

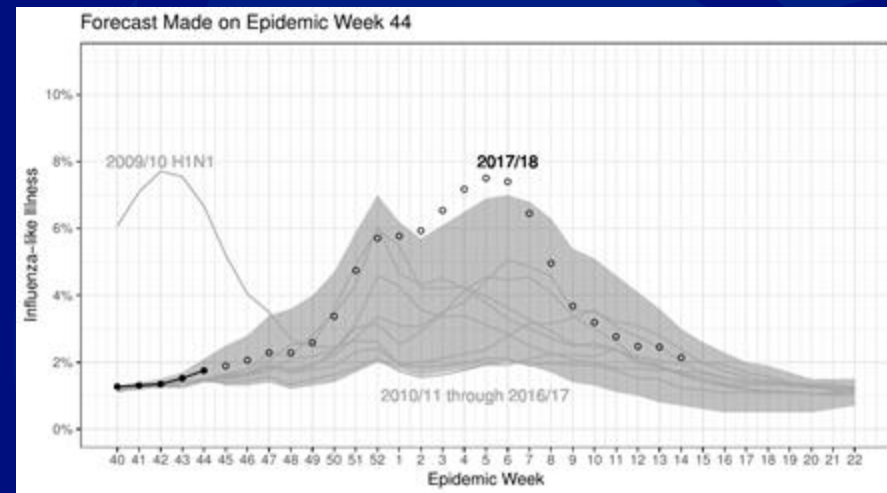
EpiFFORMA: Ensemble Weighting for Forecasting Emerging Epidemic Time Series without Historical Data

Alexander C. Murph, Lauren Beesley, Lauren Castro, Casey Gibson, Sara del Valle, Dave Osthus

January 8th, 2025

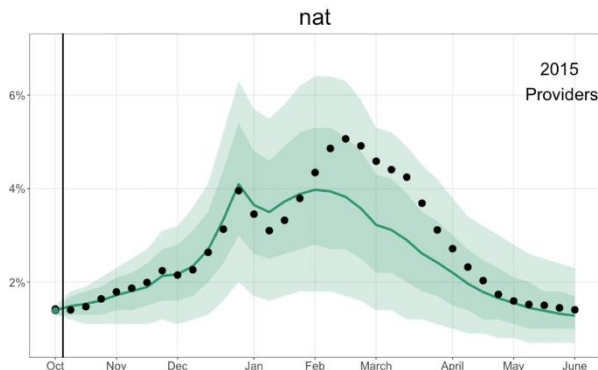
murph@lanl.gov

LA-UR-25-20054



A tale of two outbreaks: Influenza and COVID

- LANL's forecasting model **Dante** won the FluSight Challenge in 2018/19
 - Bayesian hierarchical model
 - Borrowed information across states and seasons
 - Took hours to run
- **However, when COVID hit in late 2019, Dante was all but useless to us. Why?**
 - **Dante** was tailored to forecast a seasonal disease, needing years of historical data
 - **COVID** isn't seasonal, nor were there any historical data at the onset
- LANL's COVID forecasting model had to be made from scratch. **This was stressful!**



GOAL: Develop and test a unified forecasting framework that works for:

- 1 Diseases with lots of historical data (e.g., flu)
- 2 Emerging diseases (e.g., COVID in early 2020)
- 3 Everything in between!

What properties do we want in a forecasting model?



Accurate and
well-calibrated



Fast to train,
faster to forecast



Scalable and
parallelizable



Able to ingest
large or small
amounts of data



Produce
probabilistic
forecasts (i.e.,
quantified
uncertainties)



**Able to borrow
information**
across
geographies
and/or diseases,
but **not require**
multiple
geographies/
diseases



Seamlessly
incorporate
**exogenous
information**
when available
(demographic
data, mobility
data, sequence
data, etc.), but
not require it

Model Ensembling: A Method to Improve Forecasts

$$\textit{Ensemble Forecast} = \sum_k \mathbf{w}_k \textit{Forecast}_k$$

Better Forecast

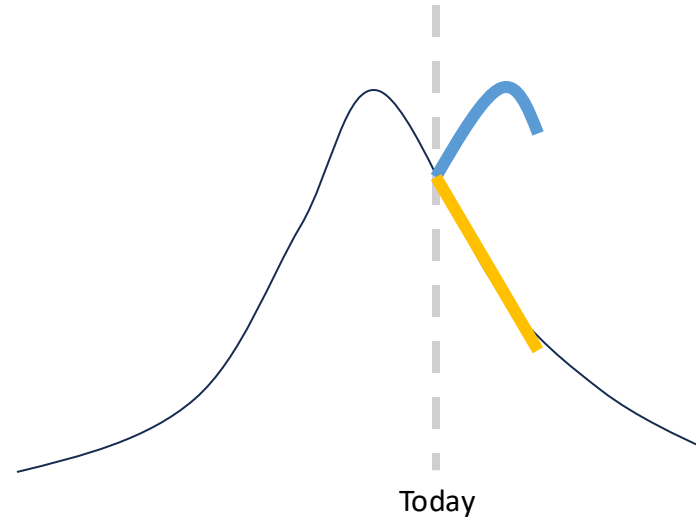
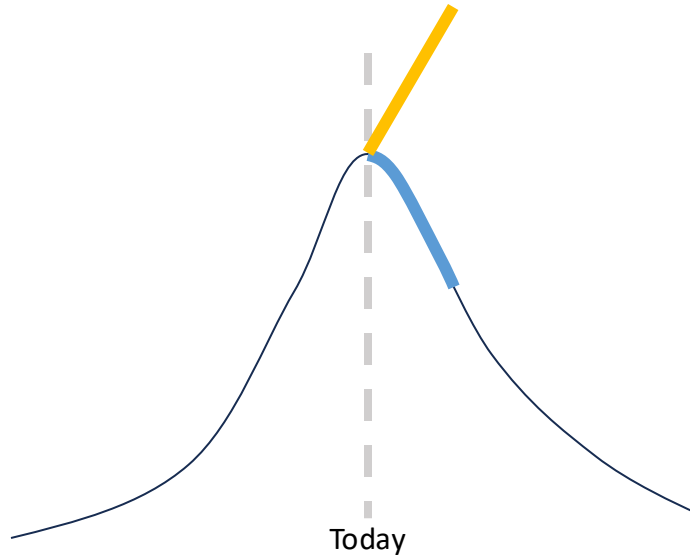
Okay forecast models

These weights are usually based on past forecast performance

What if we could identify *when* a model performs well?

Model 1: Continue linear trend

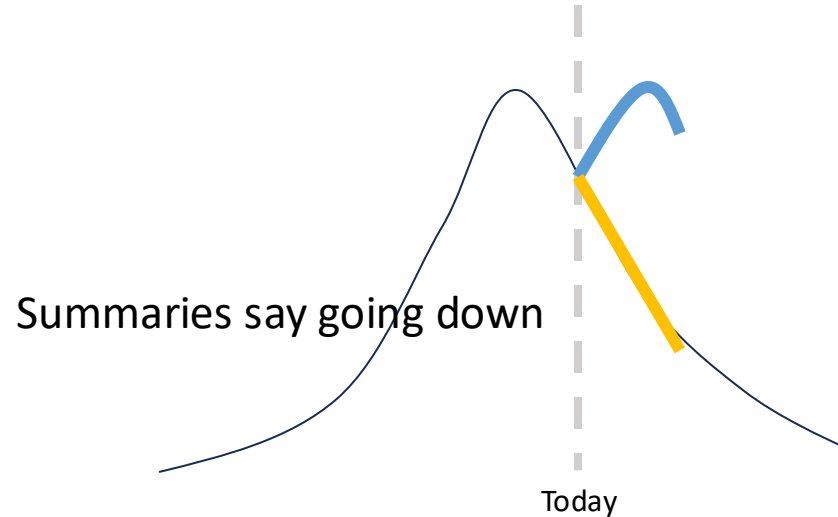
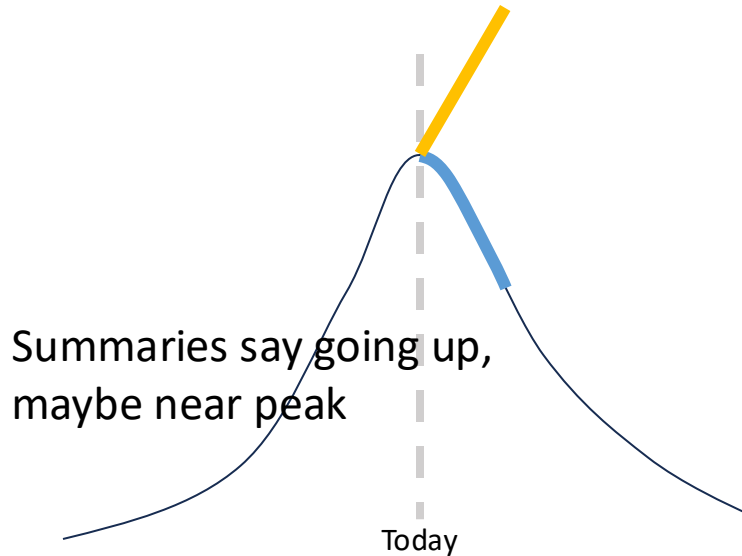
Model 2: Reverse last few observations



What if we could identify *when* a model performs well?

Model 1: Continue linear trend

Model 2: Reverse last few observations



Model Ensembling: A Method to Improve Forecasts

$$\textit{Ensemble Forecast} = \sum_k \mathbf{w}_k(\mathbf{x}) \textit{Forecast}_k$$

Problem: How do we specify the weights?

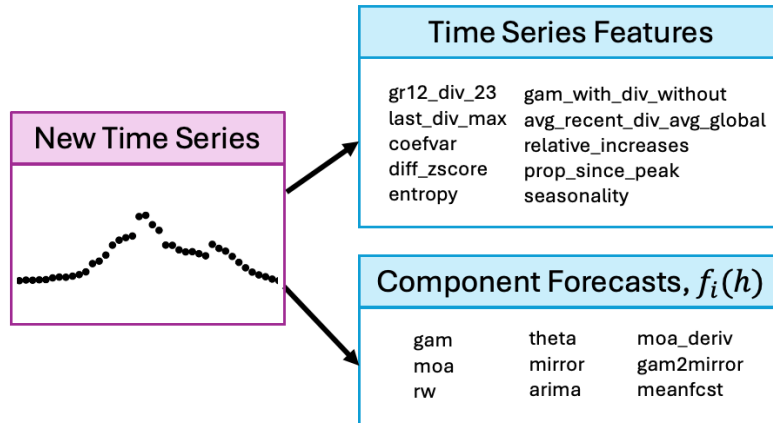
Main Idea: Automated model reweighting based on time series features, \mathbf{x}

**FFORMA: Feature-based
Forecast Model Averaging**

Pablo Montero-Manso, George Athanasopoulos,
Rob J Hyndman, Thiyaanga S Talagala

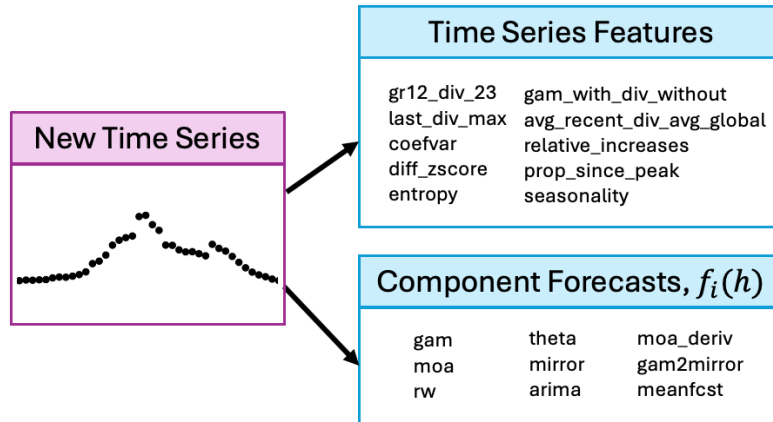
→ **epiFFORMA**

epiFFORMA: data-driven ensembling for infectious disease forecasting

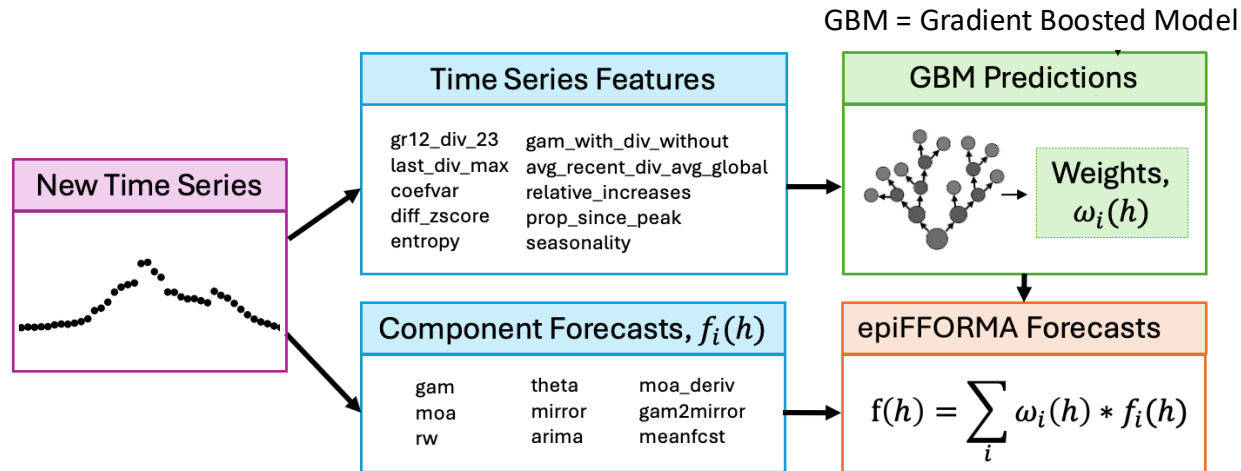


epiFFORMA: data-driven ensembling for infectious disease forecasting

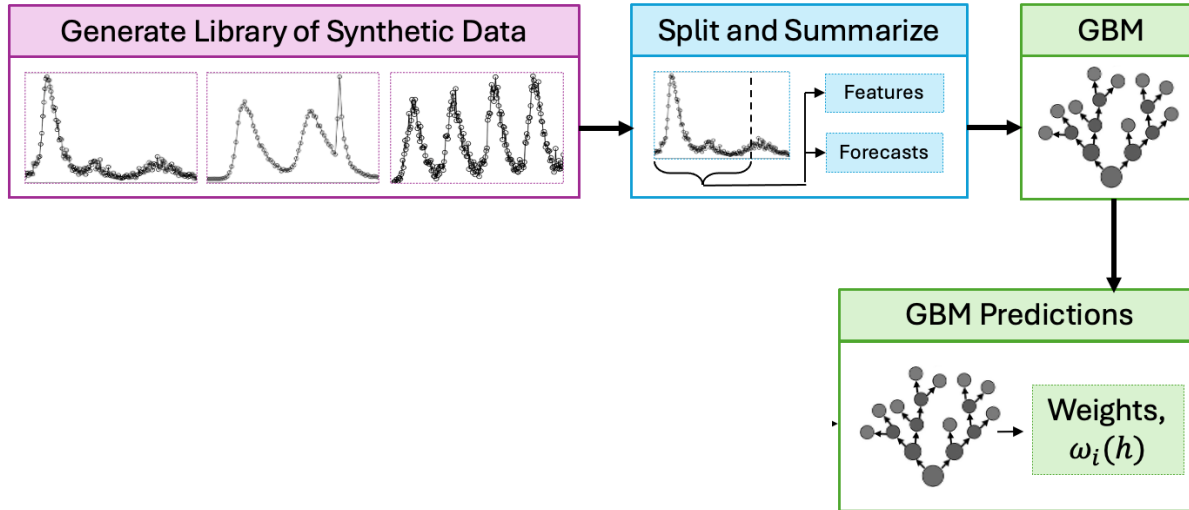
New to epiFFORMA:
Features/components tailored to infectious disease context



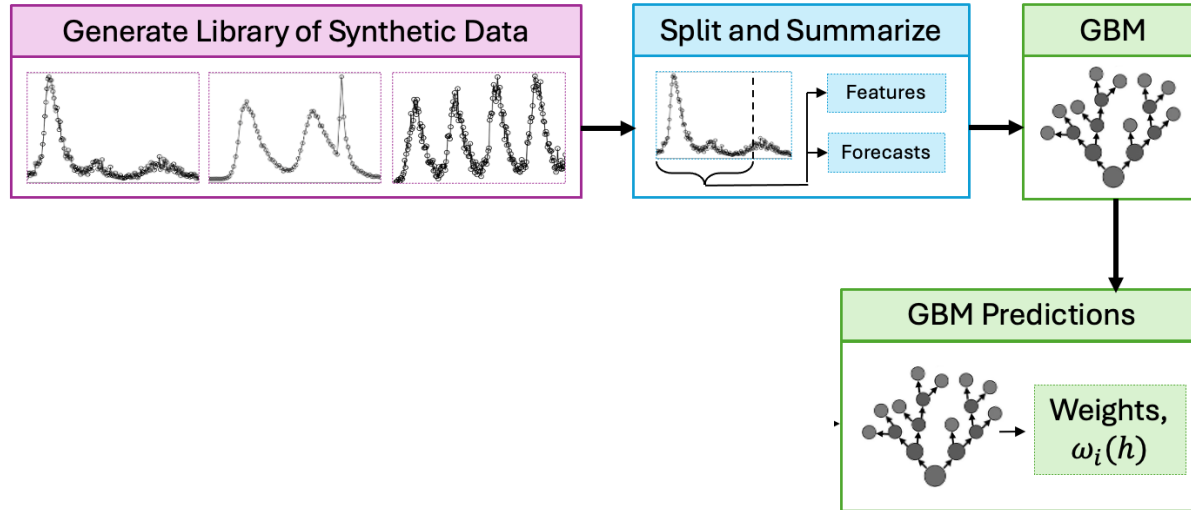
epiFFORMA: data-driven ensembling for infectious disease forecasting



epiFFORMA: data-driven ensembling for infectious disease forecasting



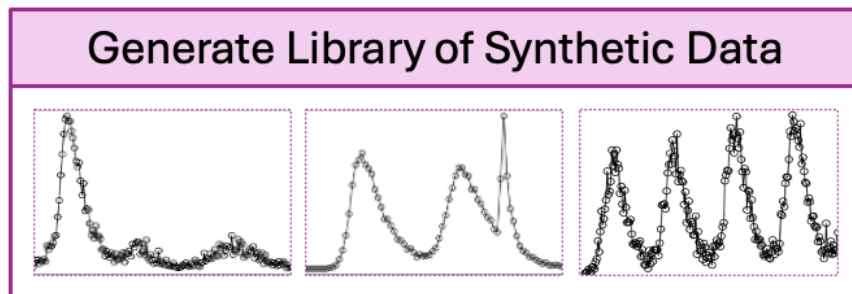
epiFFORMA: data-driven ensembling for infectious disease forecasting



New to epiFFORMA: Use of synthetic data

The Synthetic Data Library

We wish to represent many **possible disease dynamics** without catering to a specific disease.



Generate seasonality components

Generate peaks, troughs, etc.

We mostly used combinations of compartmental models and random effects via sinusoidal functions

Evaluating epiFFORMA

$$\textit{Ensemble Forecast} = \sum_k \mathbf{w}_k(\mathbf{x}) \textit{Forecast}_k$$

epiFFORMA: $w_k(x) = \text{GBM-estimated weight}$

Equal Weights: $w_k(x) = 1/K$

Component c : $w_k(x) = I(k = c)$

Will evaluate using data across **10 different diseases!!!!**

Component Models

rw:

theta:

arima:

gam:

moa:

moa-deriv:

meanfcst:

mirror:

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw: random walk forecast, with drift = FALSE;

theta:

arima:

gam:

moa:

moa-deriv:

meanfcst:

mirror:

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw:

theta: equivalent to simple exponential smoothing with drift;

arima:

gam:

moa:

moa-deriv:

meanfcst:

mirror:

gam2mirror:

From the **forecast** package
in R.

Component Models

rw:

theta:

arima: autoregressive integrated moving average;

gam:

moa:

moa-deriv:

meanfcst:

mirror:

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw:

theta:

arima:

gam: generalized additive models;

moa:

moa-deriv:

meanfcst:

mirror:

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw:

theta:

arima:

gam:

moa: method of analogues, using synthetic data;

moa-deriv:

meanfcst:

mirror:

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw:

theta:

arima:

gam:

moa:

moa-deriv: the **moa** method on the derivative scale;

meanfcst:

mirror:

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw:

theta:

arima:

gam:

moa:

moa-deriv:

meanfcst: mean of the time series up through two years before the
start of the forecast;

mirror:

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw:

theta:

arima:

gam:

moa:

moa-deriv:

meanfcst:

mirror: predictions symmetric about the start of a forecast, out through the forecast horizon;

gam2mirror:

} From the **forecast** package
in R.

Component Models

rw:

theta:

arima:

gam:

moa:

moa-deriv:

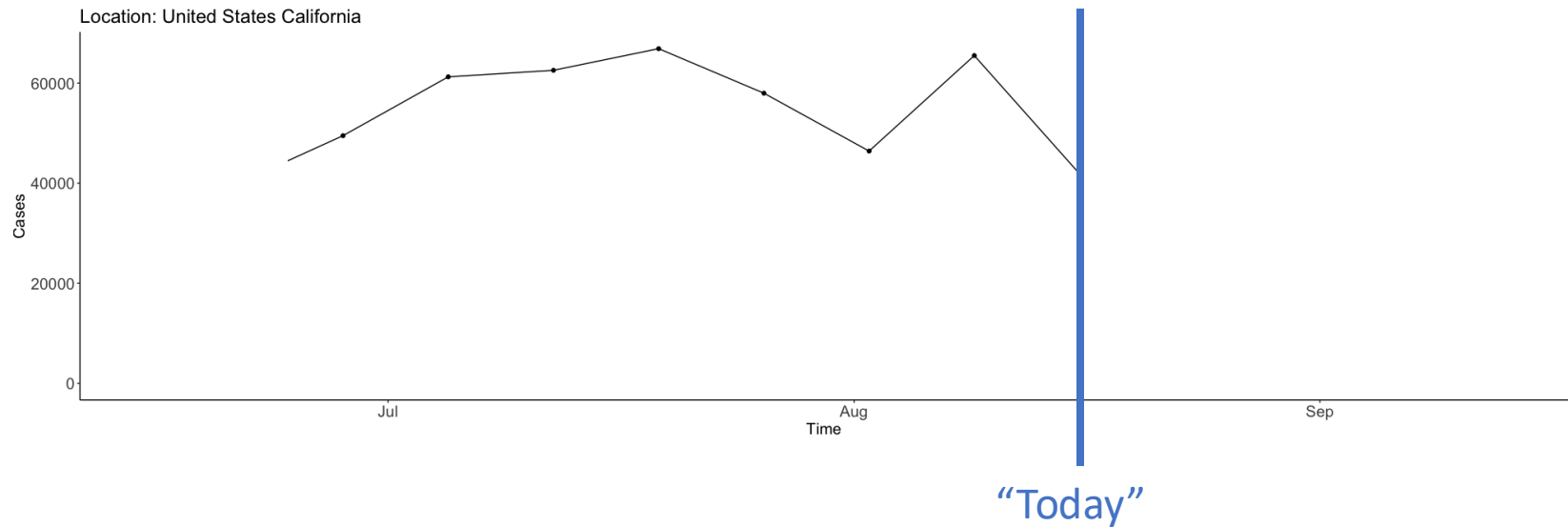
meanfcst:

mirror:

gam2mirror: a linear combination of the **gam** and the **mirror** models.

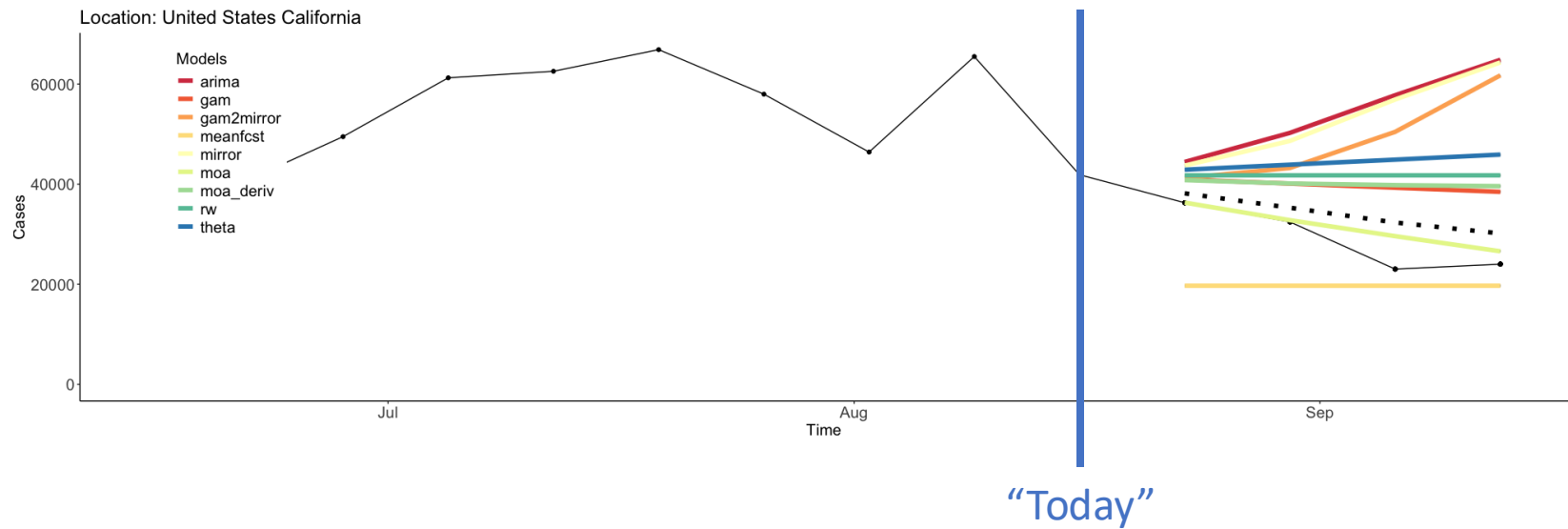
} From the **forecast** package
in R.

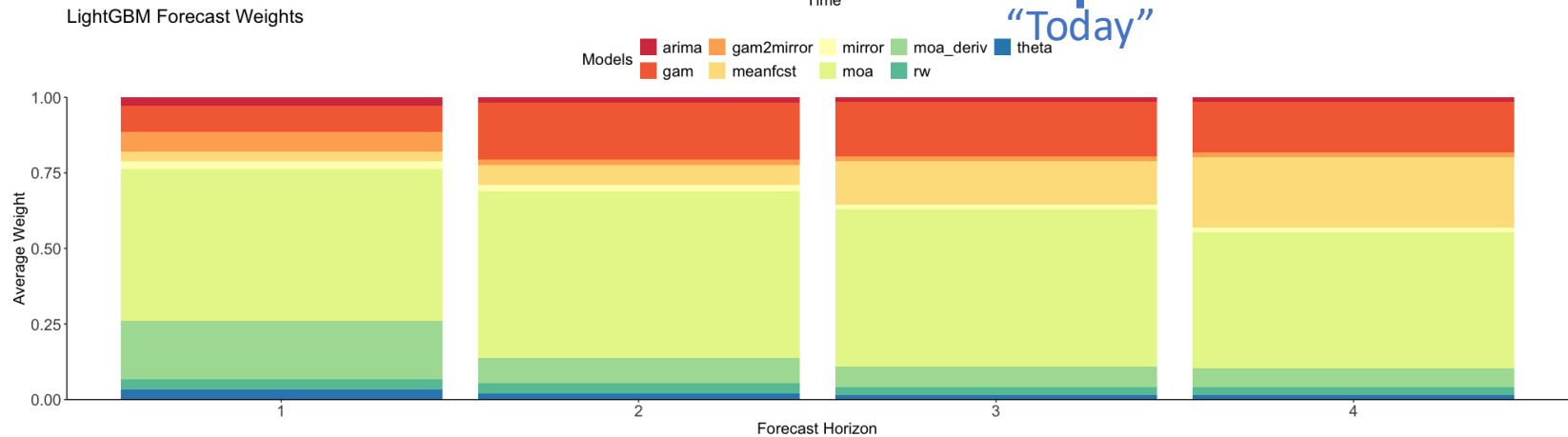
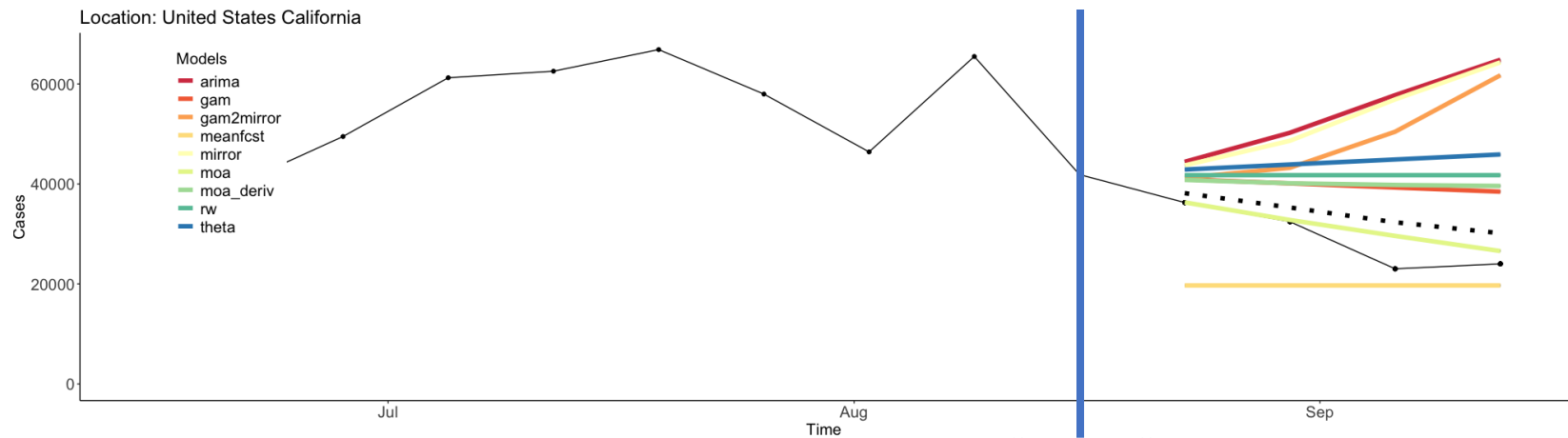
epiFFORMA Results



Initial example of epiFFORMA: COVID-19 cases in California.

- **Starting in early September 2020.**
- **Forecast 4 weeks into the future.**





epiFFORMA Results

(now across 10 diseases)

COVID-19 (US)

COVID-19 (Global)

ILI (US)

Diphtheria (US)

Measles (US)

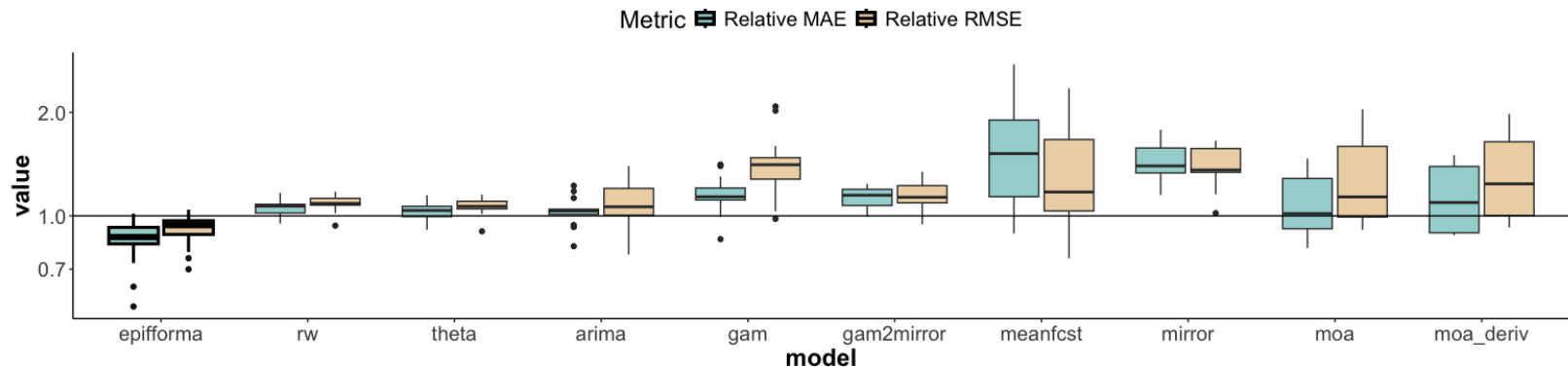
Mumps (US)

Polio (US)

Rubella (US)

Smallpox (US)

Chikungunya (Brazil)



Recall:

epiFFORMA:

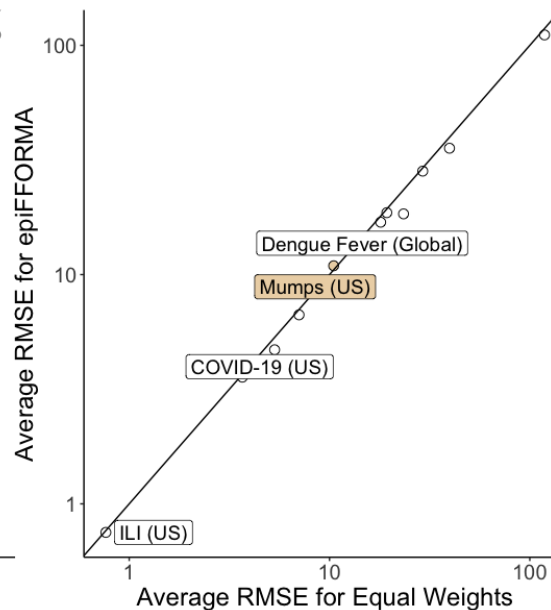
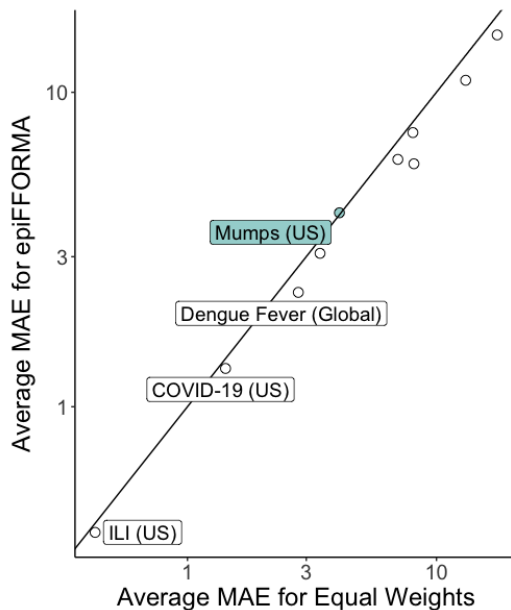
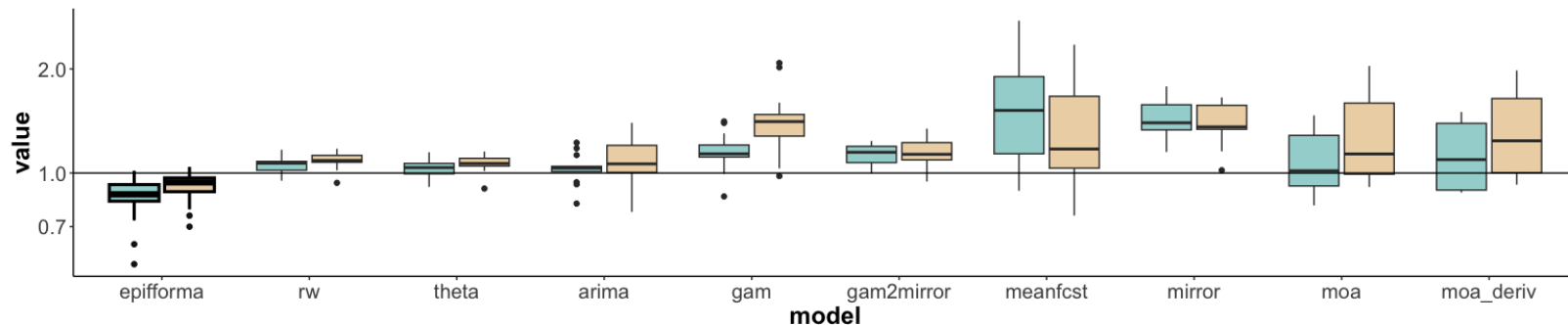
$$w_k(x) = \text{GBM-estimated weight}$$

Equal Weights:

$$w_k(x) = 1/K$$

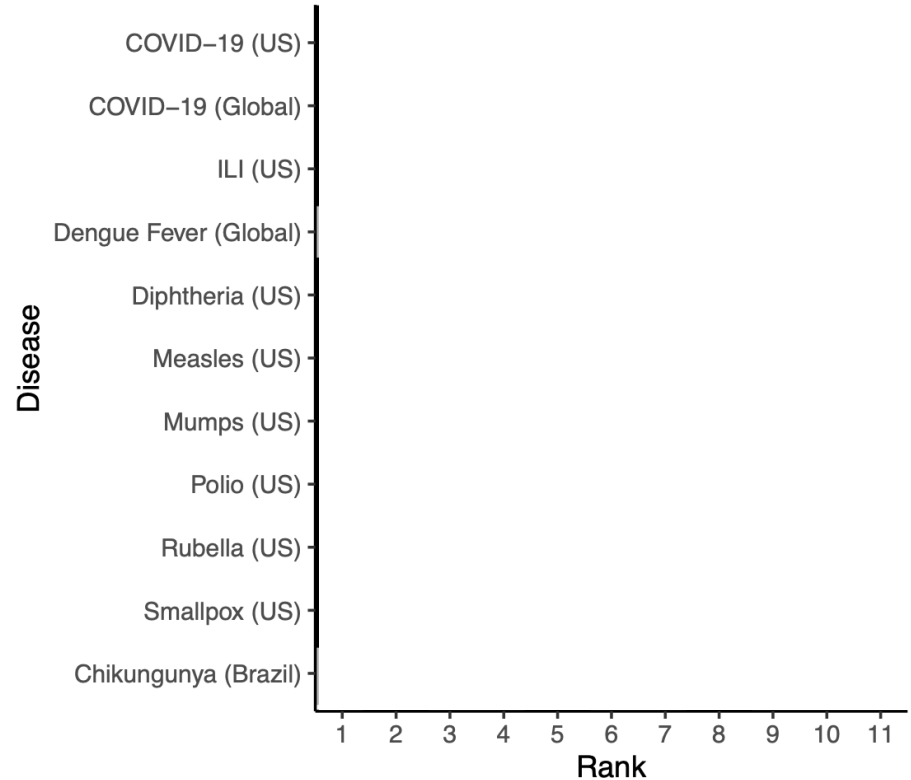
Component c :

$$w_k(x) = I(k = c)$$



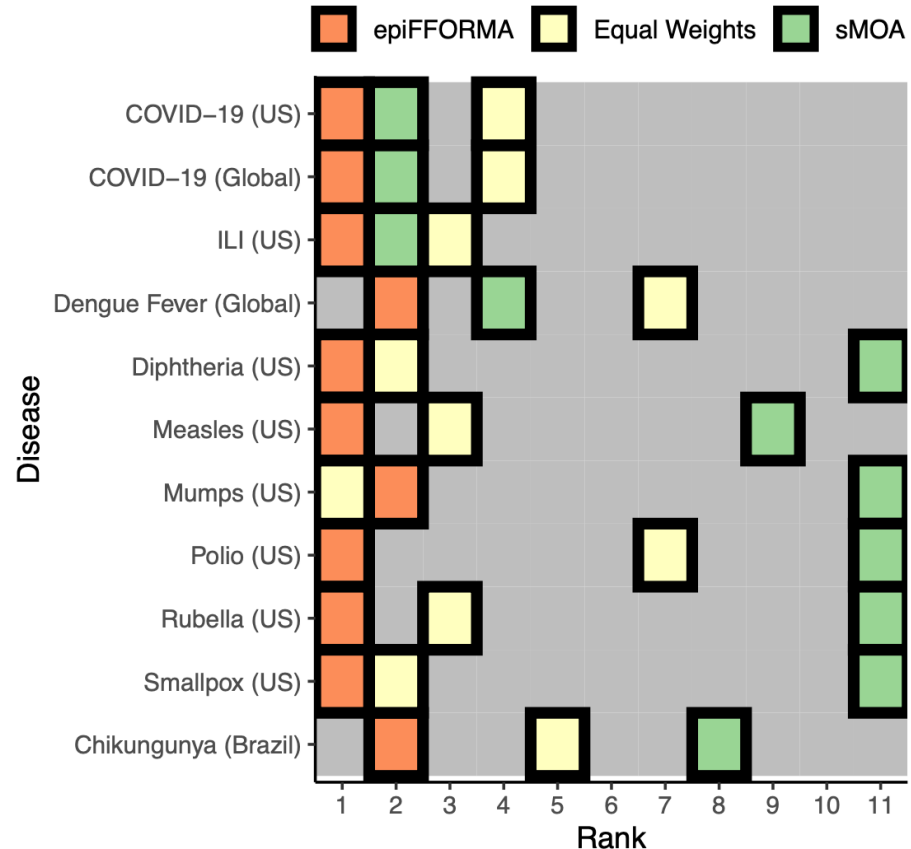
By disease, epiFFORMA outperforms the equal-weights ensemble for every disease except Mumps.

Overall Forecast Rankings (MAE)



**epiFFORMA outranks
other methods**

Overall Forecast Rankings (MAE)



Uncertainty Quantification for epiFFORMA

$$\textit{Ensemble Width} = \sum_k \omega_k(\boldsymbol{x}) \textit{Width}_k$$

Ensemble
95% Interval Width

Component Model
95% Interval Widths

Uncertainty Quantification for epiFFORMA

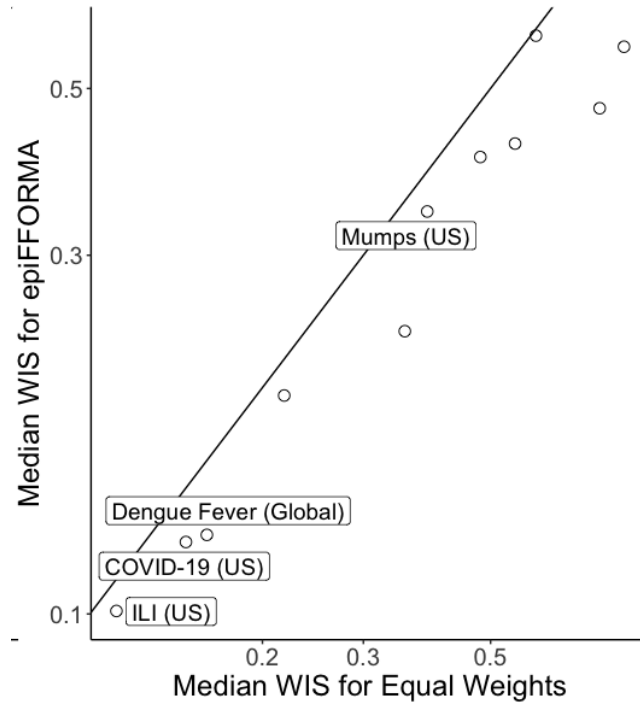
$$\textit{Ensemble Width} = \sum_k \omega_k(\mathbf{x}) \textit{Width}_k$$

Ensemble
95% Interval Width

Component Model
95% Interval Widths

**epiFFORMA has smaller WIS
than equal weights model**

Comparing UQ for epiFFORMA by Disease



Back-up Slides

