

# An iterated block particle filter for inference on coupled dynamic systems

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Slides are at <https://ionides.github.io/talks/aprm24.pdf>

Joint work with Patricia Ning, Jesse Wheeler, Kidus Asfaw, Jifan Li, Joonha Park, Aaron King, Mercedes Pascual

## Question to be addressed

- 1 When can we carry out full-information likelihood-based inference on nonlinear non-Gaussian spatiotemporal partially observed Markov process (SpatPOMP) models? In particular, models for networks of interacting biological population dynamics.
- 2 We introduce the **iterated block particle filter**, currently the most effective algorithm in the spatPomp R package.

# Inference challenges in population dynamics

- 1 Combining measurement noise and process noise.
- 2 Including covariates in mechanistically plausible ways.
- 3 Continuous time models.
- 4 Modeling and estimating interactions in coupled systems.
- 5 Dealing with unobserved variables.
- 6 **Modeling spatiotemporal dynamics.**
- 7 Studying population dynamics via genetic sequence data.

1–5 are largely solved, from a methodological perspective.

6 is our immediate topic.

7 is exciting but not the focus of this talk.

Reviews: Bjornstad & Grenfell (*Science*, 2001); Grenfell et al (*Science*, 2004)

- Consideration of arbitrary dynamic models. The limitations should be our scientific creativity and the information in the data.

Hence, **plug-and-play** methods which need a simulator from the model but not nice closed-form expressions for densities.

- Statistically efficient inference, to extract all the information in the data.

Hence, **likelihood-based** methods.

# Fitting mechanistic models to time series

- Iterated particle filtering via `mif2` in the R package `pomp` enables Masters-level statisticians to do plug-and-play likelihood-based inference for nonlinear, non-Gaussian, partially observed dynamic systems:

<https://ionides.github.io/531w22/>

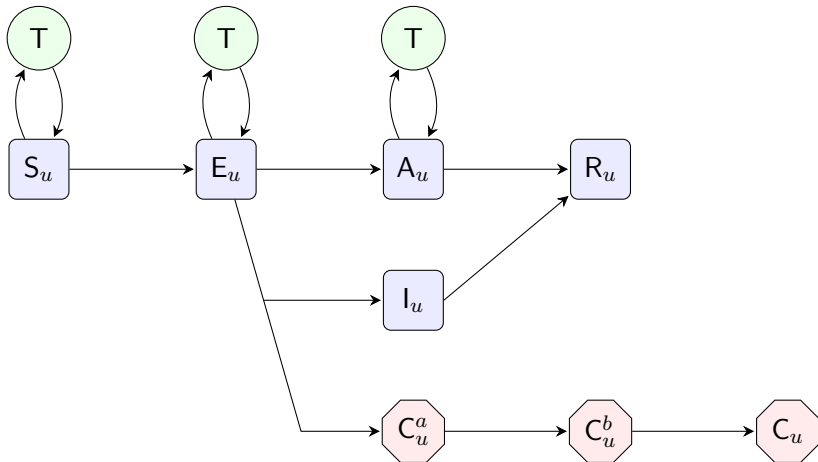
- The science may be hard, but the statistics is becoming routine.

# The curse of dimensionality

- Particle filter (PF) methods fail for high-dimensional systems. They scale exponentially badly.
- Algorithms with improved scalability include:
  - Bagged filters (BF, IBF)**
  - Ensemble Kalman filter (EnKF, IEnKF)**
  - Guided intermediate resampling filter (GIRF, IGIRF)**
  - Block particle filter (BPF, IBPF)**
- Filters estimate latent states and evaluate the likelihood.
- Iterated filters estimate parameters using stochastic parameter perturbations.
- These algorithms are all implemented in the `spatPomp` R package.

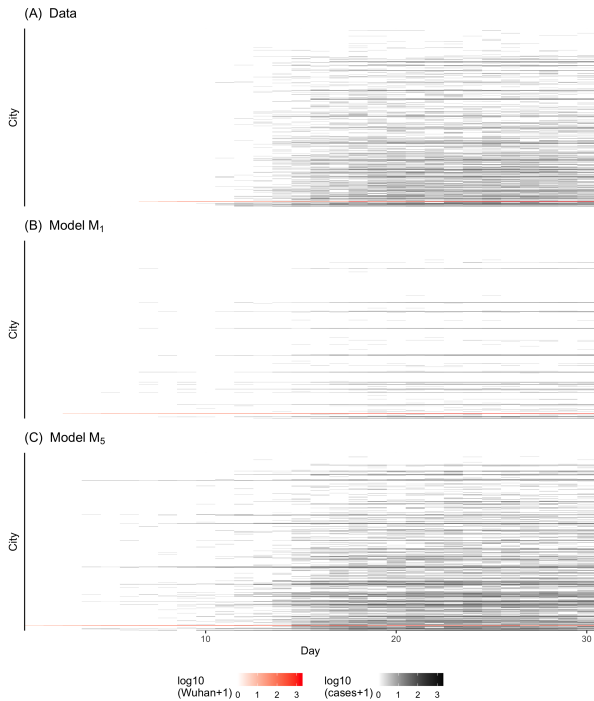
# COVID in 373 cities in China, Jan 10 to Feb 8, 2020

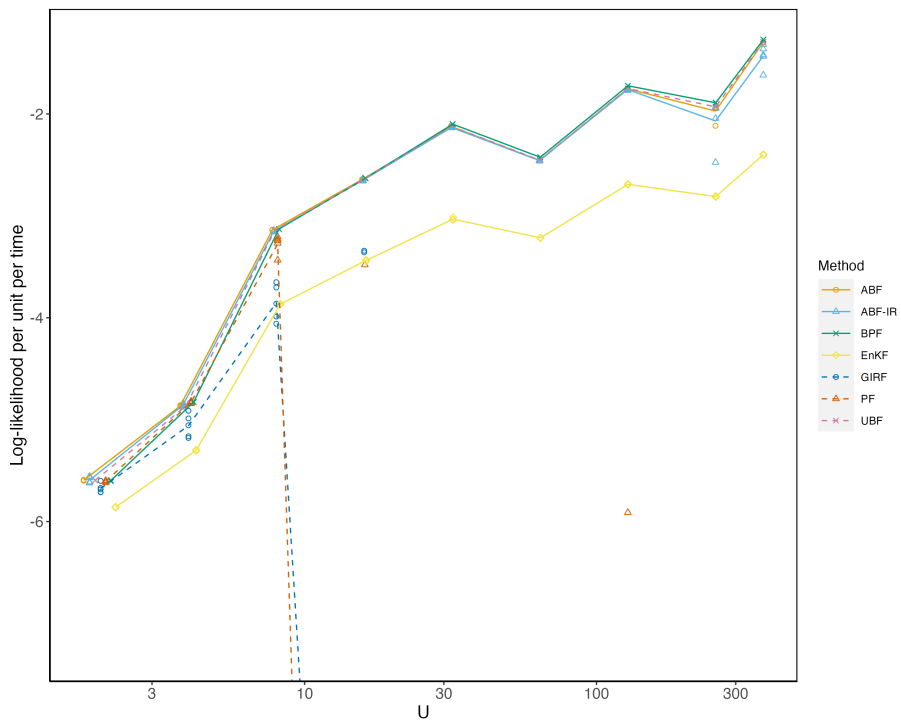
- Metapopulation data were used to infer the fraction of asymptomatic cases and their contagiousness (Li et al, *Science*, May 2020).
- SEIR (susceptible-exposed-infected-removed) model with asymptomatics, reporting delay, and coupling based on cell phone data.
- Li et al (2020) used iterated EnKF for inference.
- The time interval covers the initial China lockdown.
- We present re-analysis of this model and data, recently posted on arXiv.



- Reportably infectious individuals,  $I_u$  for city  $u$ , are included in the delayed reporting compartment,  $C_u^a$ .
- An individual arriving at  $C_u$  is a case report for city  $u$ .
- Individuals in  $A_u$  are not reportable and transmit at a reduced rate.
- Travel occurs to and from  $T$ , based on 2018 data from Tencent.







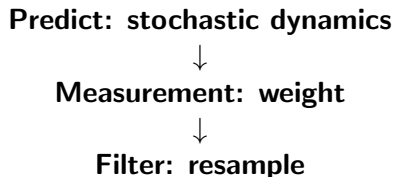
## More on the block particle filter (BPF)

- BPF also worked quickly, easily and reliably on a measles metapopulation, as well as various toy benchmark problems (Ionides et al, *JASA*, 2023).
- BPF has theoretical support in some situations (Rebeschini & Van Handel, *Annals of Applied Probability*, 2015).
- This motivated us to develop an iterated BPF (IBPF) for parameter estimation.
- IBPF has theoretical guarantees similar to BPF (Ning & Ionides, *JMLR*, 2023).
- BPF was independently proposed as the “factored particle filter” by Ng et al (2002).

## Evolutionary analogy



## Particle filter algorithm

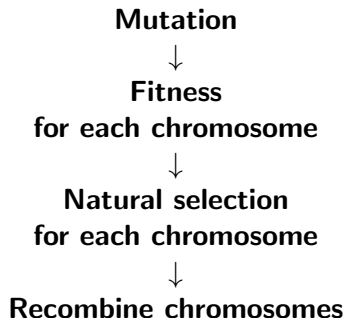


- PF is an evolutionary algorithm with good mathematical properties: an unbiased likelihood estimate and consistent latent state distribution.

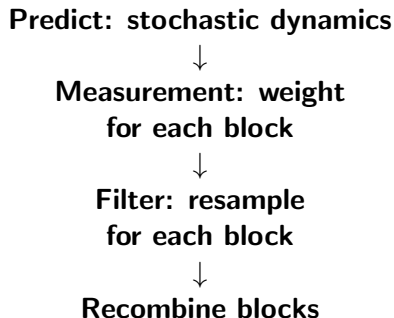
# Block particle filter (BPF)

- Blocks are a partition of the metapopulation units.
- For measles, we use each city as a block.

## Evolutionary analogy



## Block particle filter

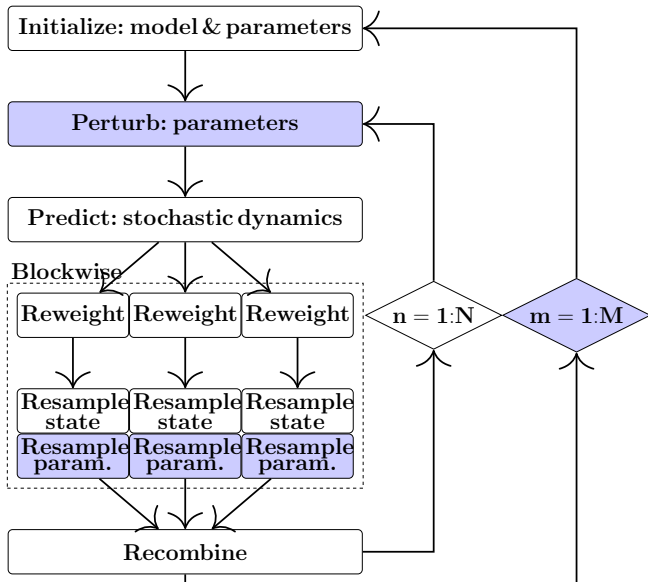


- Blocks are segments of the full state which can be reassorted between particles at the resampling step.

## Comments on the Ensemble Kalman Filter (EnKF)

- EnKF is more dependent on approximate Gaussianity than is sometimes supposed.
- The Gaussian-inspired update rule is similar to the extended Kalman filter (EKF), which has largely been superseded by particle filter methods for low-dimensional nonlinear biological dynamics.
- Simple systems can defeat EnKF: the linear Gaussian update is helpless when data inform the conditional variance rather than the conditional mean.
- Big systems need computationally tractable analysis. EnKF may sometimes be the best solution available, but be aware of its limitations.

# An iterated block particle filter for parameter estimation

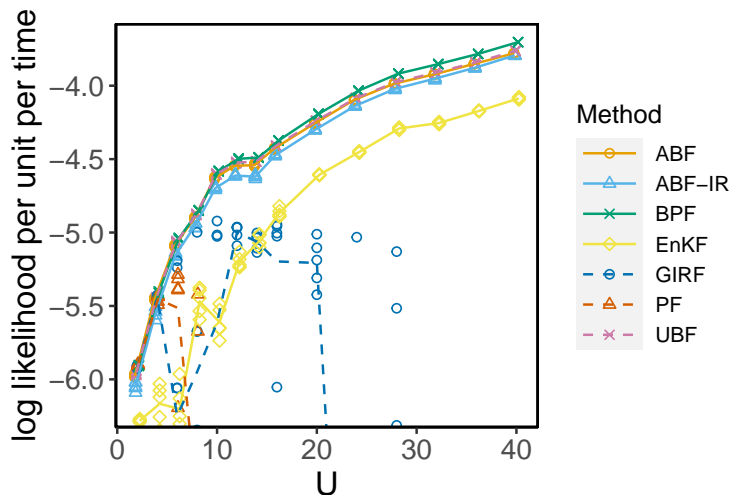


# Practical inference using IBPF

- ① Monte Carlo adjusted profile likelihood (Ionides et al., 2017) obtains confidence intervals that accommodate Monte Carlo error.
- ② Comparing the log-likelihood with an autoregressive model (or other simple statistical model) provides a check of model fit.
- ③ Comparing the block log-likelihood against the benchmark provides insight into problematic units.
- ④ Comparing the conditional log-likelihood for each observation against the benchmark helps to identify outliers.
- ⑤ Two recent case studies (Wheeler et al, 2023; Li et al, 2023) demonstrate data analysis using IBPF. Code and data are provided via R packages extending `spatPomp`.

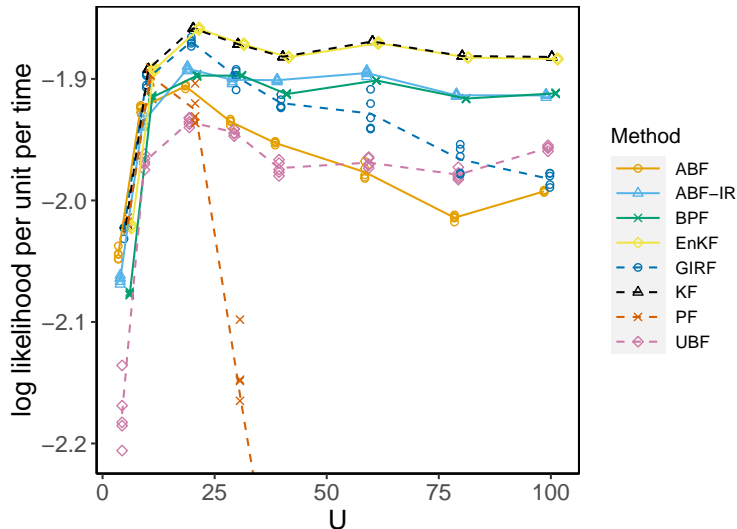


# Filtering $U$ units of a coupled measles SEIR model



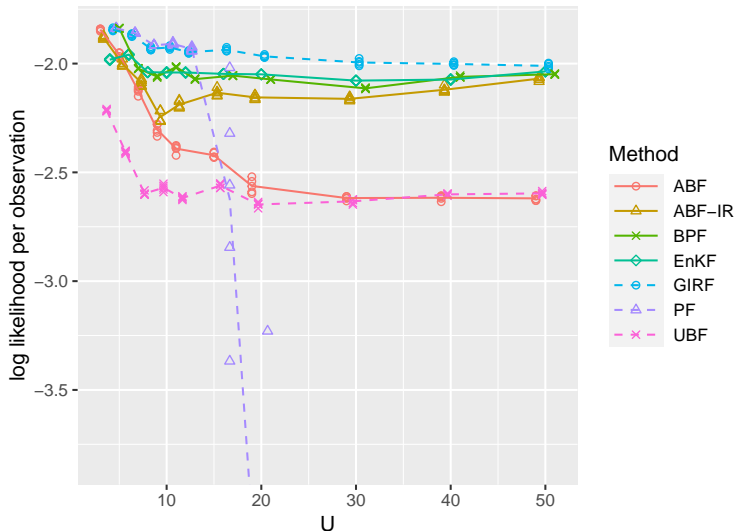
Simulated data using a gravity model with geography, demography and transmission parameters corresponding to UK pre-vaccination measles (Ionides et al, JASA, 2023).

# Filtering $U$ -dimensional correlated Brownian motion



$$\text{Cov}(X_{u,n} - X_{u,n-1}, X_{\tilde{u},n} - X_{\tilde{u},n-1}) \sim 0.4^{|u-\tilde{u}|}$$

# Filtering $U$ units of Lorenz 96 toy atmospheric model



$$dX_u(t) = \{X_{u-1}(t)(X_{u+1}(t) - X_{u-2}(t)) - X_u(t) + F\}dt + \sigma dB_u(t)$$

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