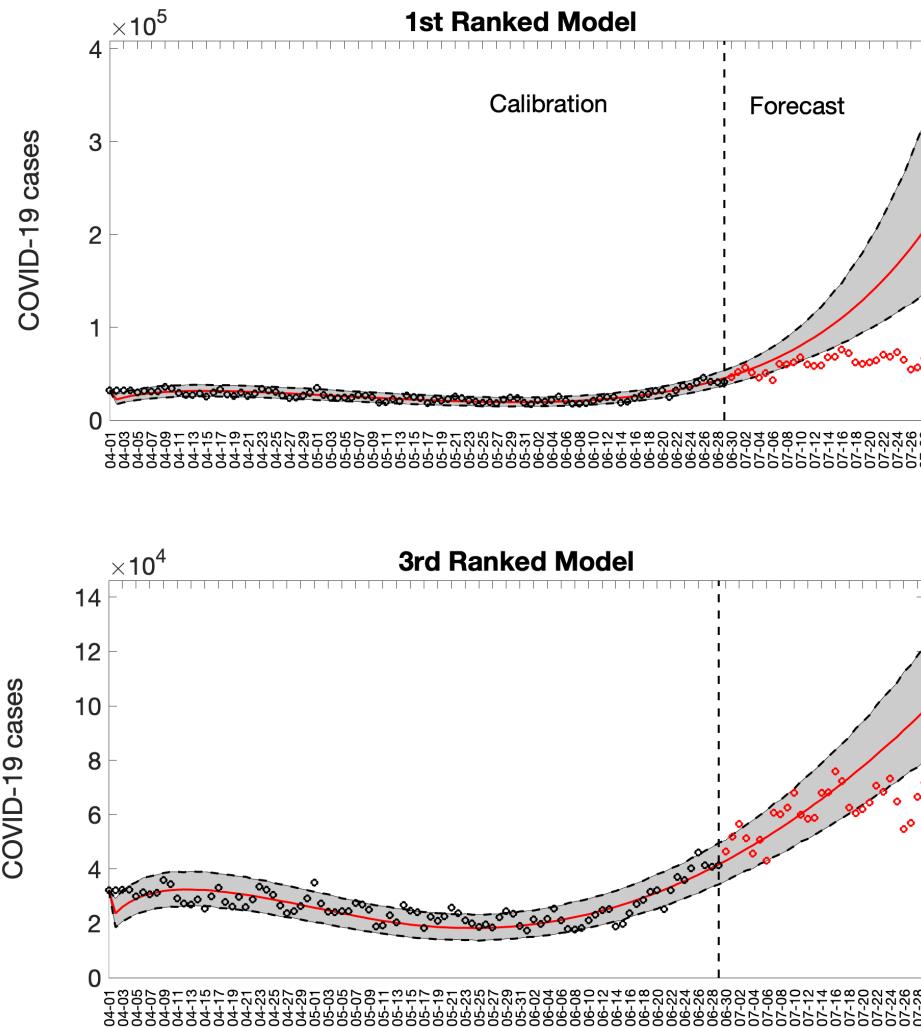
Representative fits of the top-ranking sub-epidemic models



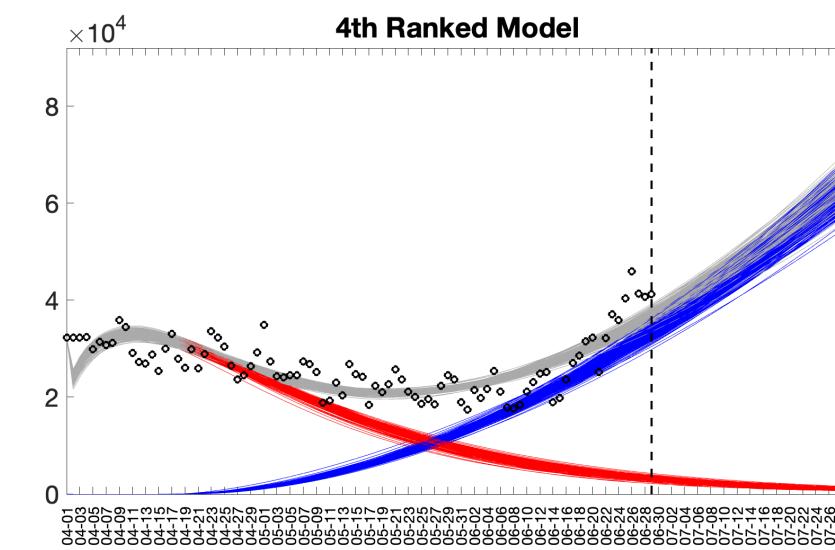
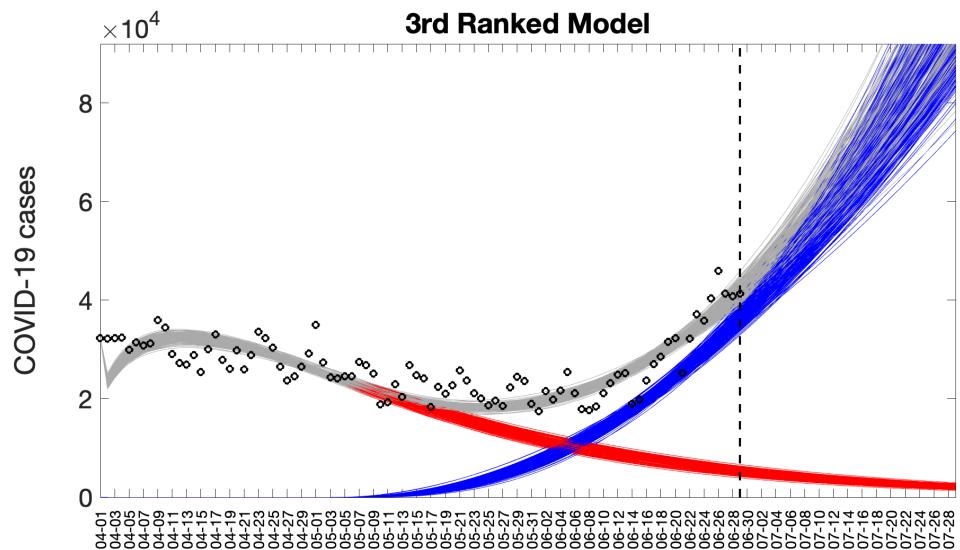
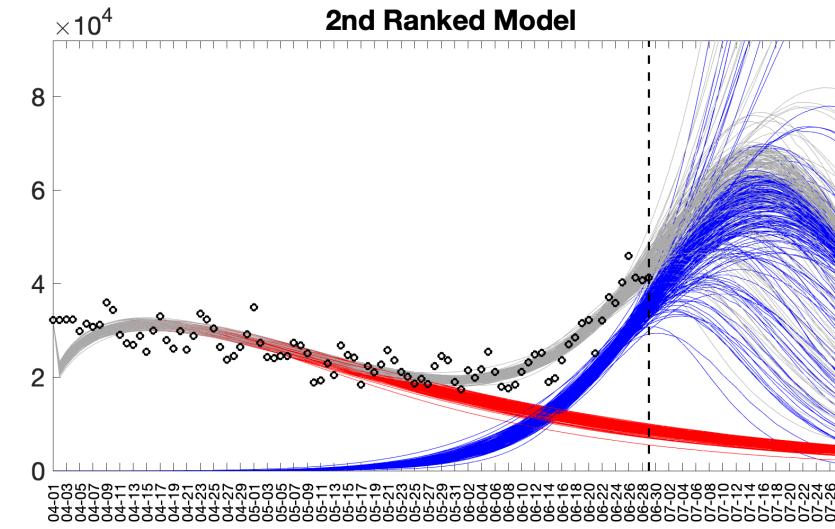
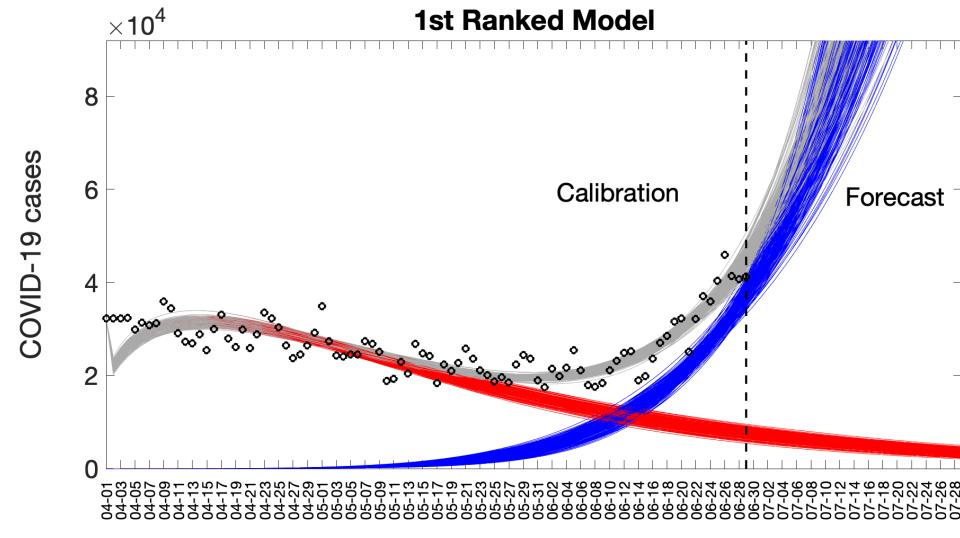
Parameter estimates for the top-ranking sub-epidemic model



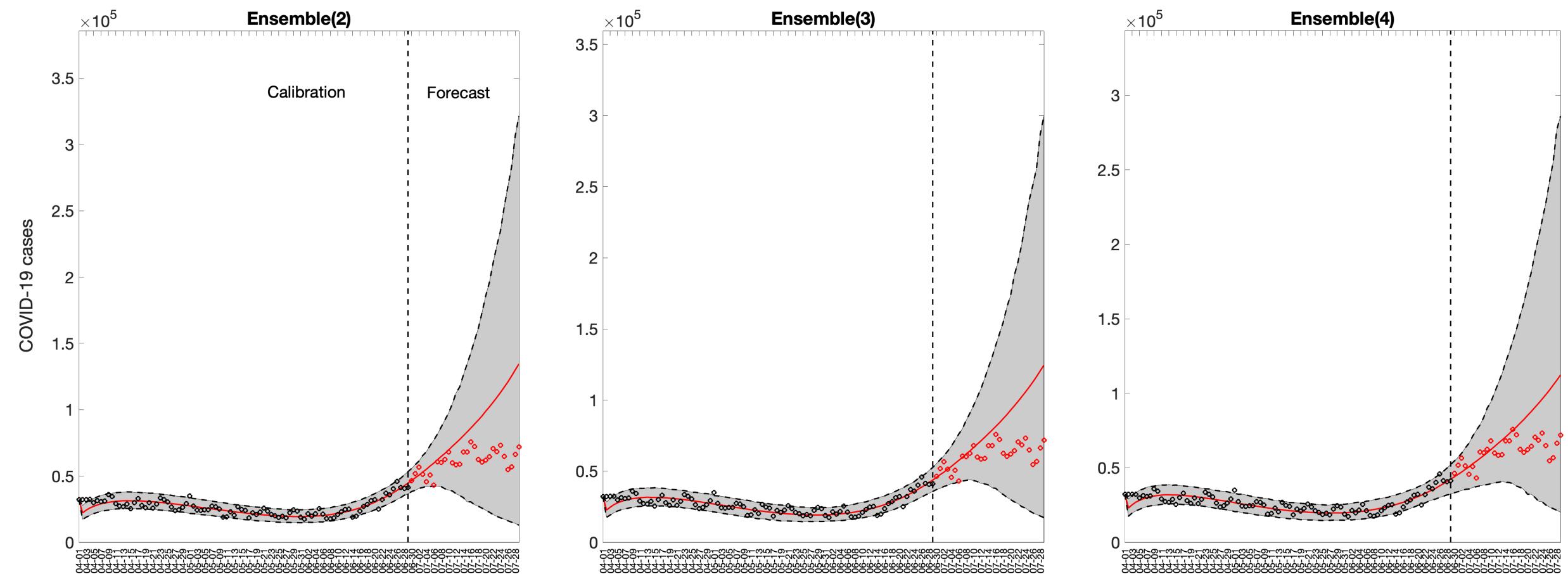
Summer resurgence, USA



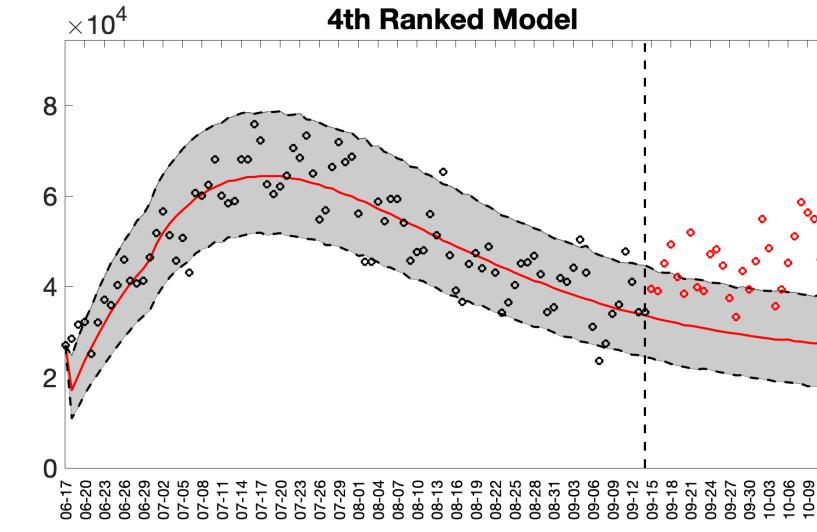
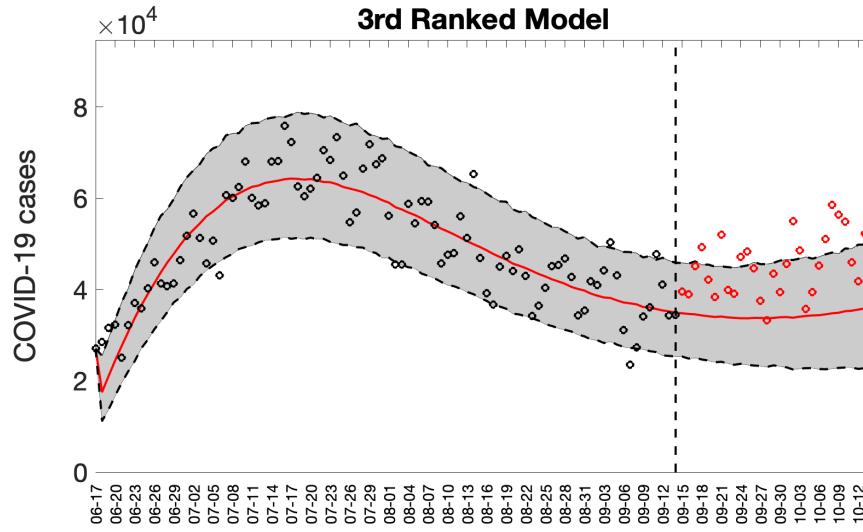
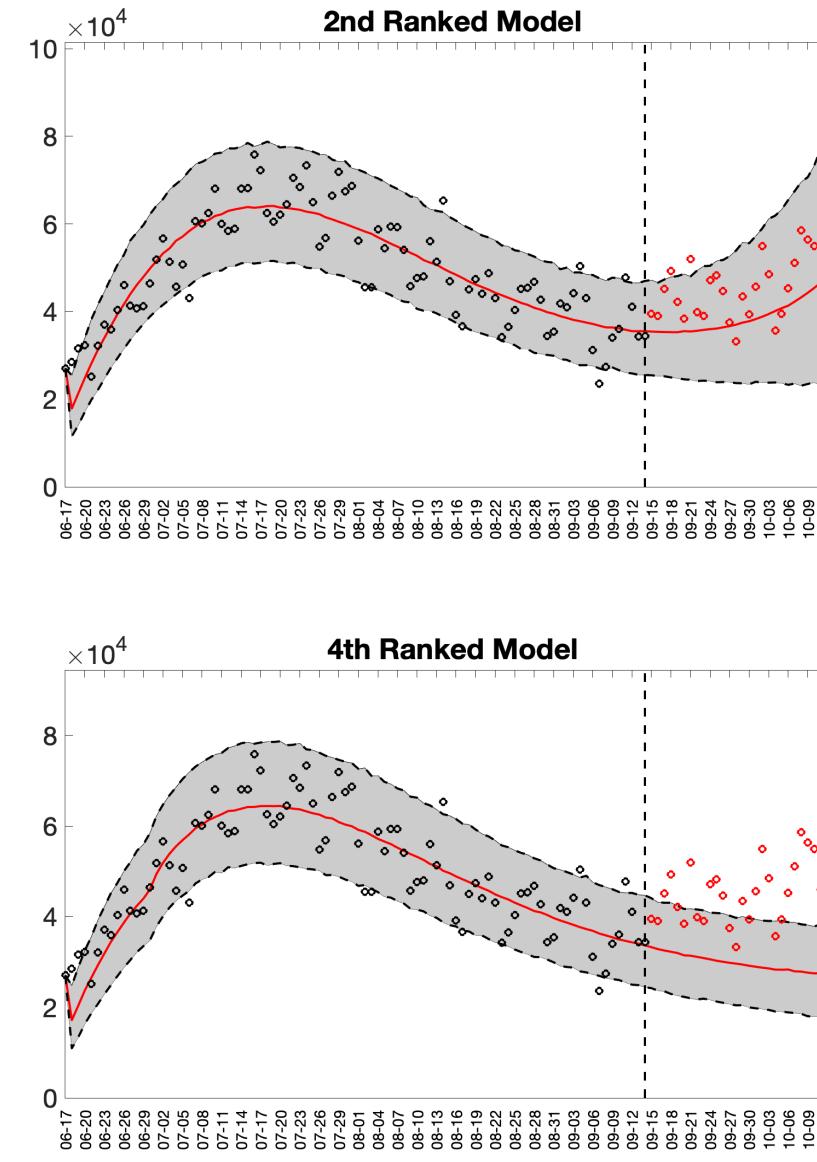
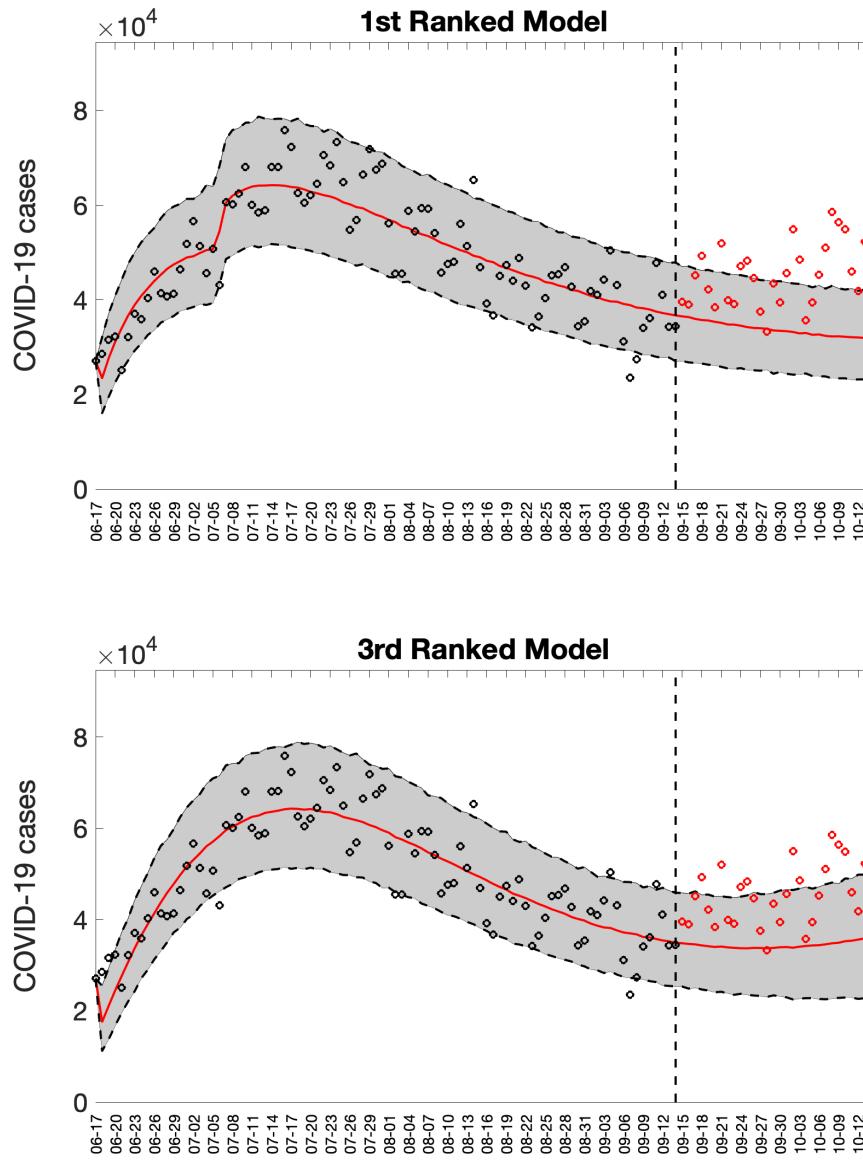
Sub-epidemic profiles



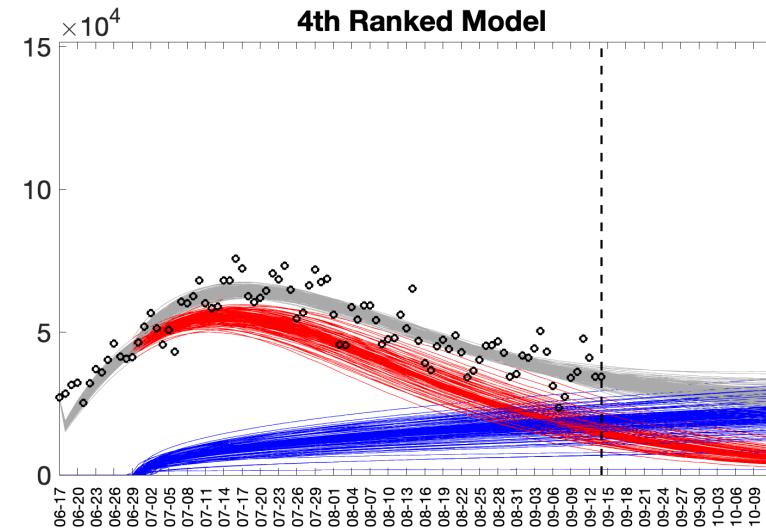
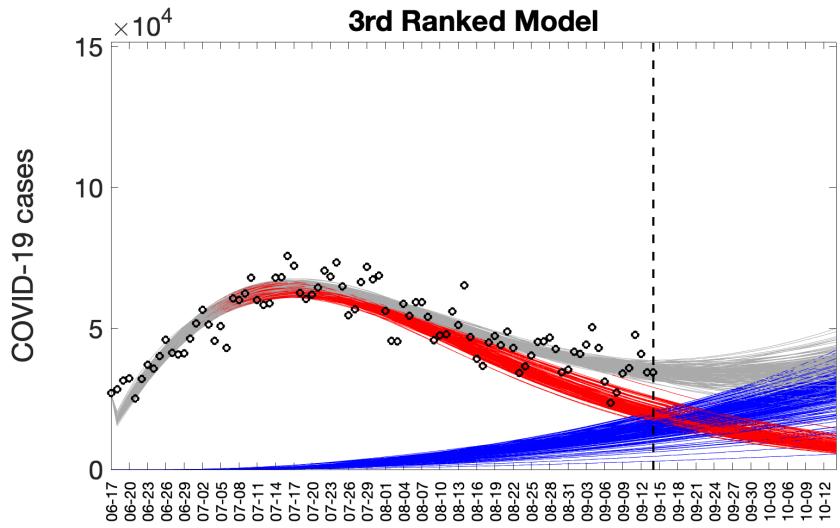
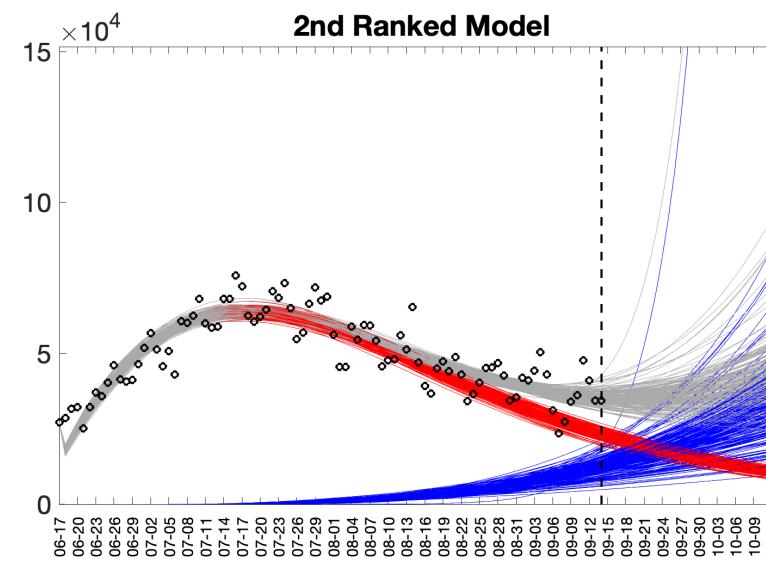
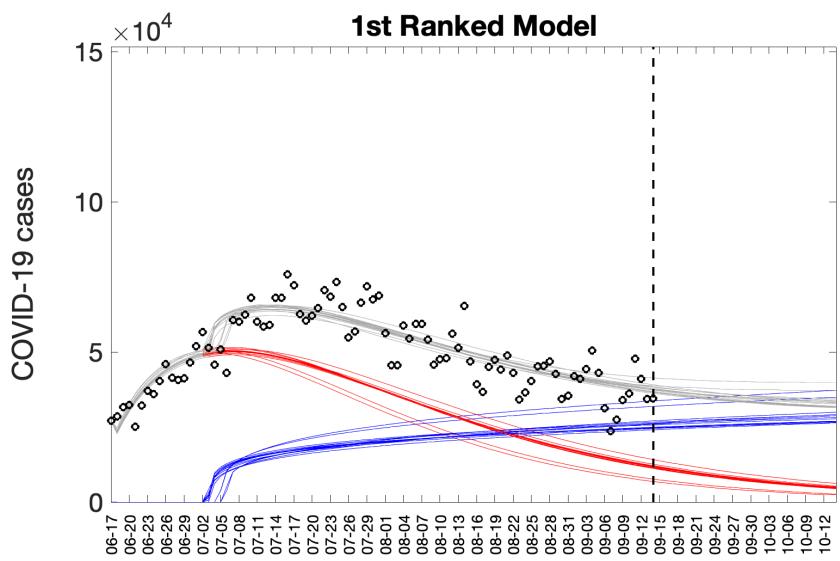
Ensemble models – Summer resurgence, USA



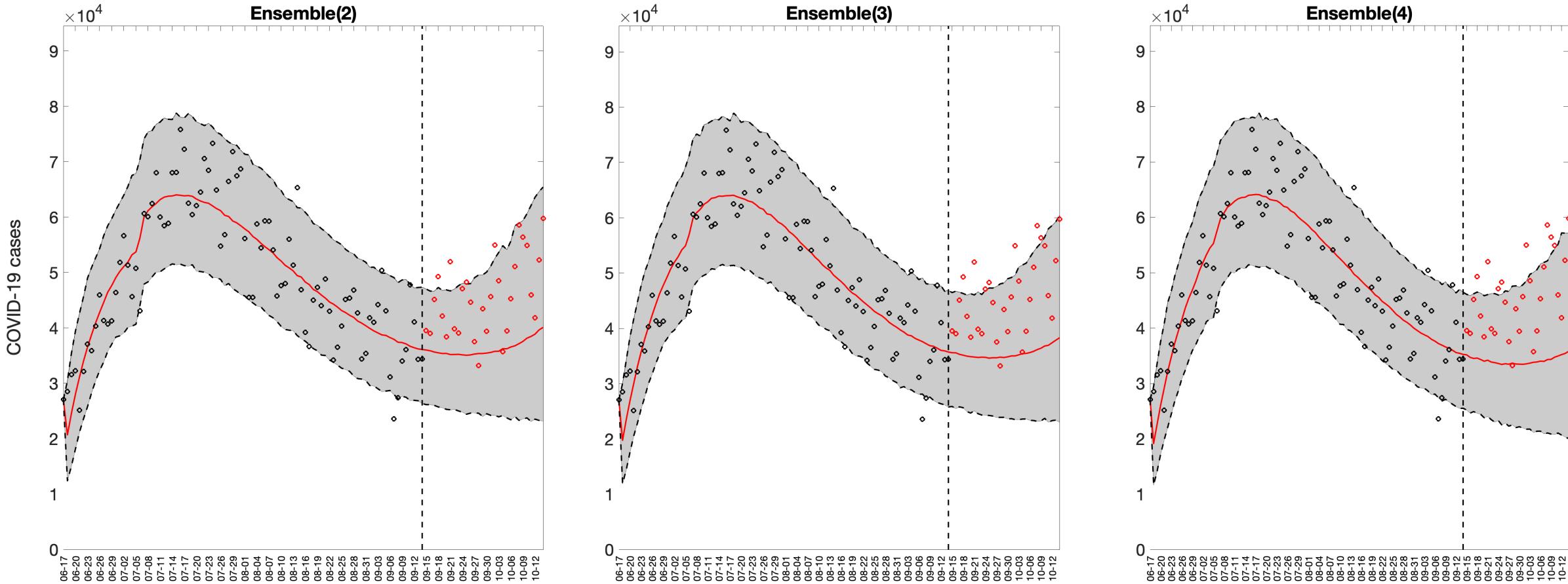
Fall resurgence, USA



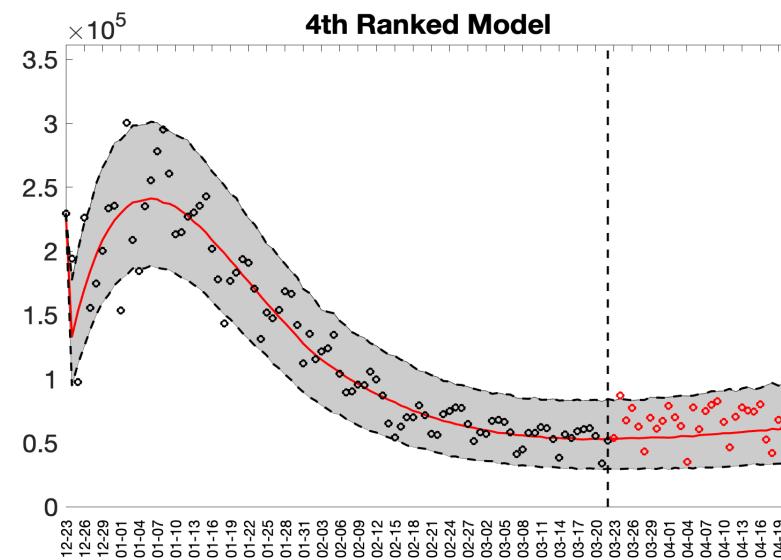
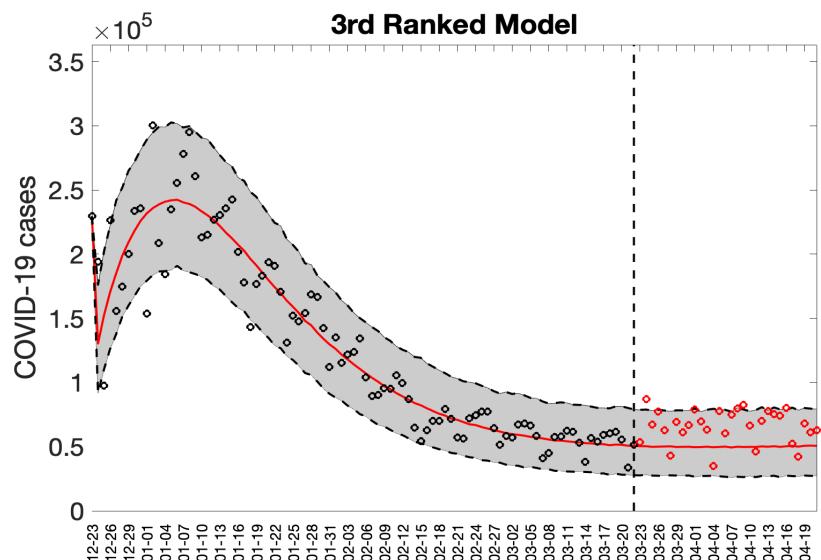
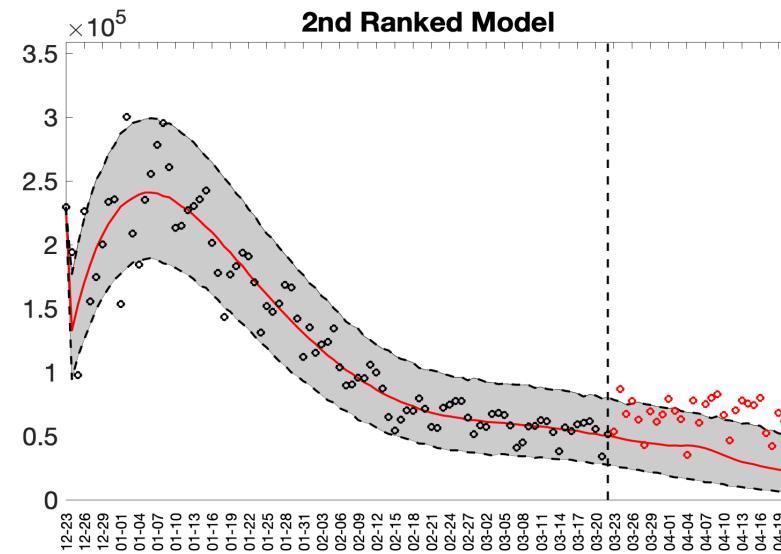
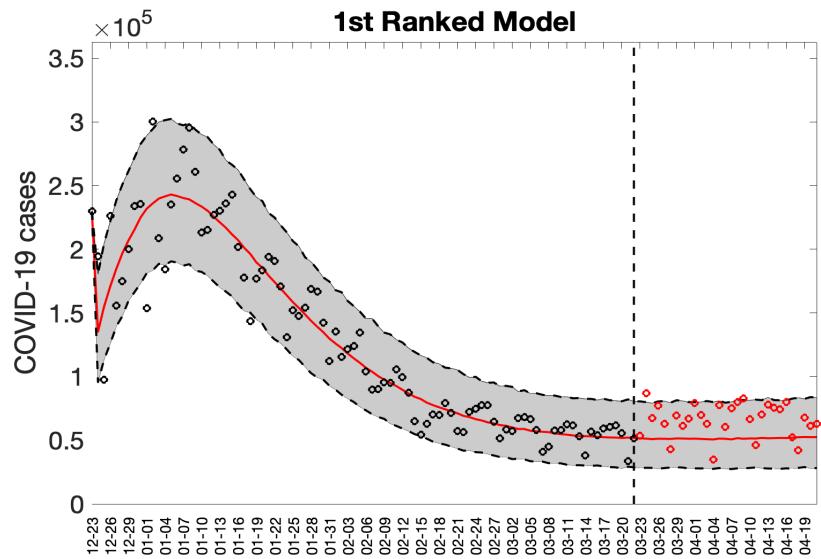
Sub-epidemic profiles



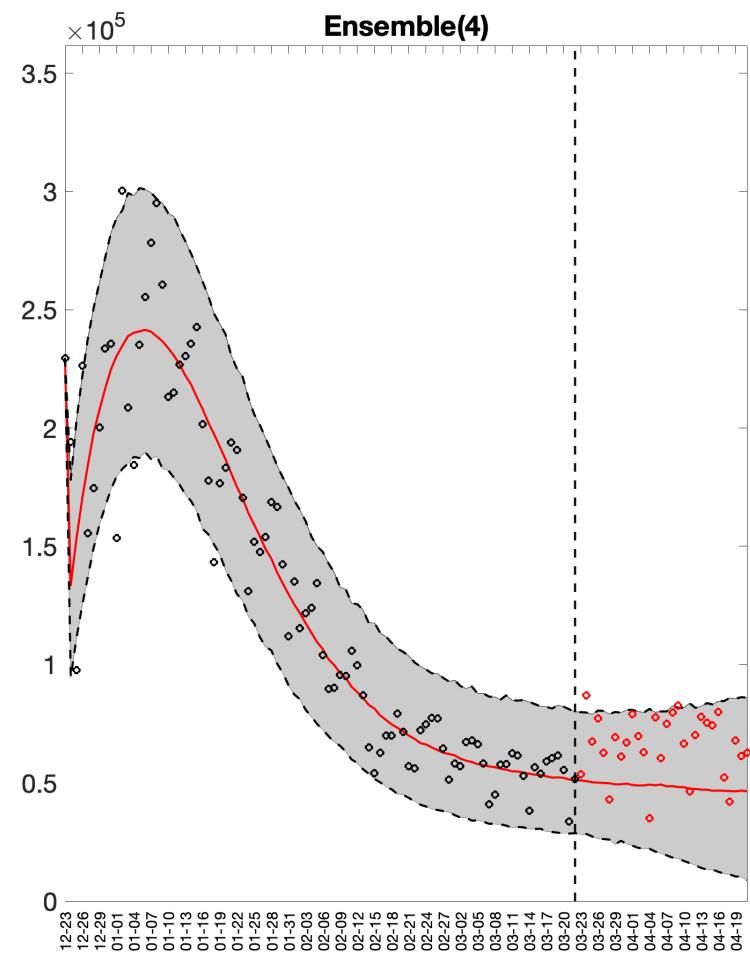
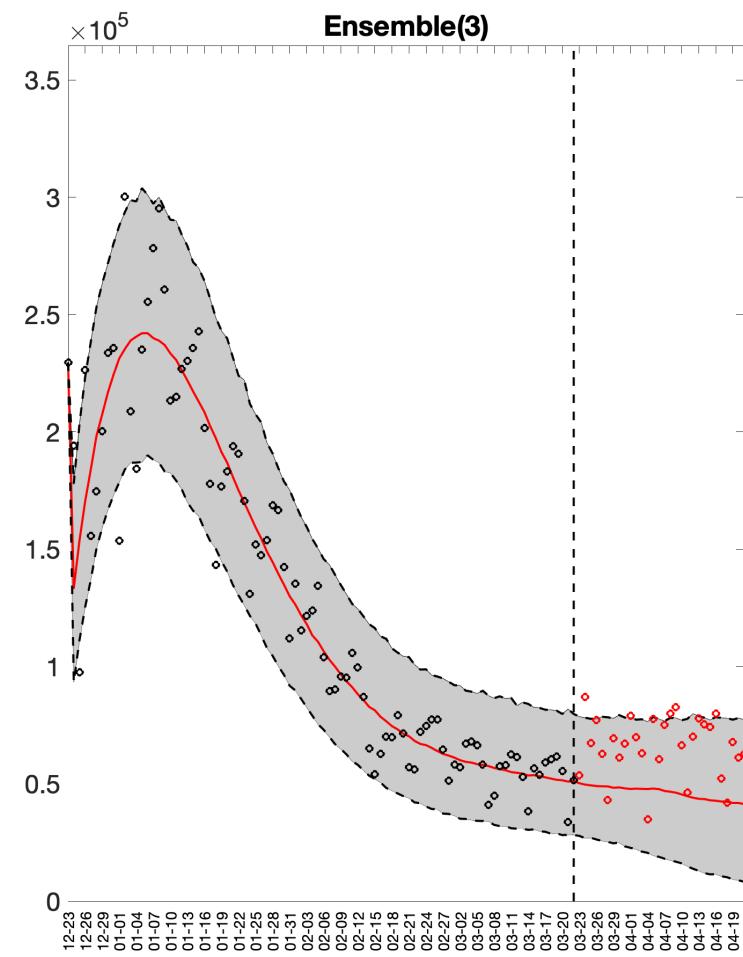
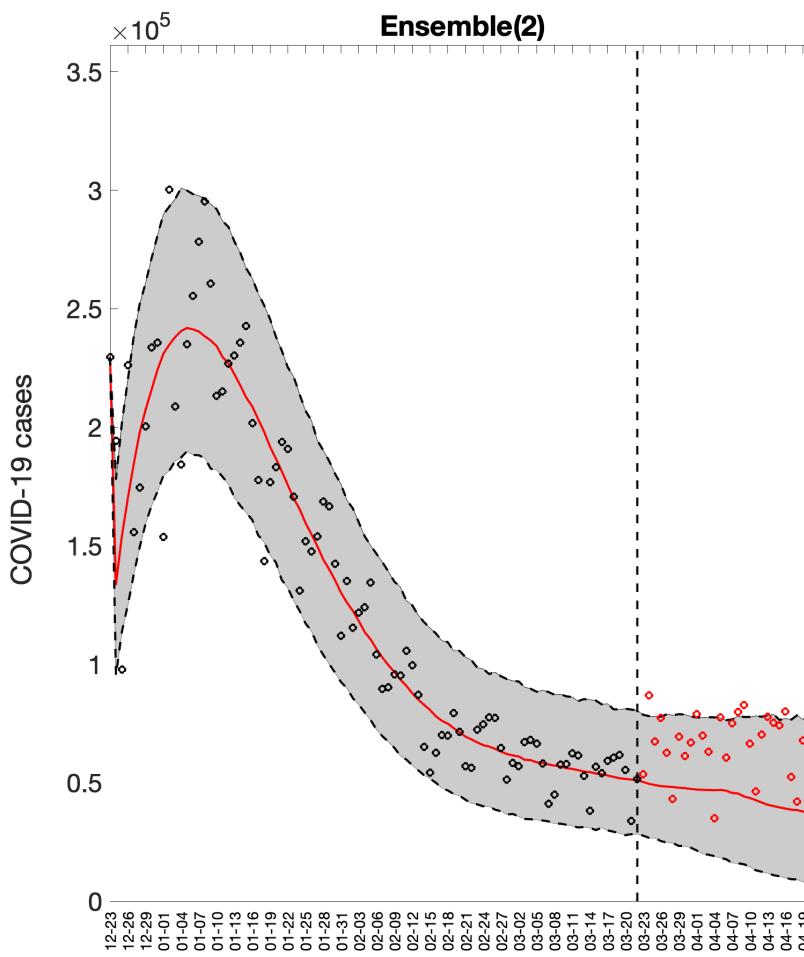
Ensemble models – Fall resurgence



Spring 2021



Ensemble models – Spring 2021



Ensemble n-subepidemic framework - MATLAB Toolbox

Screenshot of the GitHub repository page for the Ensemble n-subepidemic framework - MATLAB Toolbox.

Repository Information:

- Owner: gchowell
- Name: ensemble_n-subepidemic_framework
- Status: Public
- Code: 84 commits (d00f40c last week)
- Issues: 1
- Pull requests: 0
- Actions: 0
- Projects: 0
- Wiki: 0
- Security: 0
- Insights: 0
- Settings: 0

Code Overview:

- main branch (1 commit)
- 1 branch
- 0 tags

Commits:

Author	Commit Message	Date
gchowell	Add files via upload	d00f40c last week
	ensemble n-subepidemic code v1.0 Add files via upload	last week
	LICENSE Initial commit	2 months ago
	README.md Update README.md	last week

README.md Content:

ensemble_n-subepidemic_framework

A Matlab toolbox for fitting and forecasting epidemic trajectories using the ensemble n-subepidemic framework

It carries out the following tasks:

- fitting models to time series data,
- estimation of parameters with quantified uncertainty,
- plotting the model fits, sub-epidemic profiles, and residuals of the top-ranked models,
- plotting the empirical distributions of the model parameters associated with each sub-epidemic
- plotting the calibration performance metrics of the top-ranked models,
- plotting the AICc values, relative likelihood, and evidence ratio of the top-ranked models,
- plotting forecasts derived from the top-ranked and ensemble models,
- plotting the forecasting performance metrics of the top-ranked models and the ensemble models,
- plotting the effective reproduction number derived from the top-ranked models.

About:

A Matlab toolbox for fitting and forecasting epidemic trajectories using the ensemble n-subepidemic framework

Readme:

GPL-3.0 license:

Stars: 0

Watching: 1

Forks: 0

Releases:

No releases published

[Create a new release](#)

Packages:

No packages published

[Publish your first package](#)

Languages:

MATLAB 99.1% Objective-C 0.9%

https://github.com/gchowell/ensemble_n-subepidemic_framework

ARIMA models

>> Commonly used to forecast trends in finance and the weather.

The auto.arima function in the R package “forecast” is used to select orders and build the model. First, the degree of differencing $0 \leq d \leq 2$ is selected based on successive KPSS unit-root. Then given d , the orders p and q are selected based on the AICc for the d -times differenced data. For $d=0$ or $d=1$, a constant will be included if it improves the AICc value; for $d>1$, the constant μ is fixed as 0 to avoid the model having a quadratic or higher order trend.

1) (log)ARIMA using log-transformed data. Then we take the exponential of the forecasted values and the PI bounds to predict the incident death counts and get the PIs.

2) ARIMA. Any negative values are set as zero. Then, it is possible that the actual coverage probability of such PIs can be smaller than the nominal value (95%).

Forecasting strategy

Data. We used daily COVID-19 deaths reported in the USA from the publicly available data tracking system of the Johns Hopkins Center for Systems Science and Engineering (CSSE) from 27 February 2020 to 30 March 2022.

Models. We conducted short-term forecasts using the top-ranking n -subepidemic model ($1 \leq n \leq 2$) and three ensemble models constructed with the top-ranking sub-epidemic models namely Ensemble(2), Ensemble(3), and Ensemble(4). For comparison, we also generated short-term forecasts using the previously described ARIMA models.

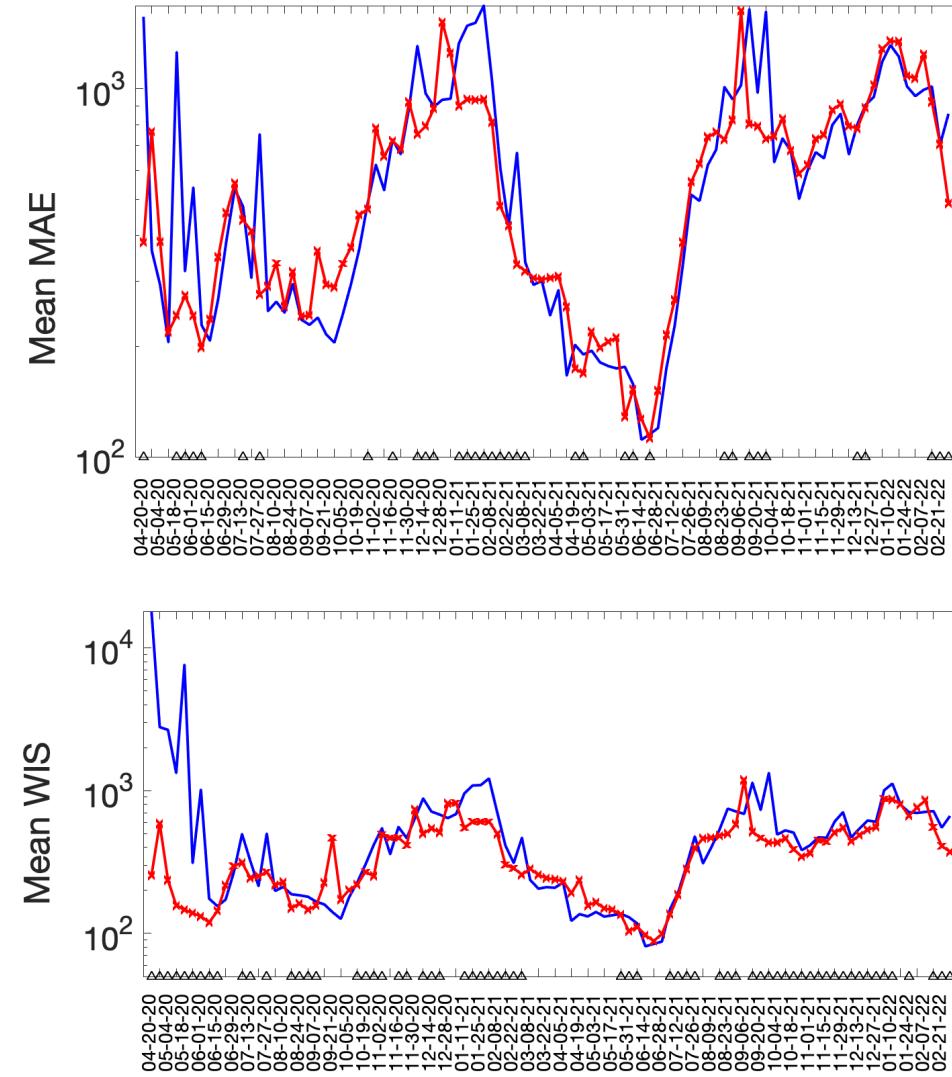
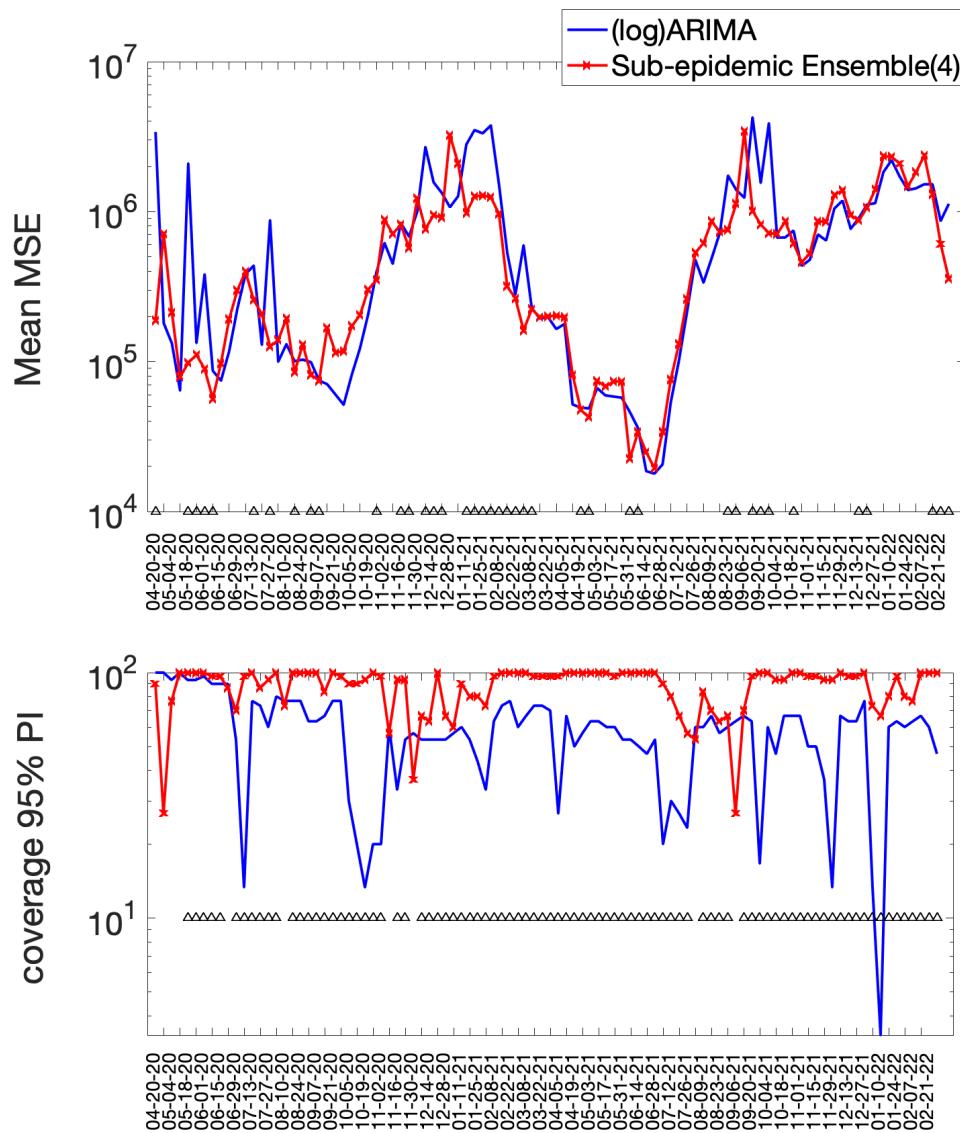
Forecasting periods. Using a 90-day calibration period for each model, we conducted a total of 98 weekly sequential 10-day, 20-day and 30-day ahead forecasts from 20 April 2020 to 28 February 2022, spanning five pandemic waves. The calibration period for each sequential forecast included seven additional days of data than the previous forecast.

Average forecasting performance metrics

20 April 2020 to 28 February 2022 (98 sequential forecasts)

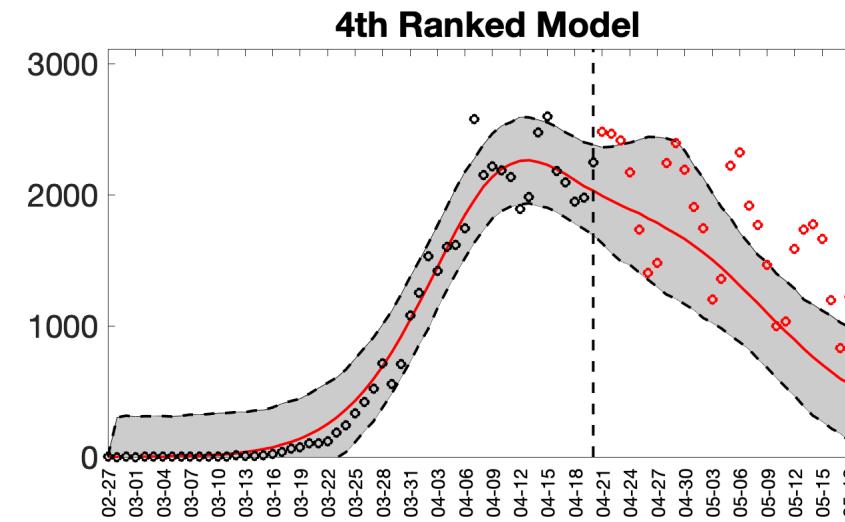
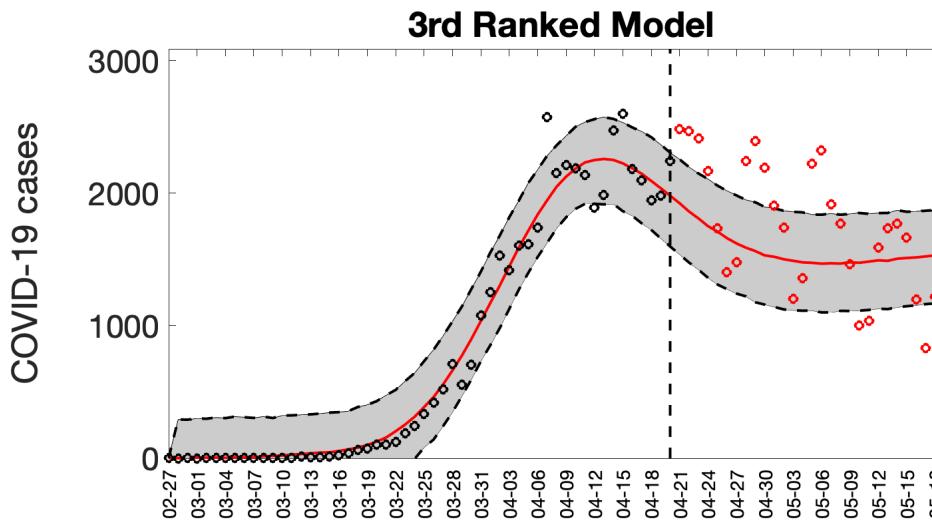
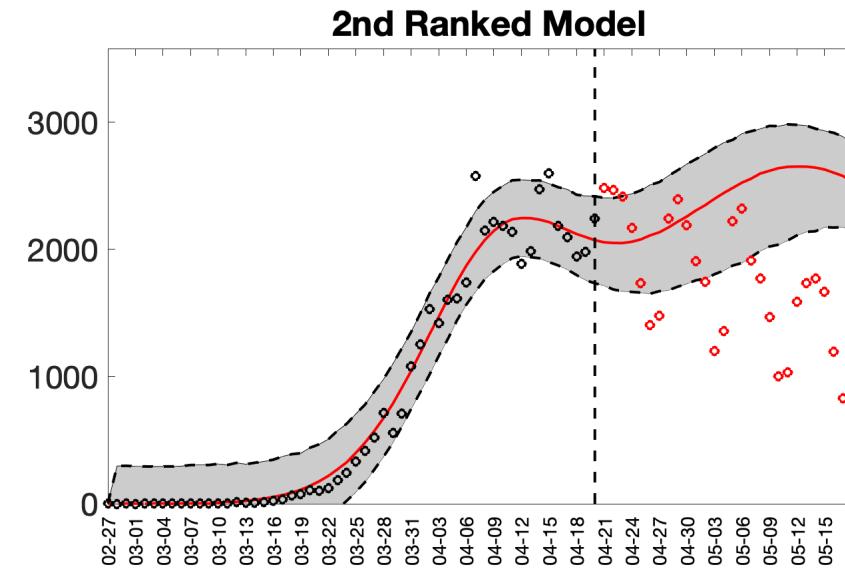
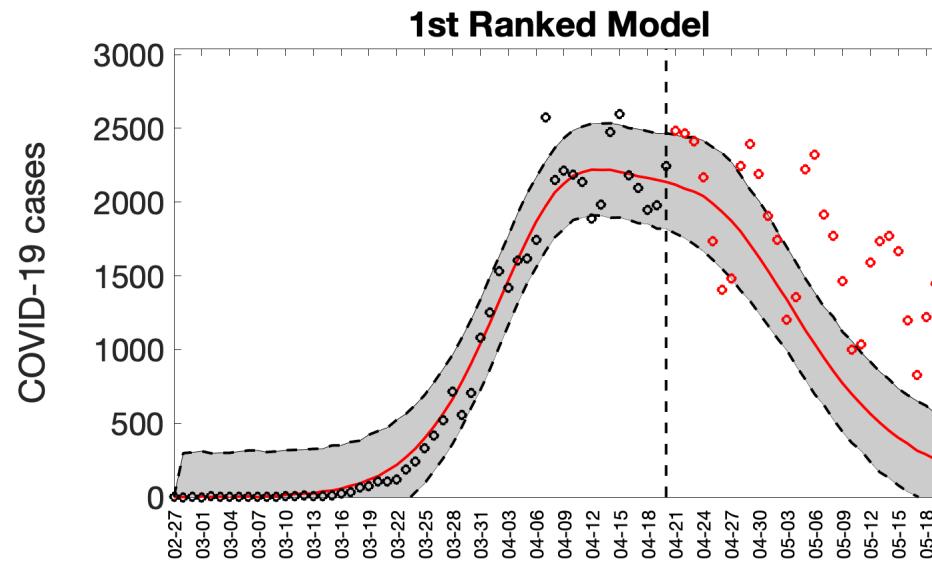
Model	Mean absolute error (MSE)	Mean squared error (MAE)	Percentage coverage of the 95% prediction interval	Weighted Interval Score (WIS)
20 days ahead				
Top ranked sub-epidemic model	646880.00	570.34	85.15	382.90
Ensemble(2) model	576700.00	544.35	88.57	354.04
Ensemble (3) model	558890.00	540.71	89.59	350.73
Ensemble (4) model	557130.00	539.30	89.44	346.83
(log) ARIMA model	591980.00	536.22	51.07	422.41
ARIMA model	538690.00	528.87	55.05	404.92
30 days ahead				
Top ranked sub-epidemic model	749560.00	613.75	82.18	421.29
Ensemble (2) model	670740.00	586.52	87.35	383.36
Ensemble (3) model	650790.00	584.20	88.20	382.79
Ensemble (4) model	644270.00	579.77	88.16	377.64
(log) ARIMA model	818530.00	621.58	57.99	767.05
ARIMA model	656480.00	591.93	60.34	439.29

Performance metrics

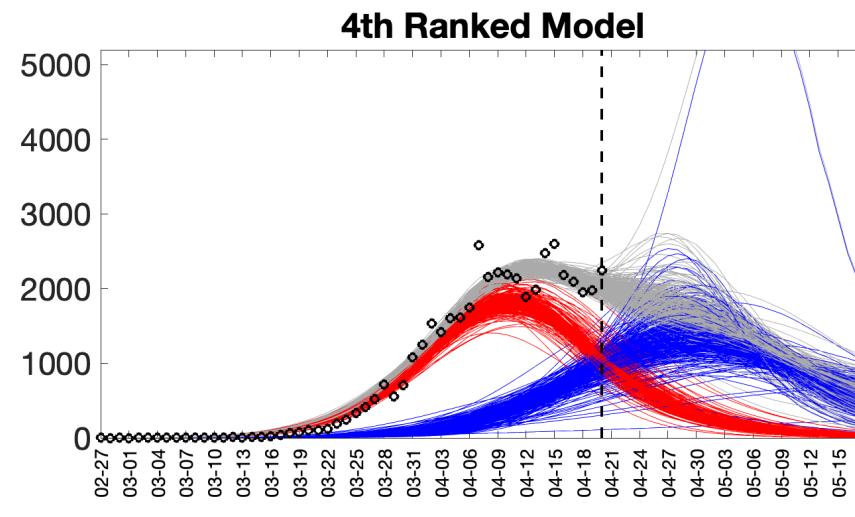
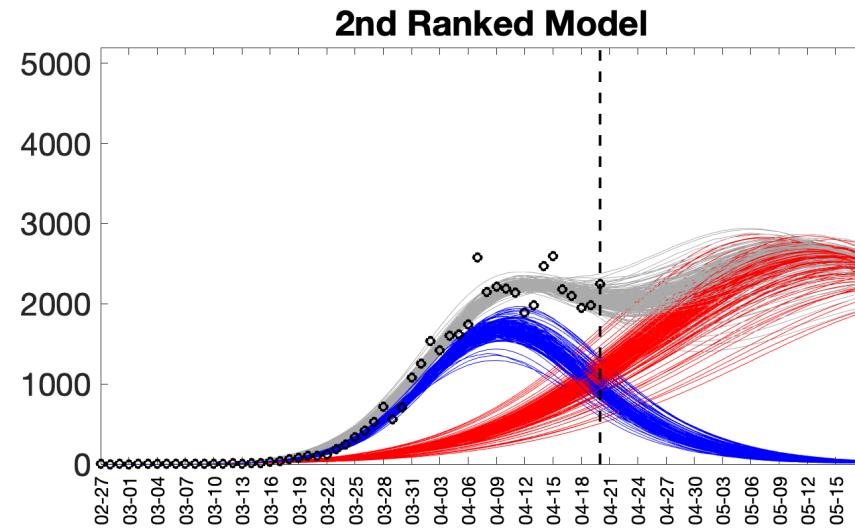
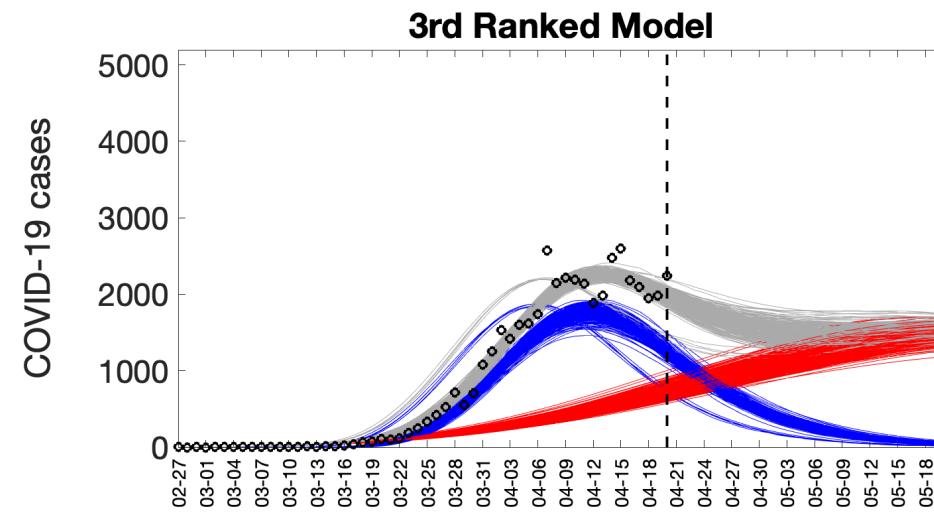
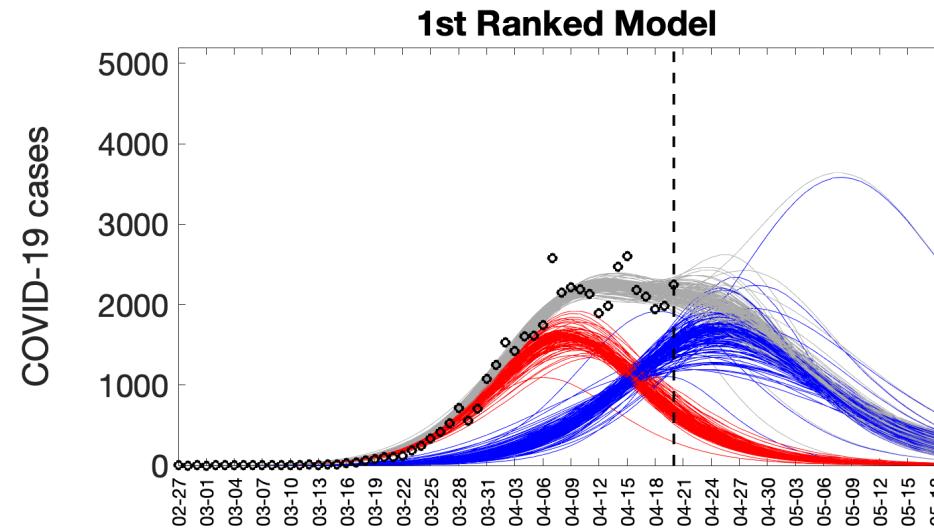


20 April 2020 to 28 February 2022

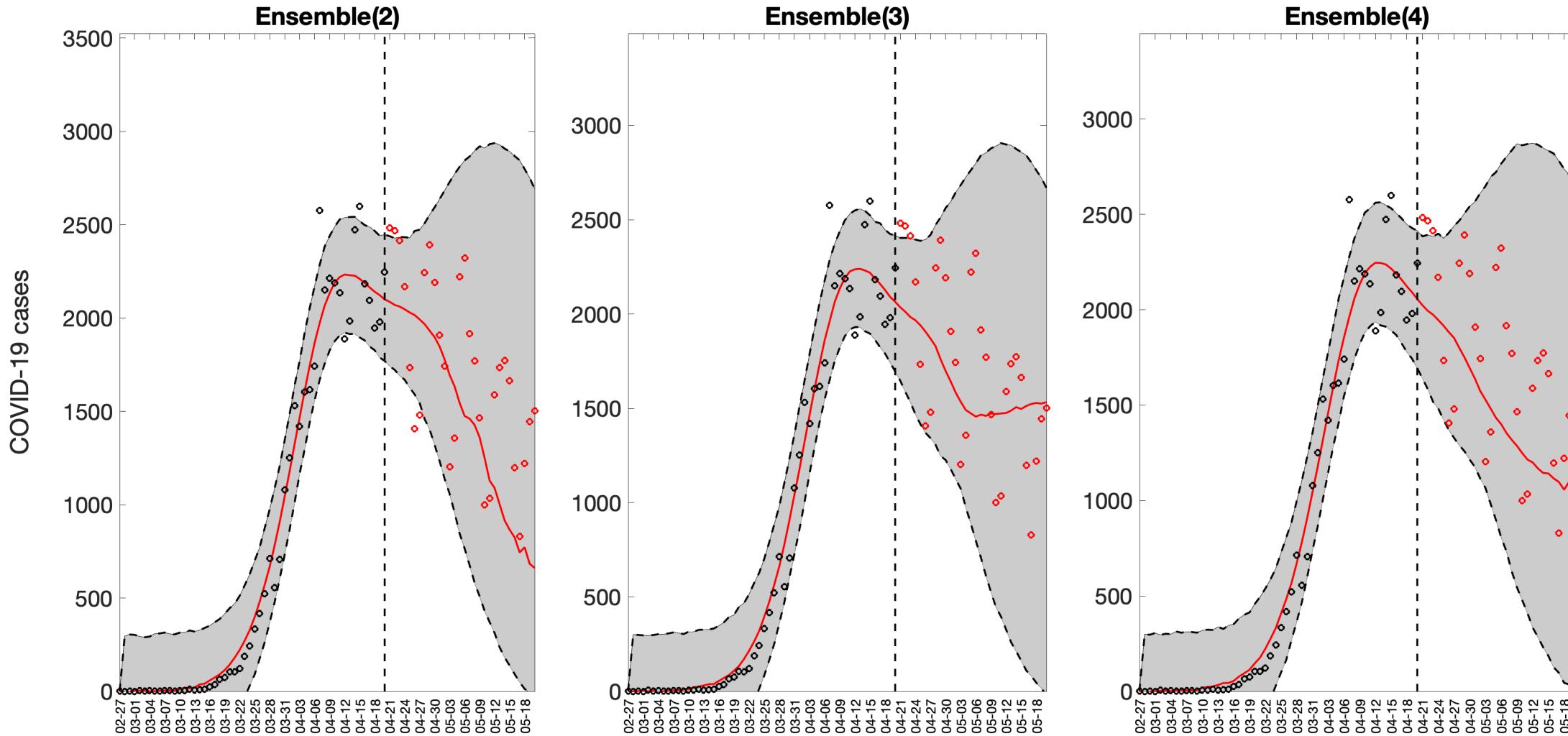
Representative top-ranking sub-epidemic forecasts



Representative top-ranking sub-epidemic profile forecasts



Representative ensemble sub-epidemic forecasts derived from top-ranking sub-epidemic models





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RESEARCH ARTICLE

An ensemble n -sub-epidemic modeling framework for short-term forecasting epidemic trajectories: Application to the COVID-19 pandemic in the USA

Gerardo Chowell Sushma Dahal, Amna Tariq, Kimberlyn Roosa, James M. Hyman, Ruiyan Luo

Version 2

Published: October 6, 2022 • <https://doi.org/10.1371/journal.pcbi.1010602>

Summary

- Our ensemble sub-epidemic models outperformed top-ranking sub-epidemic models and a set of ARIMA models in weekly short-term forecasts covering the national trajectory of the COVID-19 pandemic in the USA from the early growth phase up until the Omicron-dominated wave.
- Forecasting performance consistently improved for the ensemble sub-epidemic models that incorporated a higher number of top-ranking sub-epidemic models.
- The sub-epidemic framework could also be used to forecast other biological and social growth processes, such as the epidemics of lung injury associated with e-cigarette use or vaping and the viral spread of information through social media platforms.
- Code and performance metrics are publicly available in a GitHub repository to facilitate comparison with other modeling approaches.
- Possibilities for further development of the framework.

Real-time forecasting monkeypox, July-October 2022

- **Data.** Weekly updates of the daily confirmed monkeypox cases by date of report from the CDC and the Global.health (G.h) GitHub repository. The CDC and G.h data sources define a confirmed case of monkeypox as a person with a laboratory-confirmed case of monkeypox.
- **Scope.** At the global level and for countries that have reported the great majority of the cases including Brazil, Canada, England, France, Germany, Spain, and the United States.
- **Forecasting periods.** Data updated on Wednesday evening from both the CDC and GitHub Global.health (G.h) repository.

EPIDEMIC FORECASTING CENTER

Forecasts of national monkeypox incidence in the United States.

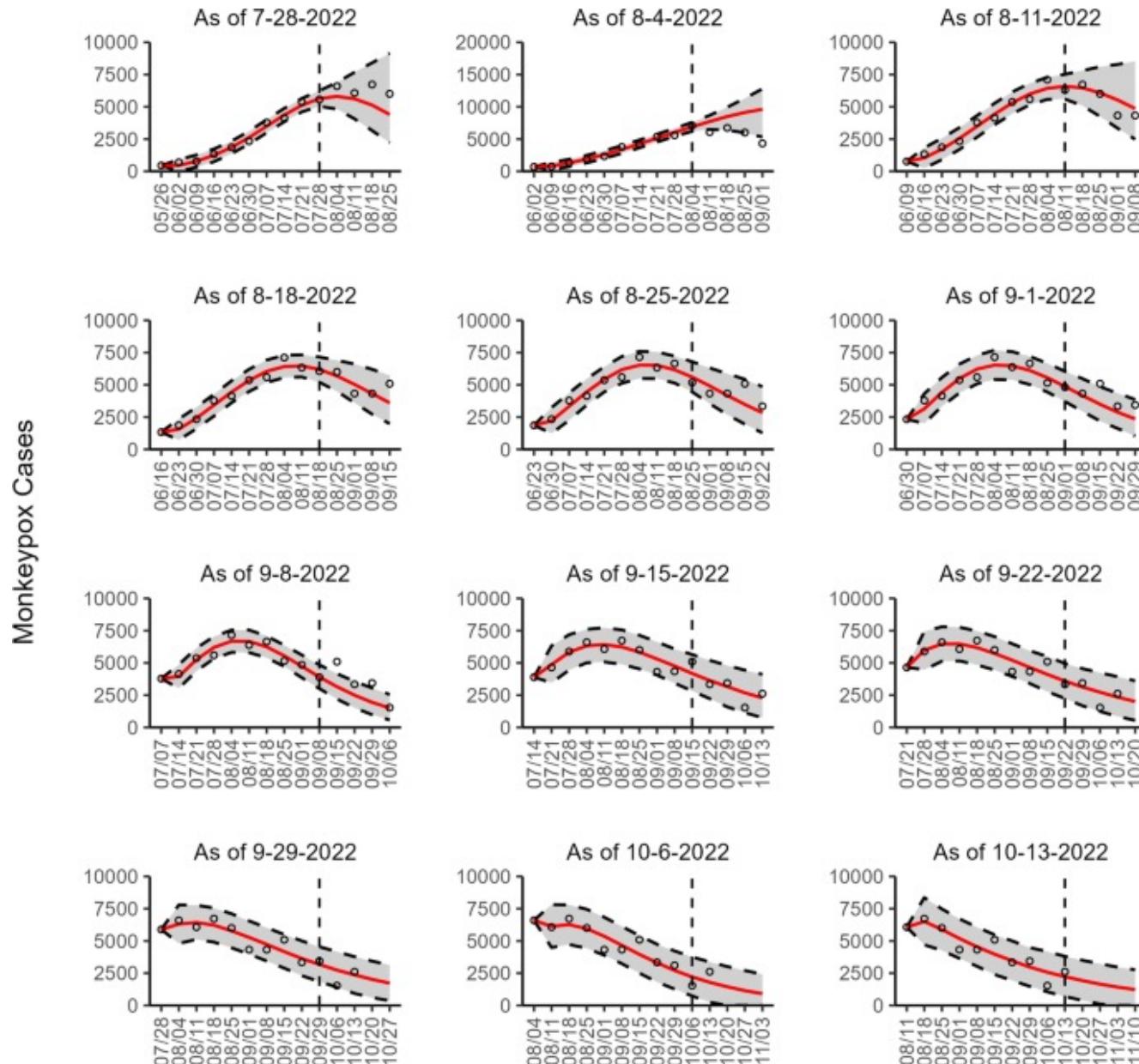
Model: [Ensemble n-sub-epidemic modeling framework](#)

Forecasts are updated every Wednesday by 9 p.m. ET. Please contact gchowell@gsu.edu for more information.

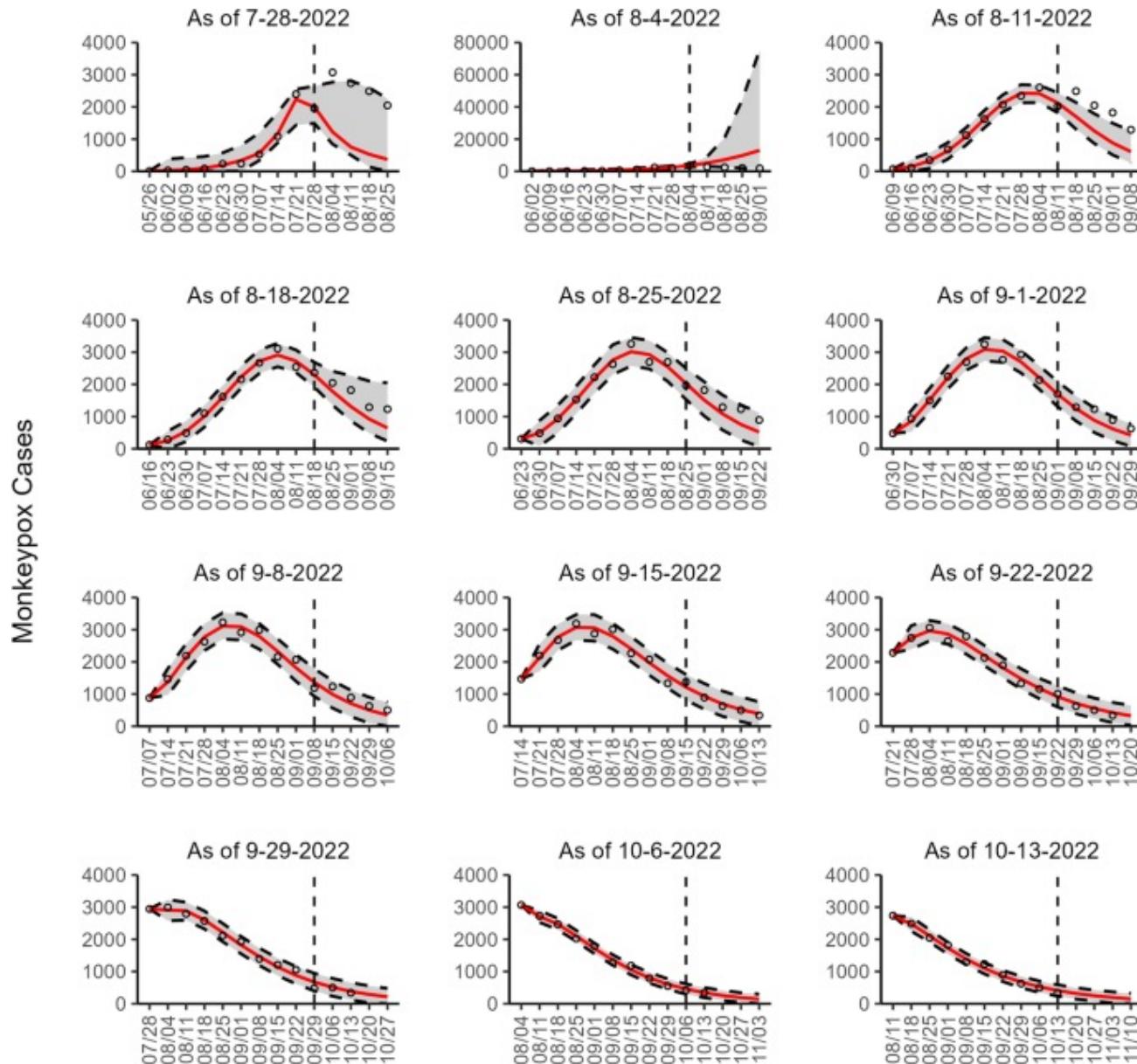
Please click on the ^ symbol to open the previous forecasts.



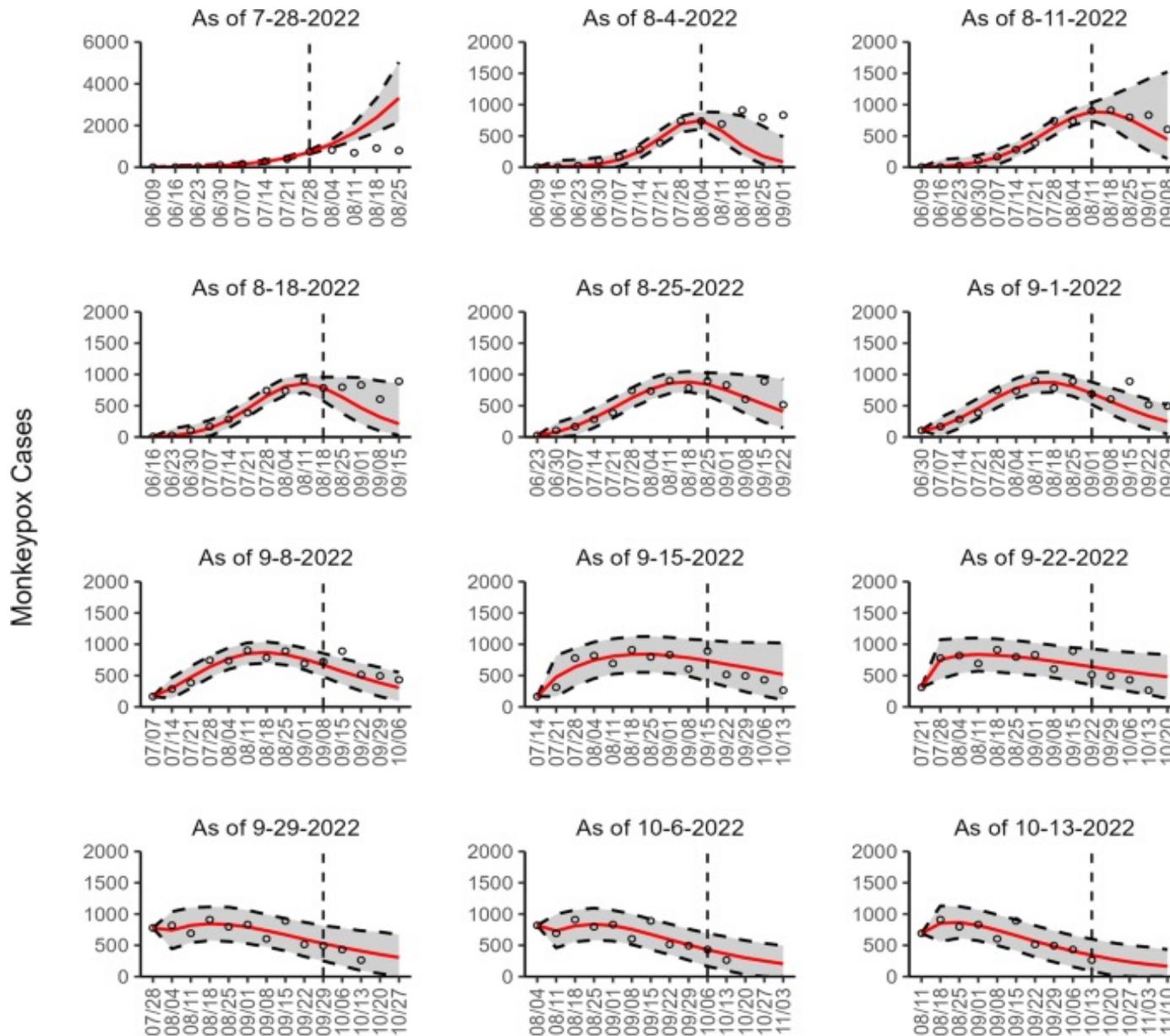
Weighted ensemble model forecasts, Global



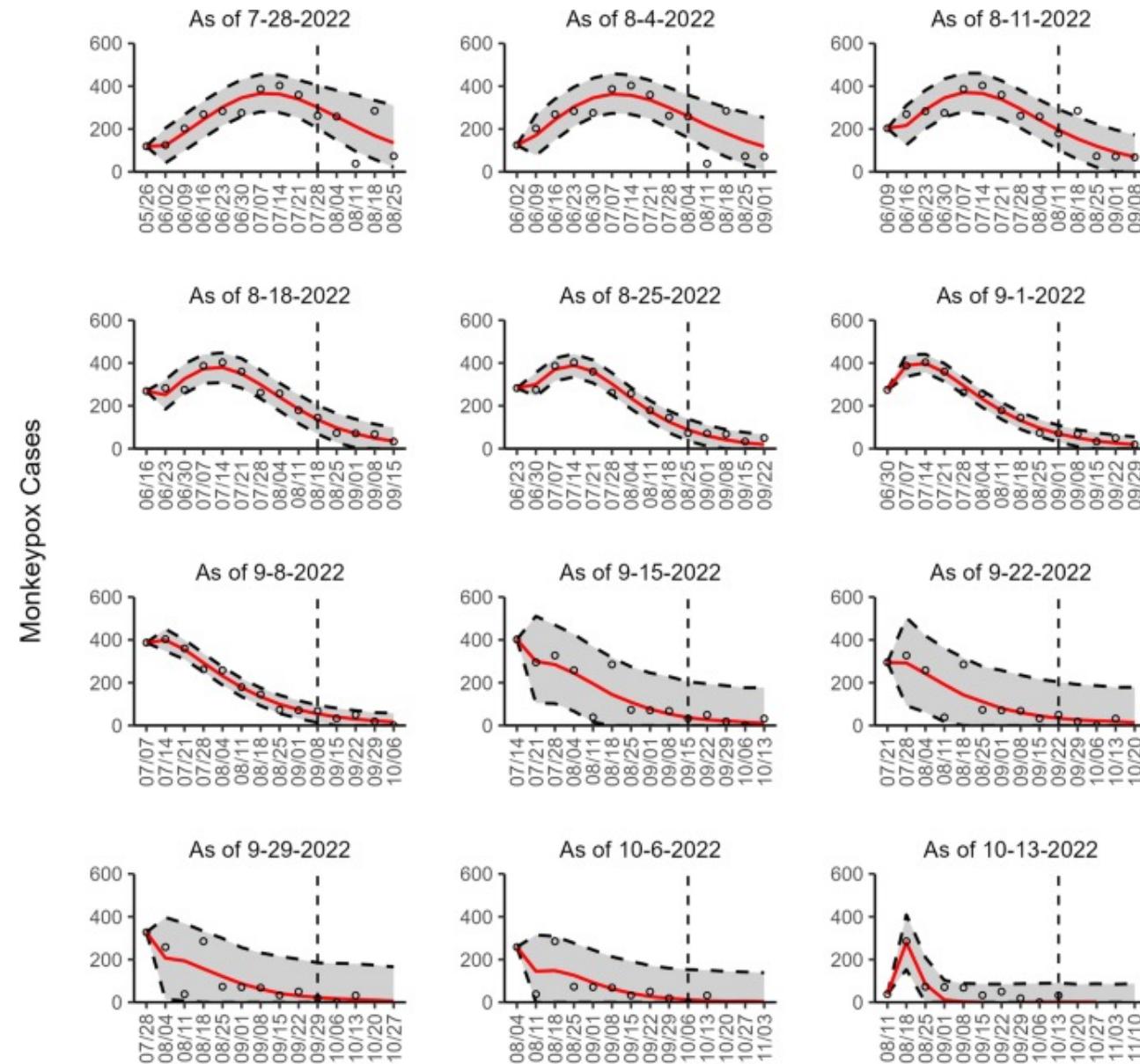
Weighted ensemble model forecasts, USA



Weighted ensemble model forecasts, Brazil



Weighted ensemble model forecasts, England





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Research article | Open Access | Published: 16 January 2023

Real-time forecasting the trajectory of monkeypox outbreaks at the national and global levels, July–October 2022

Amanda Bleichrodt, Sushma Dahal, Kevin Maloney, Lisa Casanova, Ruiyan Luo & Gerardo Chowell 

BMC Medicine 21, Article number: 19 (2023) | [Cite this article](#)

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Sections

Figures

References

Abstract

Background

Methods

Results

Collaborators



Kimberlyn Roosa
Now postdoc



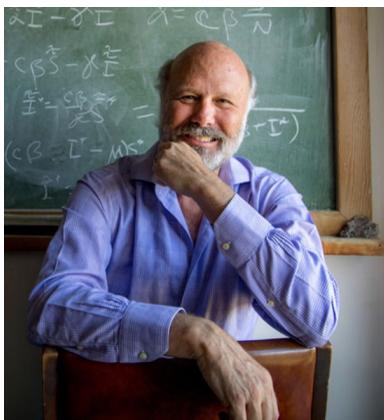
Amna Tariq
Now postdoc



Sushma Dahal
PhD student



Amanda Bleichrodt
PhD student



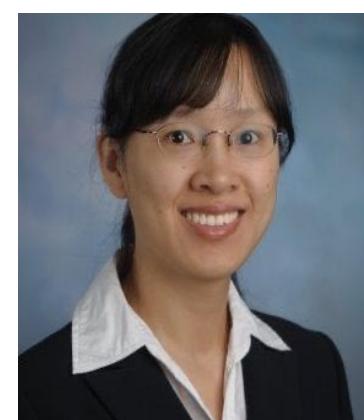
Mac Hyman (Tulane)



Kevin Maloney (GSU)



Lisa Casanova (GSU)



Ruiyan Luo (GSU)

