

# Statistical Modelling of Infectious Diseases: Influenza and the “Next Disease”

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## We want to improve disease forecasting

### Better statistical forecasting includes:

- Understanding of underlying **assumptions**
- Incorporation of **high-quality data**
- Attention to **variance** of forecasts

### Our work:

- Empirical Bayes model to forecast the flu
- Exploring agent-based models (ABMs)
- Visualizing the spread of disease

## Disease is costly

For influenza in the US alone every year,

- Tens of thousands of lives are lost
- Nearly **\$80B** in healthcare costs
- 31.4 million outpatient visits

**Improved forecasting can alleviate these costs!**

## Forecasting the flu in the US

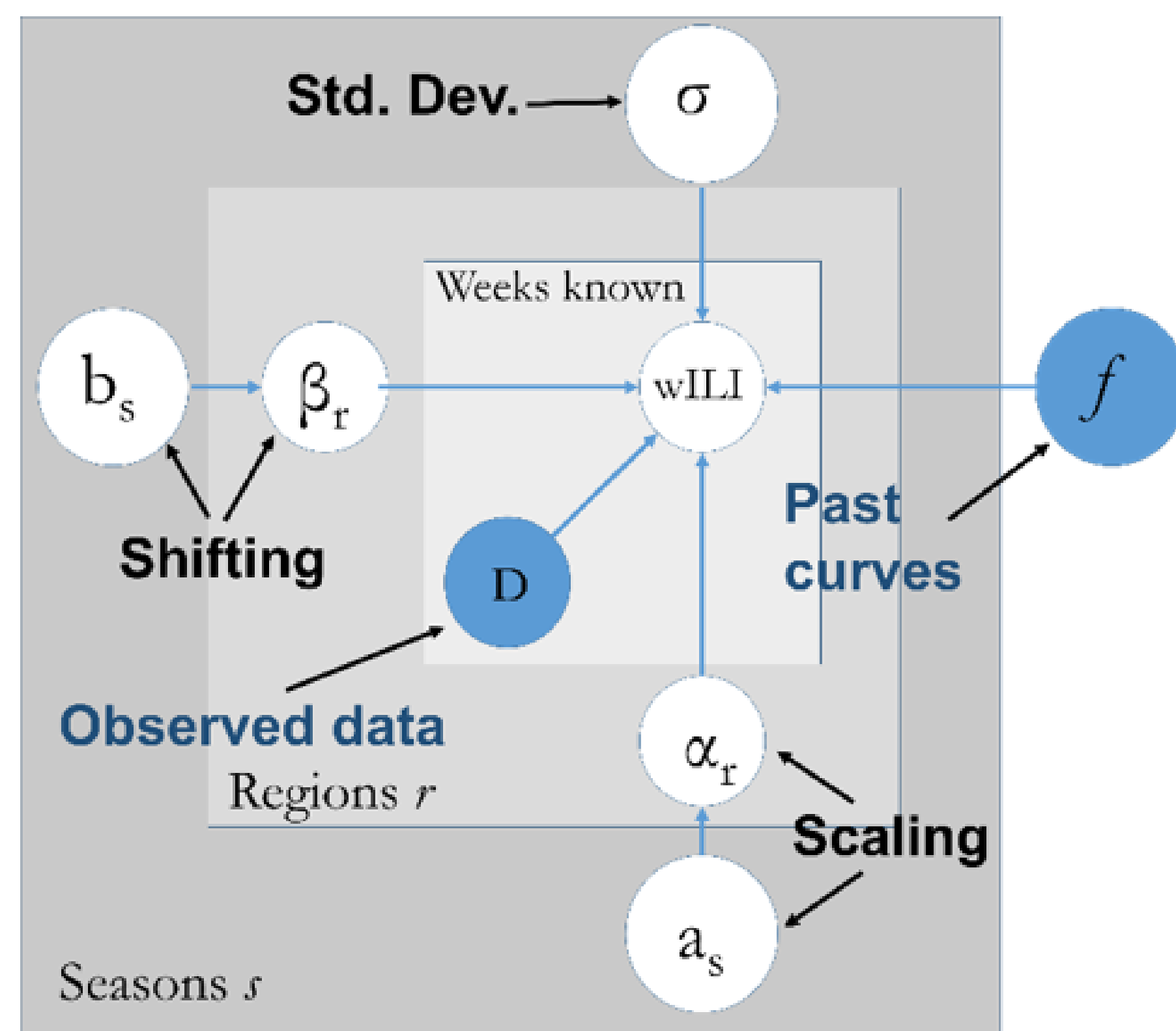
**Goal:** Forecast **wILI** (weighted influenza-like illness),

- For every week in the season (with new data each week)
- For 10 regions in the US
- Emphasis on peak week and peak wILI

**Idea:** A new flu season is going to look like a past one, with some scaling and shifting, *and* regions are dependent on one another

**The model:** Empirical Bayes with Regional Effects

**Figure:** Illustration of the model, consisting of seasonal and regional parameters. New data is incorporated each week.



The model:  $wILI_t^{(r,s)} = [a_s \cdot \alpha_r] \cdot f(t - b_s - \beta_r) + \epsilon_t$ , with  $\epsilon_t \stackrel{iid}{\sim} N(0, \sigma^2)$ , for  $r = 1, 2, \dots, R$  and  $s = 1, \dots, S$  and uniform priors for the parameters.

## Forecasting the flu in the US (continued)

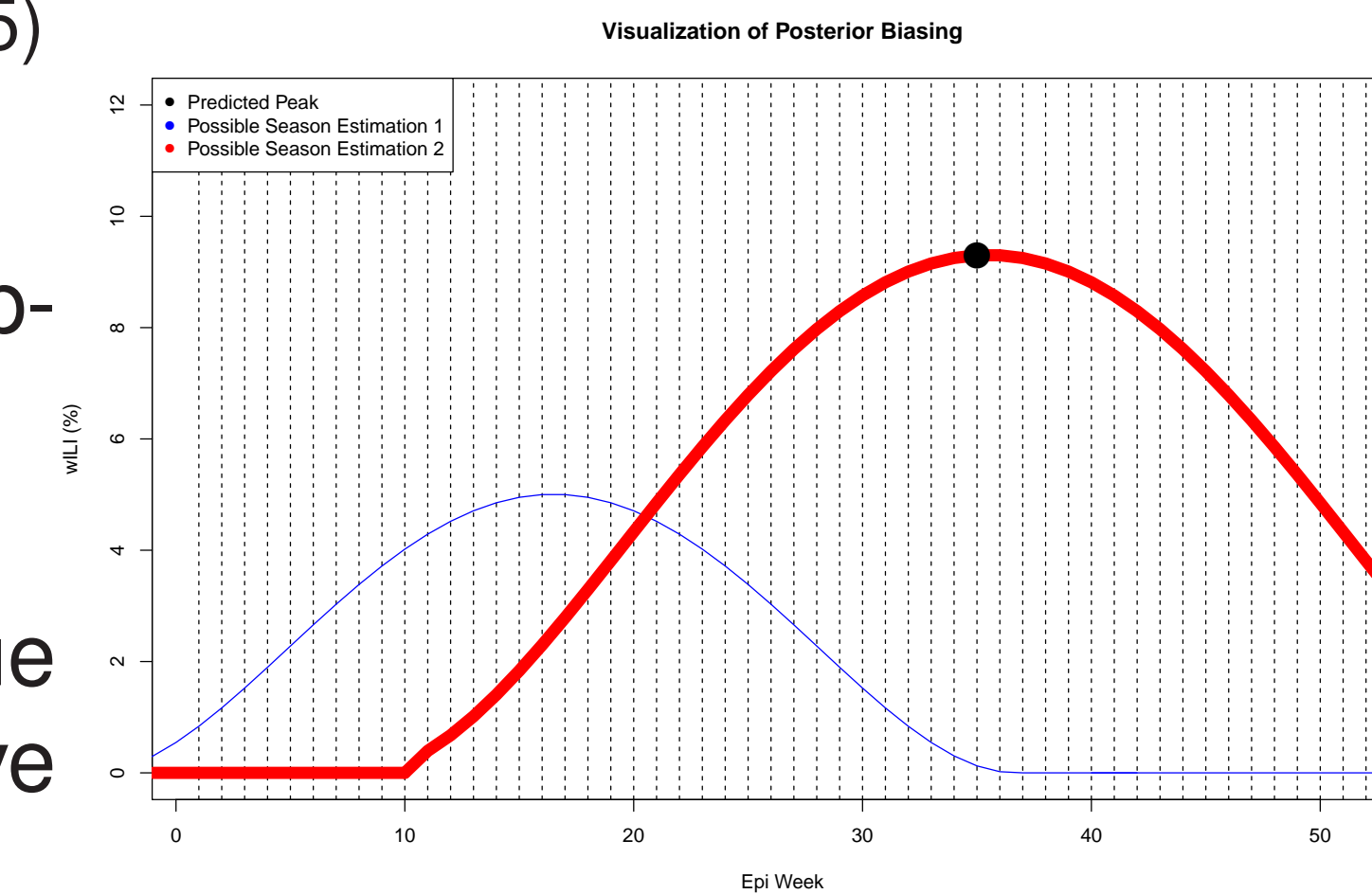
**EB:** Empirical Bayes (let  $\alpha_r, \beta_r = 1$  for all  $r$ ) (Brooks et al., 2015)

**EBR:** Empirical Bayes with regional effects

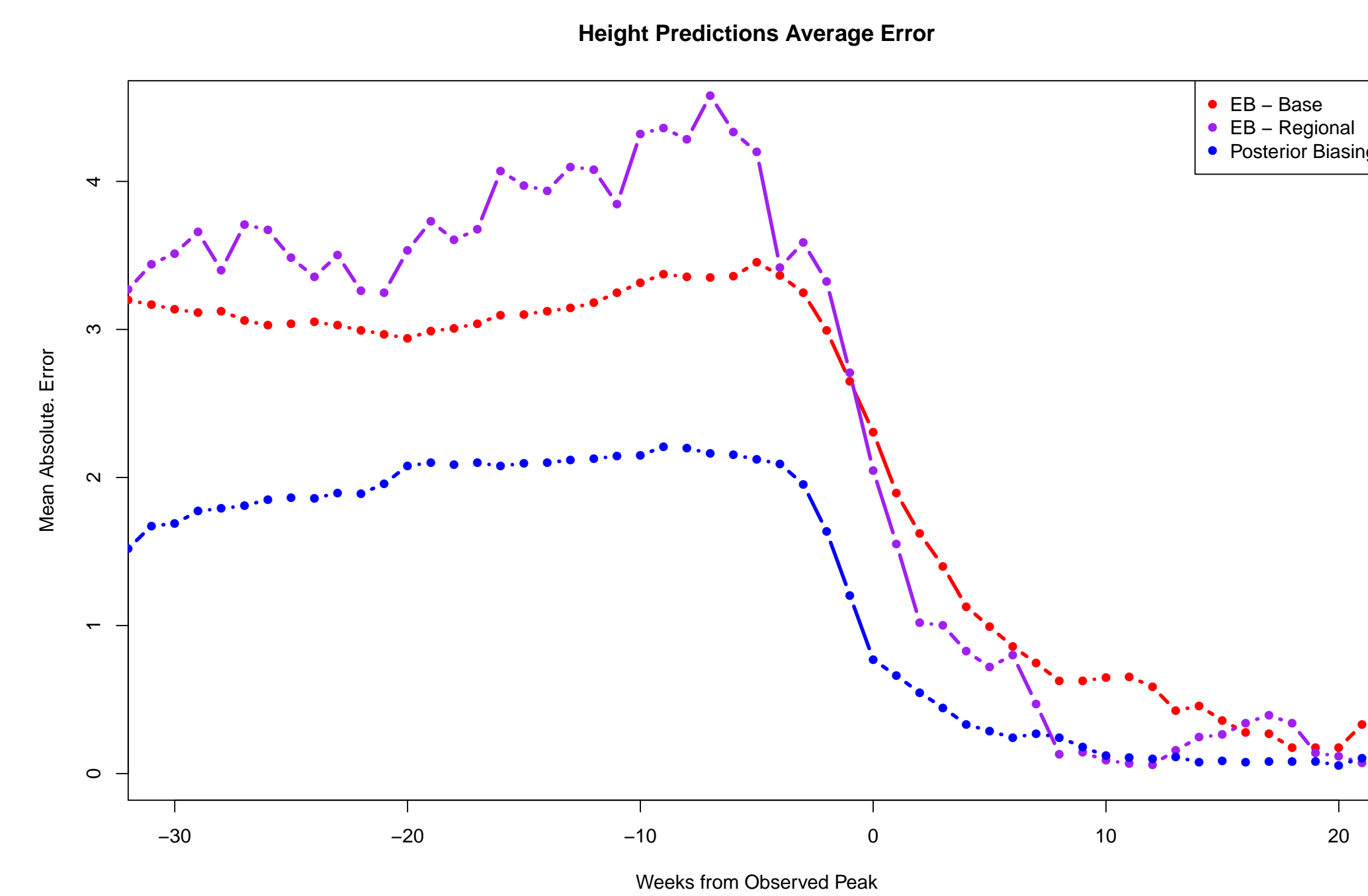
**Problem:** EBR is intractable and we are forced to make approximations for  $\beta_r$

**Solution:** New approach: posterior biasing (**PB**)

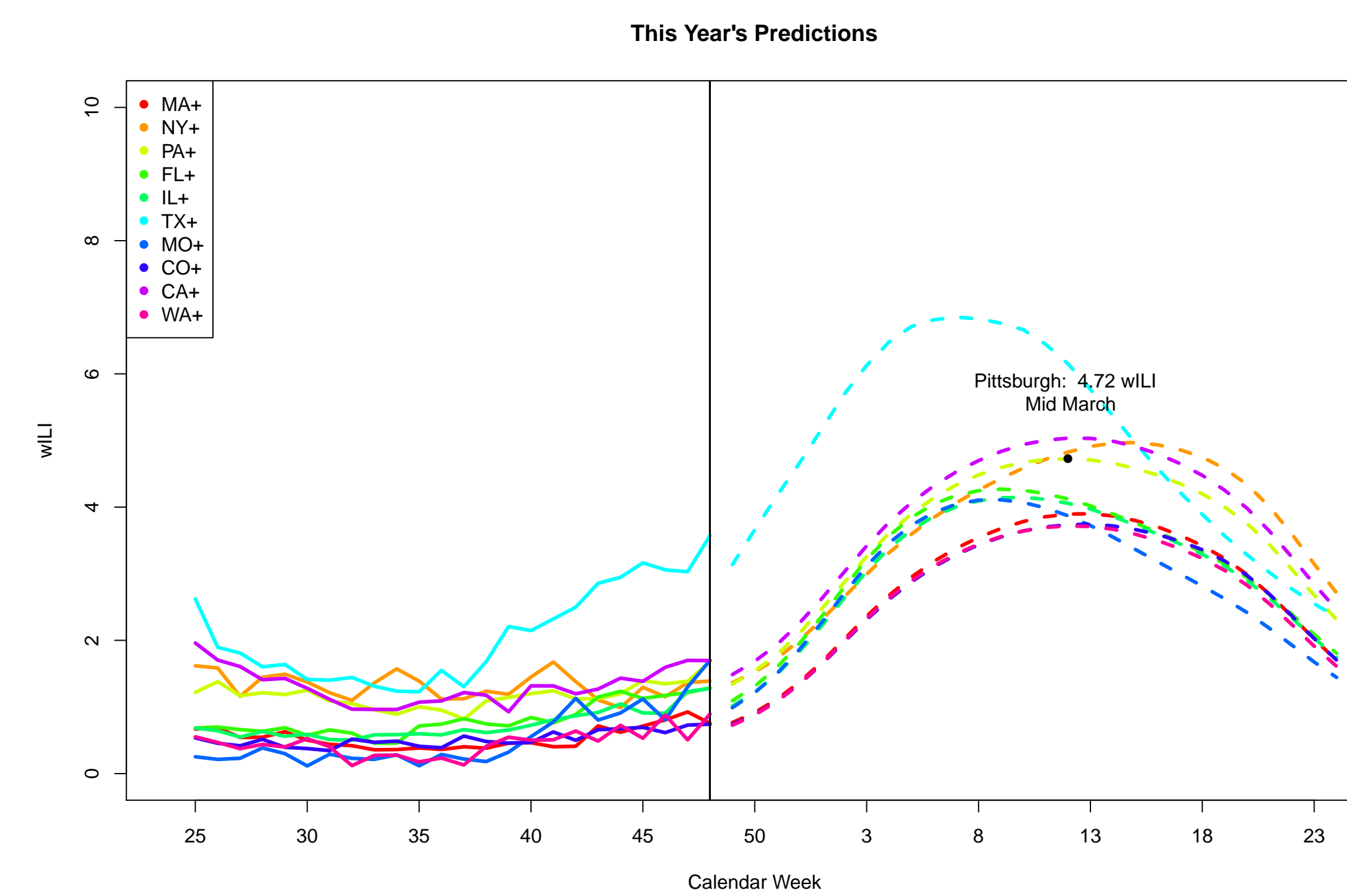
**Figure:** Illustration of PB. The black dot is the peak. The blue curve is given less weight than the red one because the curve goes through the predicted peak.



**Results:** PB yields improved results, while EBR suffers from the approximation we made



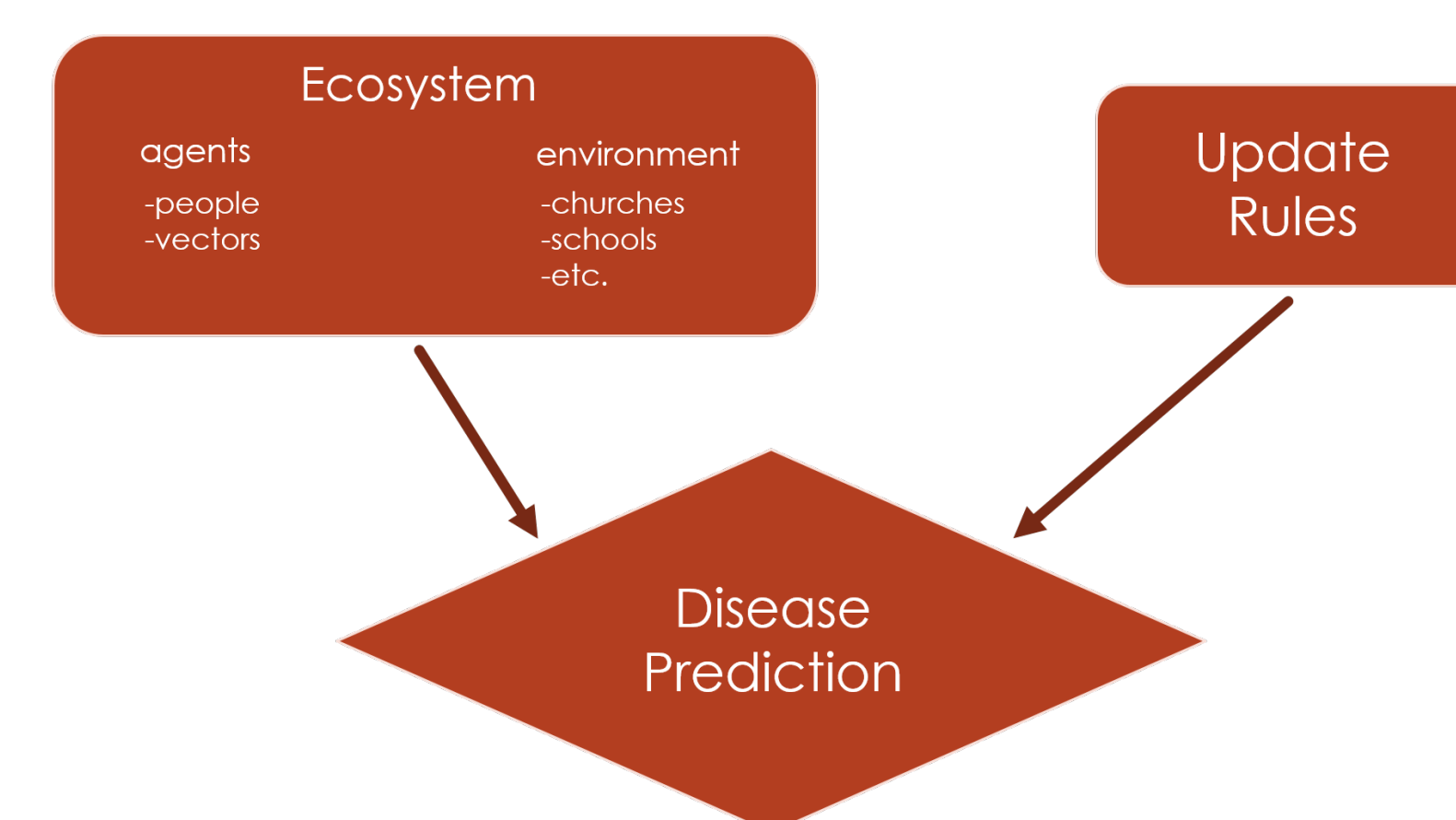
**Figure:** Leave one-season-out cross-validation (CV) of the three models. We calculate the CV for each week from the observed week as we expect better results the closer we are to the peak week



**Figure:** Current predictions for the (new) 2015-2016 Flu season with 25 weeks observed. The Texas region is predicted to be effected more intensely and sooner than the other regions

## The “next” disease and agent-based models (ABMs)

ABMs are a viable way to forecast new diseases (e.g. Dengue, Ebola, Zika, the “next” one)



For new infectious diseases, we have

- Little data
- Less knowledge
- Frenzied awareness
- Few if any models

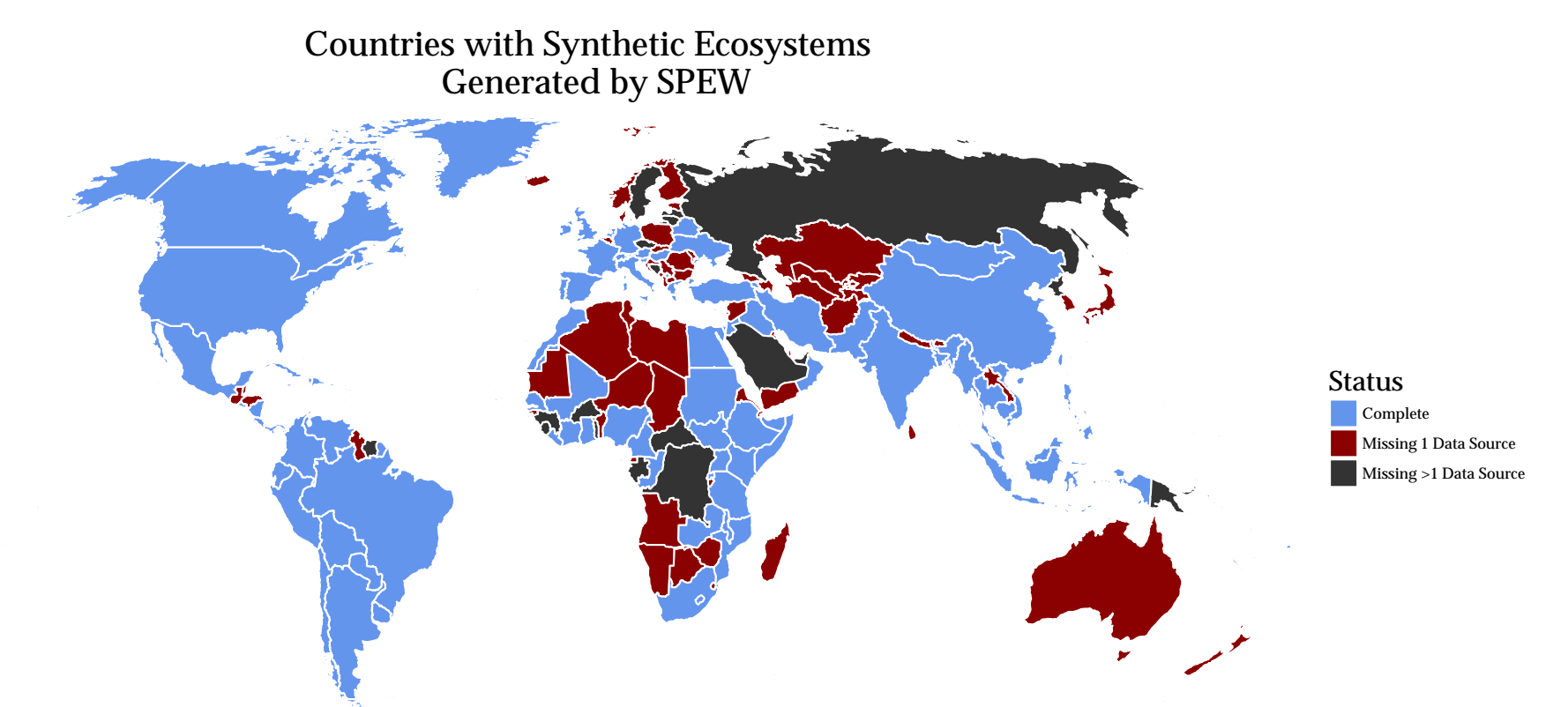
**Solution:** Simulate the spread of disease using ABMs!

## SPEW: Synthetic Pop. & Ecosystems of the World

ABMs require **synthetic ecosystems** as input!

**SPEW** has generated high-quality input for ABMS:

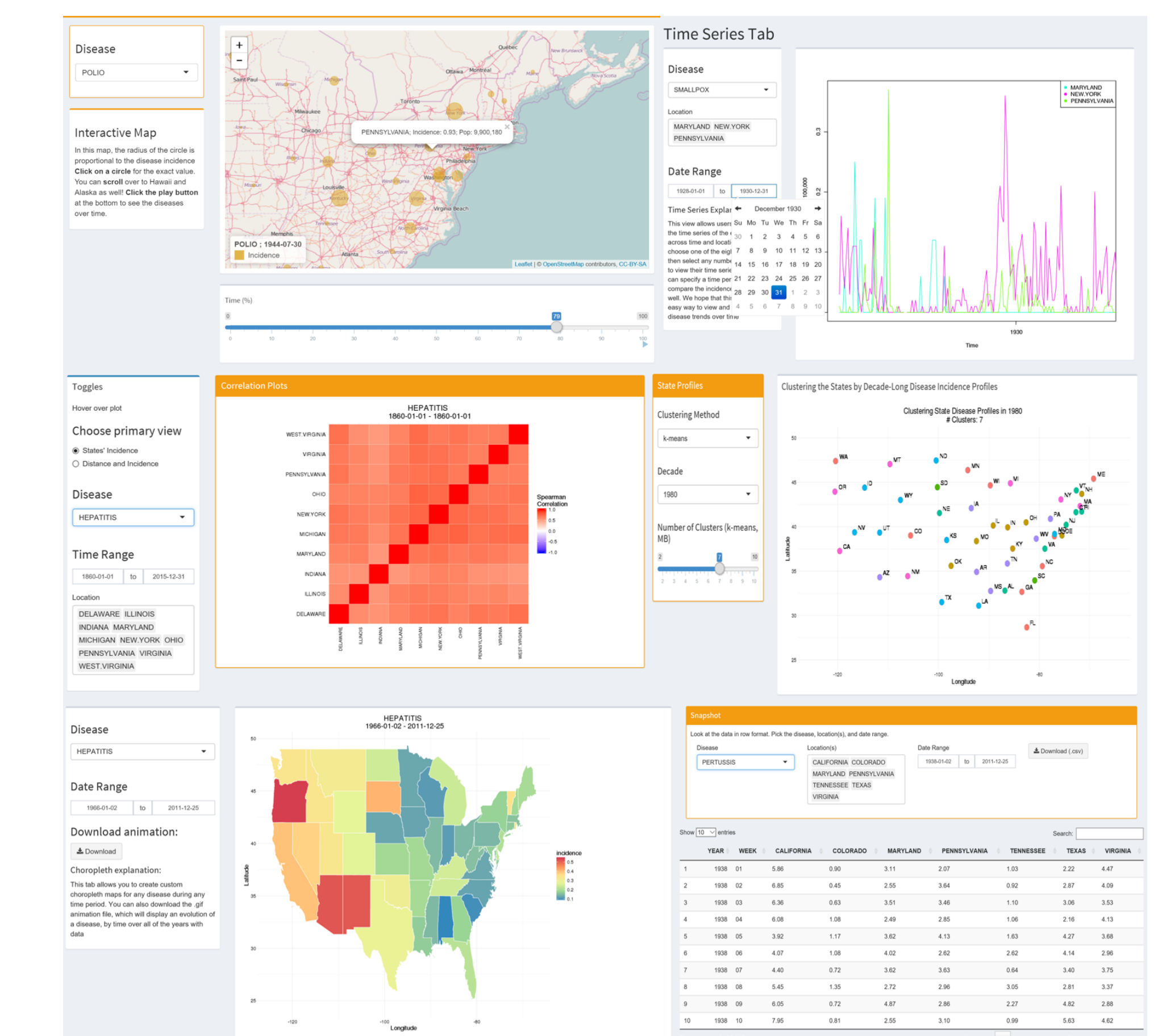
- nearly 5 billion human agents
- 70+ countries and counting!
- automatic diagnostic reports



**Figure:** Synthetic ecosystems available at [epimodels.org](http://epimodels.org)

## SPEW View

[shiny.stat.cmu.edu:3838/sgallagh/spewview/](http://shiny.stat.cmu.edu:3838/sgallagh/spewview/)



**Figure:** SPEW View. Data visualization of historical diseases in the US. **1st place** in Pittsburgh Supercomputing Center Public Health Hackathon

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