

# EPIDEMIC FORECASTING

## Review of the state of the art

---

Dexter Barrows

April 9 2015

Department of Mathematics and Statistics  
McMaster University

Introduction

Techniques

- Phenomenological

- Mechanistic / semimechanistic

Data assimilation

Measuring prediction accuracy

## INTRODUCTION

---

## Basics

- Prediction of future values in a time series
- Based on mechanistic understanding, data, mix

## Outbreak type

- New disease
  - Scarcity of information is key concern
  - Forecasting extremely difficult
- Established disease
  - Long time series, likely better biological understanding
  - Short-term forecasting is easiest (information plentiful)
  - Long-term forecasting possible, integration of weather/socio-economic factors important

## TECHNIQUES

---

3 main families

- Phenomenological - pure inference from data
- Mechanistic - capture “drivers” of disease spread
- Semi-mechanistic - integration of data into model

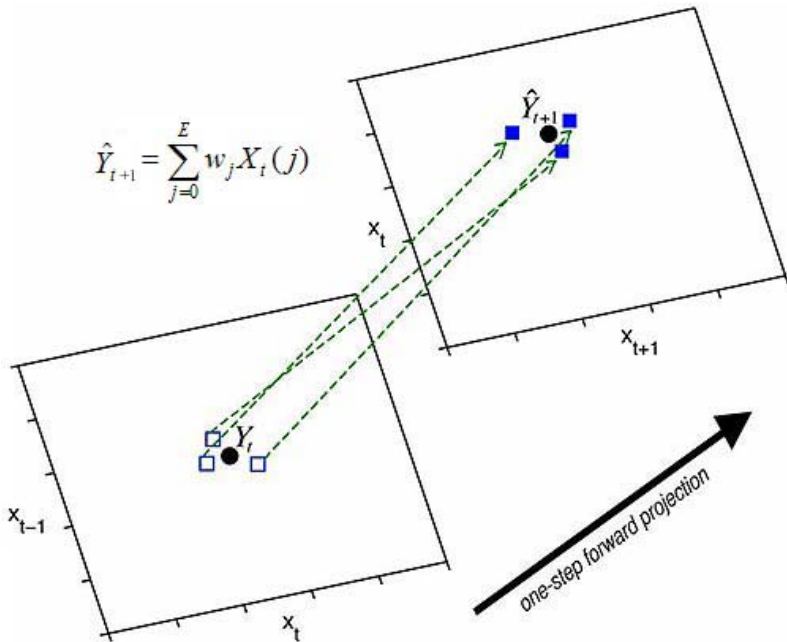
- **AutoRegressive Integrated Moving Average**
- Purely phenomenological
- Assumes linear process, Gaussian distributions
- 3-parameter process  $ARIMA(p, d, q)$ , indicating order of
  - $p$  - Autoregressive
    - Linear combination of past terms
  - $d$  - Integrated
    - Used to remove the trend - makes the series stationary
  - $q$  - Moving average
    - Dependence on past error
- Orders are determined using
  - $ACF$  - works on consecutive elements in series (correlation)
  - $PACF$  - works on additional predictor variables (conditional correlation)

- Adaptation of ARIMA used to capture seasonal effects
- Usually expressed as  $SARIMA(p, d, q) \times (P, D, Q)_s$ 
  - $P, D, Q$  are *seasonal* orders
- Orders are determined using
  - $ACF$  and  $PACF$  as before
  - Also Periodic  $ACF$  (every  $k$  elements)



- Construct a “library” of consecutive time lag vectors  $\{x_i\}$  of some length  $E$  and corresponding forward trajectories  $\{y_i\}$
- Use similar past system states with **known** outcomes to project to **unknown** future state
  - A weighted linear combination of closest vectors
- Weightings are exponential, function of distance

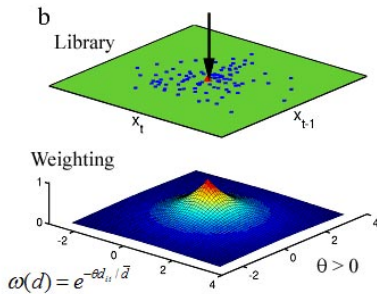
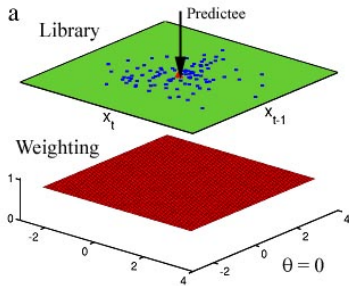
[2]



<http://simplex.ucsd.edu/>

- Sequentially locally weighted global linear maps (S-map)
- Designed to handle linear, locally nonlinear time series
- Similar to Simplex projection
  - But **all** vectors are used for projection
- Weightings are again exponential

[2][11]

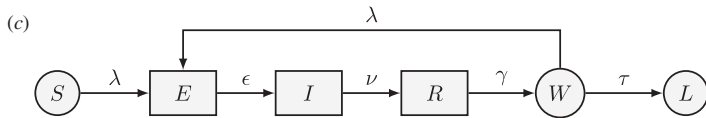
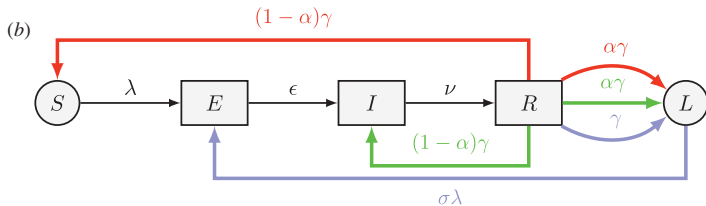
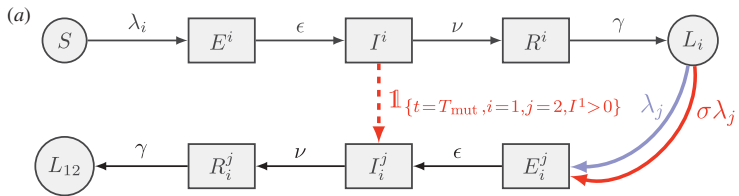


<http://simplex.ucsd.edu/>

- Extensively used model in epidemiology
- Division into classes: **S**usceptible-**I**nfected-**R**emoved
- Transition between states

$$\begin{aligned}\frac{dS}{dt} &= -\frac{\beta IS}{N} \\ \frac{dI}{dt} &= \frac{\beta IS}{N} - \gamma I \\ \frac{dR}{dt} &= \gamma I\end{aligned}$$

- Many extensions exist
  - Additional classes
  - Additional mechanistic terms

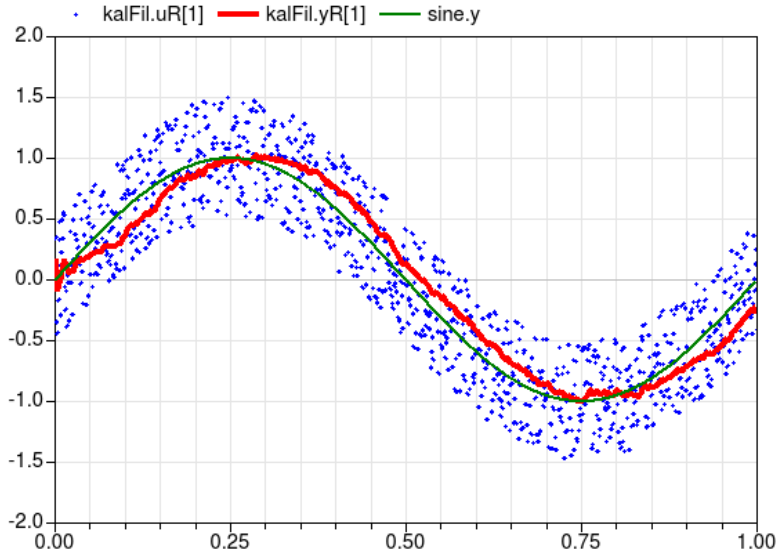


Camacho et al, 2011

- SIR-based models *may* require many parameters to be estimated
  - Not a problem if statistical caution is exercised
- Over-fitting a particular problem - can reduce forecasting ability
- More model complexity = longer time series required
- Iterated filtering methods can estimate parameters in addition to producing forecasts

- Designed to operate on linear models, assumptions:
  - Underlying dynamics are linear
  - Error distributions are normal (or close to it)
- Uses knowledge of underlying dynamics (ex. SIR model)
- Operation in cyclical phases
  - Prediction → projection forward
  - Update → observed data used to refine estimation mechanism

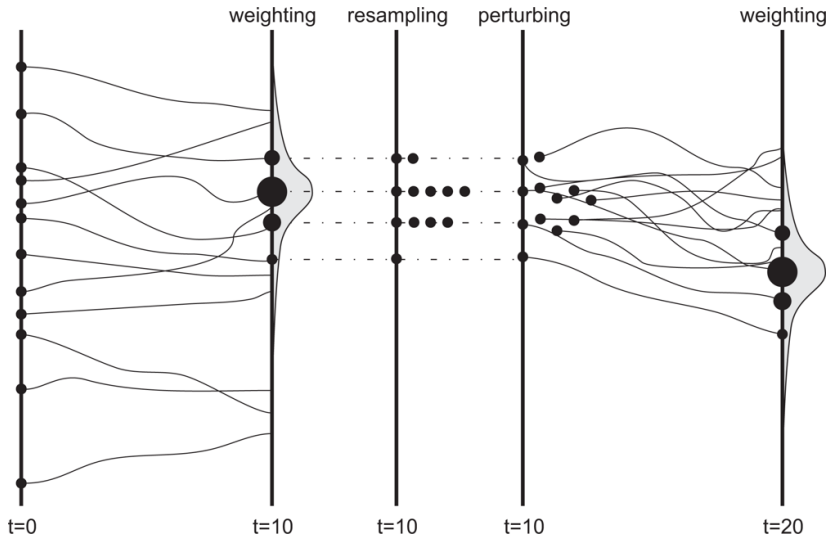




[http://simulationresearch.lbl.gov/modelica/releases/latest/help/Buildings\\_Uilities\\_IO\\_Python27\\_](http://simulationresearch.lbl.gov/modelica/releases/latest/help/Buildings_Uilities_IO_Python27_)

- Extended Kalman filter (*EKF*)
  - Linearises about the estimate of current mean and covariance
- Ensemble Kalman filter (*EnKF*)
  - Uses a cohort of ensemble members, their sample mean and covariance
  - Still assumes linear process / Gaussian distributions
  - Useful for large number of parameters
- Ensemble Adjustment Kalman filter (*EAKF*)
  - Combination of *EKF* and *EnKF*
    - Linearises as in *EKF*
    - Ensemble members as in *EnKF*

- Uses a set of particles, similar to *EnKF* cohort
- Makes no assumption about the distributions involved in the system
- Particle importance using weights
- Problem: Particle degeneracy
  - When one particle accumulates most of the weight
  - Avoided via resampling at each iteration



[http://www.mdpi.com/sensors/sensors-12-16291/article\\_deploy/html/images/sensors-12-16291f2-1024.png](http://www.mdpi.com/sensors/sensors-12-16291/article_deploy/html/images/sensors-12-16291f2-1024.png)

- Maximum likelihood via iterated filtering (MIF or IF1)
  - Uses multiple rounds of particle filtering
  - Stochastic perturbation of parameters
  - Each round pushes the parameter estimates toward ML
- Particle Markov chain Monte Carlo (pMCMC)
  - Uses an MCMC method constrain model parameters
  - Particle filter between each MCMC iteration
- IF2 (MIF2)
  - Evolution of MIF (IF1)
  - Uses stochastic perturbation as before, also data cloning
  - Looks to consistently outperform IF1 and pMCMC

# DATA ASSIMILATION

---

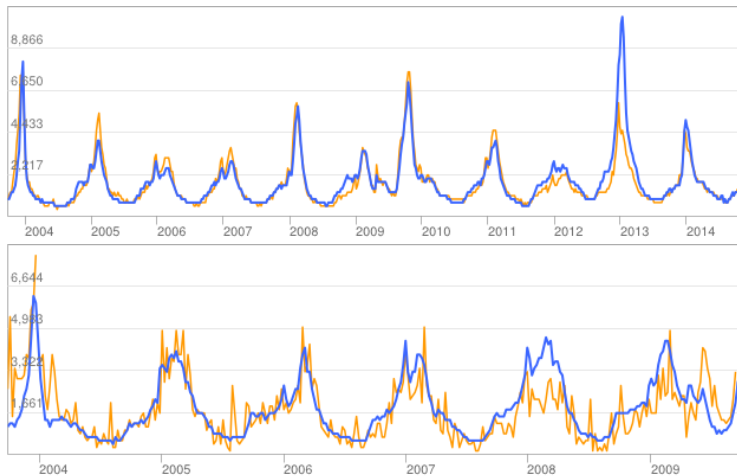
## Primary Sources

- Google Flu Trends (GFT)
  - Uses search trend data to infer incidence rates
  - Almost instantaneous, but less accurate
  - Currently up to 29 countries
- Governments ex. Centres for Disease Control (CDC)
  - Regional data (10 regions across the US)
  - Broken down further by age
  - More accurate than GFT, but lag of 1-2 weeks
- WHO

## Social Media

- Twitter: Influenza, Korea, 2012
- Social media and informal news: Haiti, Cholera, 2010

# GFT VS CDC FLUNET



<https://www.google.org/flutrends/about/how.html>



- Nearly all infectious disease affected by seasonality
  - Contact
  - Susceptibility
  - Influx of susceptibles
  - Reservoir dynamics / vector dynamics
- Weather data sources
  - National Oceanic and Atmospheric Administration (NOAA)
  - NASA Jet Propulsion Laboratory (JPL)

[12][13]

- El Niño Southern Oscillation
- Sustained anomalous ocean surface temperature in the Pacific
- Unpredictable
- Many effects on local populations
- Relevant to epidemic outbreaks in Southeast Asian locales
  - Cholera in Bangladesh
  - Dengue fever in Singapore

[4][8]

## MEASURING PREDICTION ACCURACY

---

- What to measure
  - Peak timing / intensity
  - Magnitude
  - Duration
- How to measure
  - Correlation coefficients
  - RMSE
  - Confidence intervals
  - Receiver operating characteristic (ROC) curves

- AIC - Akaike Information Criterion
  - Measures relative model quality
  - Rewards goodness-of-fit, penalizes for number of parameters
- BIC - Bayesian Information Criterion
  - Similar to AIC
  - Tends to penalize many parameters more than AIC
- DIC - Deviance Information Criterion
  - Particularly useful when comparing MCMC-based models
- WAIC - Watanabe-Akaike (widely applicable) Information Criterion
  - More “tuned” to prediction

- [1] S. Cook, C. Conrad, A. L. Fowlkes, and M. H. Mohebbi. Assessing Google Flu trends performance in the United States during the 2009 influenza virus A (H1N1) pandemic. *PLoS ONE*, 6(8):1–8, 2011.
- [2] S. M. Glaser, H. Ye, and G. Sugihara. A nonlinear, low data requirement model for producing spatially explicit fishery forecasts. *Fisheries Oceanography*, 23:45–53, 2014.
- [3] A. L. Graham, A. Camacho, F. Carrat, O. Ratmann, B. Cazelles, and L. E.-e. Mathe. Explaining rapid reinfections in multiple- wave influenza outbreaks : Tristan da Cunha 1971 epidemic as a case study. *Proc. R. Soc. B*, 2016.
- [4] Y. Hii, J. Rocklöv, S. Wall, and L. Ng. Optimal lead time for dengue forecast. *PLoS neglected tropical ...*, 6(10), 2012.
- [5] E. L. Ionides, D. Nguyen, Y. Atchadé, S. Stoev, and A. a. King. Inference for dynamic and latent variable models via iterated, perturbed Bayes maps. *Proceedings of the National Academy of Sciences*, 112(3):719–724, 2015.
- [6] E.-K. Kim, J. H. Seok, J. S. Oh, H. W. Lee, and K. H. Kim. Use of hangeul twitter to track and predict human influenza infection. *PloS one*, 8(7):e69305, Jan. 2013.
- [7] E. O. Nsoesie, J. S. Brownstein, N. Ramakrishnan, and M. V. Marathe. A systematic review of studies on forecasting the dynamics of influenza outbreaks. *Influenza and other respiratory viruses*, 8(3):309–16, May 2014.
- [8] R. C. Reiner, A. a. King, M. Emch, M. Yunus, a. S. G. Faruque, and M. Pascual. Highly localized sensitivity to climate forcing drives endemic cholera in a megacity. *Proceedings of the National Academy of Sciences of the United States of America*, 109(6):2033–6, Feb. 2012.
- [9] R. Reyburn, D. R. Kim, M. Emch, A. Khatib, L. von Seidlein, and M. Ali. Climate variability and the outbreaks of cholera in Zanzibar, East Africa: a time series analysis. *The American journal of tropical medicine and hygiene*, 84(6):862–9, June 2011.
- [10] J. Shaman, W. Yang, S. Kandula, K. S. Inference, and S. Leone. Inference and Forecast of the Current West African Ebola Outbreak in Guinea , Sierra Leone and Liberia. pages 1–17, 2014.
- [11] G. Sugihara. Nonlinear Forecasting for the Classification of Natural Time Series. *Philosophical Transactions of the Royal Society A: Mathematical, Physical and Engineering Sciences*, 348:477–495, 1994.
- [12] J. D. Tamerius, J. Shaman, W. J. Alonso, K. Bloom-Feshbach, C. K. Uejio, A. Comrie, and C. Viboud. Environmental Predictors of Seasonal Influenza Epidemics across Temperate and Tropical Climates. *PLoS Pathogens*, 9(3), 2013.
- [13] W. Yang, A. Karspeck, and J. Shaman. Comparison of filtering methods for the modeling and retrospective forecasting of influenza epidemics. *PLoS computational biology*, 10(4):e1003583, Apr. 2014.

THANKS FOR COMING!

QUESTIONS?