# Bagging and blocking: Inference via particle filters for interacting dynamic systems

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### Outline<sup>1</sup>

**The curse of dimensionality**. Particle filter (PF) methods are effective for inference on low-dimensional nonlinear partially observed stochastic dynamic systems. They scale exponentially badly.

Bagged filters. Combining independent Monte Carlo filters.

- Unadapted bagged filter (UBF)
- Adapted bagged filter (ABF)
- Adapted bagged filter with intermediate resampling (ABF-IR)

**Blocked particle filter** (BPF). Theory by Rebeschini and van Handel (2015). Independently proposed by Ng et al. (2002).

**From filtering to inference**. Iterated filtering using stochastically perturbed parameters.

**Metapopulation dynamics**. Bagged and blocked filters work on collections of weakly coupled populations, in theory and practice.

# What is a SpatPOMP?

**POMP** models are partially observed Markov processes, also known as state space models or hidden Markov models.

**SpatPOMP** models are POMP models with a unit structure.

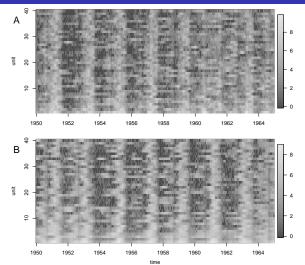
Latent Markov process:  $X_{u,n} = X_u(t_n)$ ,  $u \in 1:U$ ,  $n \in 1:N$ 

**Observation process**:  $Y_{u,n}$  depends only on  $X_{u,n}$ 

The units could be a metapopulation, say cities in an epidemic model.

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### U=40 units for a coupled measles SEIR model



- **A**. Simulated Susceptible-Exposed-Latent-Recovered dynamics coupled with a gravity model.
- B. Measles UK pre-vaccination case reports for the 40 largest cities.

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# Particle filter (PF)



Mutation
↓
Fitness
↓
Natural selection

### Particle filter algorithm

Predict: stochastic dynamics

Measurement: weight

Filter: resample

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

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# Block particle filter (BPF)

### **Evolutionary analogy**

Mutation

↓
Fitness
for each chromosome

↓
Natural selection
for each chromosome

↓

Recombine chromosomes

#### Block particle filter

Predict: stochastic dynamics

Measurement: weight for each block

Filter: resample for each block

Recombine blocks

- Blocks in BPF allow recombination (reassortment of chromosomes in sexual reproduction) in the evolutionary analogy.
- Blocks are a partition of the spatial units.

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### Anticipated limitations of BPF

Rebeschini and van Handel (2015) proved an asymptotic limit where BPF beats the curse of dimensionality but were modest in their applied hopes since blocks small enough to be practical might give unacceptable bias.

- "not anticipated to be applicable to real high-dimensional problems"
- "it is far from clear whether this simple algorithm is of immediate practical utility in the most complex real-world applications"

Thus we look for algorithms without this weakness which also have provable scalability.

# Plug-and-play methods for implicit models

- We address stochastic dynamic models where a simulator is available, but transition densities are not readily accessible.
- These models have been called implicit (Diggle and Gratton, 1984).
- An algorithm that uses a simulator but not transition densities is called plug-and-play (Bretó et al., 2009; He et al., 2010).
- Plug-and-play methods can be applied to implicit models.
- Similar ideas have been called equation-free and likelihood-free.
- BPF is plug-and-play.
- We now consider another scalable simple plug-and-play filter with different strengths and weaknesses to BPF.

### Algorithm 1: Unadapted bagged filter (UBF).

```
input: simulator for f_{X_n|X_{n-1}}(x_n|x_{n-1}) and f_{X_0}(x_0); evaluator for
            f_{Y_{u,n}|X_{u,n}}(y_{u,n} \mid x_{u,n}); data, \boldsymbol{y}_{1:N}^*; number of replicates, \mathcal{I};
            neighborhood structure, B_{u,n}
for i in 1:\mathcal{I} do
     initialize simulation, X_{0,i} \sim f_{X_0}(\cdot)
     for n in 1:N do
           simulate, \boldsymbol{X}_{n,i} \sim f_{\boldsymbol{X}_n | \boldsymbol{X}_{n-1}} (\cdot | \boldsymbol{X}_{n-1,i})
           measurement weights, w_{u,n,i}^M = f_{Y_{u,n}|X_{u,n}}(y_{u,n}^*|X_{u,n,i})
           prediction weights, w_{u,n,i}^P = \prod_{(\tilde{u},\tilde{n}) \in R_{u,n}} w_{\tilde{u},\tilde{n},i}^M
     end
end
```

$$\begin{array}{l} \ell_{u,n}^{\,\mathrm{MC}} = \log \left( \sum_{i=1}^{\mathcal{I}} w_{u,n,i}^{M} w_{u,n,i}^{P} \right) - \log \left( \sum_{i=1}^{\mathcal{I}} w_{u,n,i}^{P} \right) \\ \mathbf{output:} \ \ \log \ \mathrm{likelihood} \ \mathrm{estimate}, \ \ell^{\,\mathrm{MC}} = \sum_{n=1}^{N} \sum_{u=1}^{U} \ell_{u,n}^{\,\mathrm{MC}} \end{array}$$

### Bagged filters

- **Bagging** is bootstrap aggregating. The goal is to gain strength from many boostrap replicates.
- Simulating from a postulated model is a simple parametric bootstrap.
- To obtain scalability, we use local weights to aggregate the bootstrap replicates.
- The unadapted bagged filter is a fancy name for a simple algorithm. We view it a starting point for adapted bagged filters.

# The unadapted bagged filter is not entirely naive

- ullet UBF seems naive. Particle filter (PF) method are well known to scale better with N than unconditional simulations.
- $\bullet$  With modern computers, large numbers of simulations are feasible even when U and N are not small.
- $\bullet$  Initially we studied UBF as a theoretical toy, since it is relatively easy to show theoretically that it can beat the curse of dimensionality as U increases, for weakly coupled systems. Then we found it is competitive in practice on some models of interest.

# Adapted simulation: An easier problem than filtering

- We aim to make each replicate track the data in a weak sense, easier and more scalable than solving the full filtering problem.
- ullet The adapted simulation problem is to draw from  $f_{m{X}_n|m{Y}_n,m{X}_{n-1}}ig(m{x}_n\,|\,m{y^*}_n,m{x}_{n-1}ig).$
- ullet The adapted bagged filter (ABF) algorithm uses importance sampling to carry out adapted simulation on each replicate, with a sample size J.
- Importance sampling for adapted simulation does NOT beat the curse of dimensionality. We combine it with intermediate resampling to give scalability.
- ABF calculates the likelihood using the proper weight restricted to a neighborhood.

#### ABF. Adapted bagged filter.

Initialize adapted simulation:  $m{X}_{0,i}^{ ext{A}} \sim f_{m{X}_0}(m{x}_0)$ 

For n in 1:N

Proposals: 
$$m{X}_{n,i,j}^{\mathrm{P}} \sim f_{m{X}_n | m{X}_{n-1}} ig( m{x}_n \, | \, m{X}_{n-1,i}^{\mathrm{A}} ig)$$

Measurement weights: 
$$w_{u,n,i,j}^M = f_{Y_{u,n}|X_{u,n}} \left(y_{u,n}^* \mid X_{u,n,i,j}^P \right)$$

Adapted resampling weights:  $w_{n,i,j}^{\mathrm{A}} = \prod_{u=1}^{U} w_{u,n,i,j}^{M}$ 

Resampling: 
$$\mathbb{P}\big[r(i)=a\big]=w_{n,i,a}^{\mathrm{A}}\Big(\sum_{k=1}^{J}w_{n,i,k}^{\mathrm{A}}\Big)^{\mathrm{T}}$$

$$\begin{split} \boldsymbol{X}_{n,i}^{\mathrm{A}} &= \boldsymbol{X}_{n,i,r(i)}^{\mathrm{P}} \\ w_{u,n,i,j}^{\mathrm{P}} &= \prod_{\tilde{n}=1}^{n-1} \left[ \frac{1}{J} \sum_{k=1}^{J} \prod_{(\tilde{u},\tilde{n}) \in B_{u,n}^{[\tilde{n}]}} w_{\tilde{u},\tilde{n},i,k}^{M} \right] \prod_{(\tilde{u},n) \in B_{u,n}^{[n]}} w_{\tilde{u},n,i,j}^{M} \end{split}$$

End for

$$\ell_{u,n}^{\text{MC}} = \log \left( \frac{\sum_{i=1}^{\mathcal{I}} \sum_{j=1}^{J} w_{u,n,i,j}^{M} w_{u,n,i,j}^{P}}{\sum_{i=1}^{\mathcal{I}} \sum_{j=1}^{J} w_{u,n,i,j}^{P}} \right)$$

### Intermediate resampling

- ullet Intermediate resampling splits the time interval between observations into S subintervals.
- Reweighting and/or sampling at each subinterval uses a revised estimate
  of the anticipated measurement density at the end of the interval called a
  guide function.
- This is applicable to continuous time models.
- Intermediate resampling has useful theoretical and empirical properties (Del Moral and Murray, 2015; Park and Ionides, 2020).
- Intermediate resampling for adapted simulation within ABF gives the ABF-IR algorithm.
- Intermediate resampling within PF gives the guided intermediate resampling filter (GIRF) of Park and Ionides (2020), a generalization of the auxiliary particle filter of Pitt and Shepard (1999).

### A guide function for intermediate resampling

- Intermediate resampling with an ideal guide function can beat the curse of dimensionality (Park and Ionides, 2020).
- It is consistent for any guide function, but scalability is limited in practice since the ideal guide is generally intractable.
- In practice, we use moment-matching to approximate the ideal guide for Gaussian models.
- Additional algorithmic parameters: number of intermediate timesteps, S measurement variance parameterizations,  $\overset{\leftarrow}{\mathbf{v}}_{u,n}$  and  $\overset{\rightarrow}{\mathbf{v}}_{u,n}$  approximate process and observation mean functions,  $\boldsymbol{\mu}$  and  $h_{u,n}$
- Guided intermediate resampling is plug-and-play: it does not need evaluation of transition densities.

#### ABF-IR. ABF with intermediate resampling.

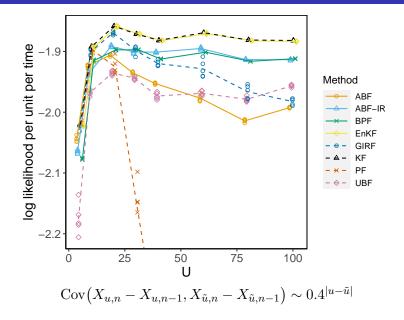
Initialize adapted simulation:  $\boldsymbol{X}_{0,i}^{\mathrm{A}} \sim f_{\boldsymbol{X}_0}(\boldsymbol{x}_0)$ For n in 1:NGuide simulations:  $oldsymbol{X}_{n,i,j}^G \sim f_{oldsymbol{X}_n | oldsymbol{X}_{n-1}} ig( oldsymbol{x}_n \, | \, oldsymbol{X}_{n-1,i}^{\mathrm{A}} ig)$ Guide variance:  $V_{u,n,i} = \operatorname{Var}\{h_{u,n}(X_{u,n,i,j}^G), j \text{ in } 1:J\}$  $g_{n,0,i,j}^{R} = 1$  and  $X_{n,0,i,j}^{IR} = X_{n-1,i}^{A}$ For s in 1:SIntermediate proposals:  $X_{n,s,i,j}^{\text{IP}} \sim f_{X_{n,s}|X_{n,s-1}} \left( \cdot | X_{n,s-1,i,j}^{\text{IR}} \right)$  $\mu_{n,s,i,j}^{\text{IP}} = \mu(X_{n,s,i,j}^{\text{IP}}, t_{n,s}, t_n)$  $egin{align*} oldsymbol{\mu}_{n,s,i,j} &= oldsymbol{\mu}(oldsymbol{A}_{n,s,i,j}, t_{n,s}, \ell_n) \ V_{u,n,s,i,j}^{ ext{proc}} &= \overrightarrow{V}_{u}( heta, \mu_{u,n,s,i,j}^{ ext{pro}}) \ , \end{array} \quad V_{u,n,s,i}^{ ext{proc}} &= V_{u,n,i} \left(t_n - t_{n,s}\right) \Big/ \left(t_n - t_{n,0}\right) \end{aligned}$  $\theta_{u,n,s,i,j} = \stackrel{\leftarrow}{\nabla}_{u} \left( V_{u,n,s,i,j}^{\text{meas}} + V_{u,n,s,i}^{\text{proc}}, \mu_{u,n,s,i,j}^{\text{IP}} \right)$  $\begin{array}{l} g_{n,s,i,j} = \prod_{u=1}^{U} f_{Y_{u,n}|X_{u,n}} \left(y_{u,n}^* \mid \mu_{u,n,s,i,j}^{\mathrm{IP}}; \theta_{u,n,s,i,j}\right) \\ \text{Guide weights: } w_{n,s,i,j}^G = g_{n,s,i,j}/g_{n,s-1,i,j}^{\mathrm{R}} \end{array}$ Resampling:  $\mathbb{P}[r(i,j)=a] = w_{n,s,i,a}^G \left(\sum_{k=1}^J w_{n,s,i,k}^G\right)^{-1}$  $X_{n,s,i,j}^{\text{IR}} = X_{n,s,i,r(i,j)}^{\text{IP}}$  and  $g_{n,s,i,j}^{\text{R}} = g_{n,s,i,r(i,j)}$ End For Set  $X_{n,i}^{A} = X_{n,S,i,1}^{IR}$ Measurement weights:  $w_{u,n,i,i}^M = f_{Y_{u,n}|X_{u,n}}(y_{u,n}^*|X_{u,n,i,i}^G)$  $w_{u,n,i,j}^{\mathrm{P}} = \prod_{\tilde{n}=1}^{n-1} \Big[ \frac{1}{J} \sum_{a=1}^{J} \prod_{\substack{(\tilde{u},\tilde{n}) \in B_{i}^{[\tilde{n}]}, \\ \tilde{n}}} w_{\tilde{u},\tilde{n},i,a}^{M} \Big] \prod_{\substack{(\tilde{u},n) \in B_{u,n}^{[n]}}} w_{\tilde{u},n,i,j}^{M}$ End for

$$\ell_{u,n}^{\text{MC}} = \log \left( \frac{\sum_{i=1}^{\mathcal{I}} \sum_{j=1}^{J} w_{u,n,i,j}^{M} w_{u,n,i,j}^{P}}{\sum_{i=1}^{\mathcal{I}} \sum_{j=1}^{J} w_{u,n,i,j}^{P}} \right)$$

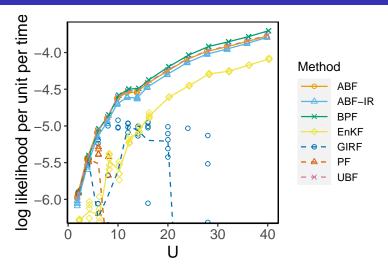
# Software for SpatPOMP models

- We use the asif, asifir, bpfilter, enkf and girf implementations in the R package spatPomp (Asfaw et al., 2019).
- All these algorithms are plug-and-play. This facilitates implementations applicable to a wide class of models: SpatPOMPs that can be simulated.
- spatPomp offers a class 'spatPomp' that extends the 'pomp' class for POMP models in the R package pomp (King et al., 2016).
- All methods available in pomp can formally be applied to 'spatPomp' objects, though they may not be practically effective for spatiotemporal POMPs.

### Filtering *U*-dimensional correlated Brownian motion

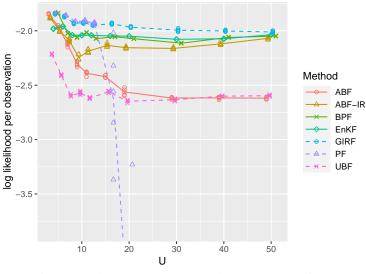


# Filtering U units of a coupled measles SEIR model



Simulated data using a gravity model with geography, demography and transmssion parameters corresponding to UK pre-vaccination measles.

# Filtering U units of Lorenz 96 toy atmospheric model

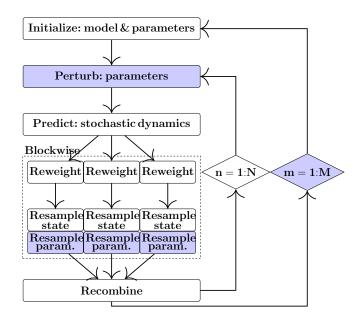


$$dX_u(t) = \{X_{u-1}(t)(X_{u+1}(t) - X_{u-2}(t)) - X_u(t) + F\}dt + \sigma dB_u(t)$$

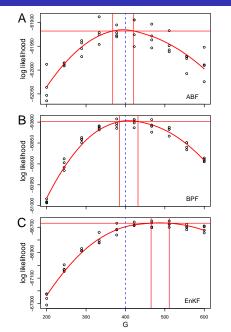
# From filtering to parameter inference

- Log likelihood evaluation in principle enables likelihood-based or Bayesian inference.
- Iterated filtering for PF (Ionides et al., 2015) and GIRF (Park and Ionides, 2020) maximizes the likelihood by randomly perturbing the parameters.
- Particle Markov chain Monte Carlo can be applied with any likelihood estimate (Andrieu et al., 2010). It is numerically intractable when Monte Carlo estimates are costly and noisy.
- Iterated filtering is harder for bagged filters; it is possible but expensive (lonides et al., 2021).
- Iterated filtering works well for BPF when parameters are unit-specific, i.e., each city has its own parameters (Ning and Ionides, 2021). It also can work with shared parameters (current unpublished work).

### An iterated block particle filter for unit-specific parameters



# Measles likelihood slices for coupling parameter, G



Simulating 15 year of data from U=40 cities for the measles model. Slice likelihood, varying G with other paramters fixed at the truth.

- A. Evaluation using ABF.
- **B**. Evaluation using BPF.
- **C**. Evaluation using EnKF.

# Convergence of UBF, ABF & ABF-IR (Ionides et al., 2021)

#### Theorem

Let  $\ell^{\text{MC}}$  denote the Monte Carlo likelihood approximation constructed by UBF, ABF or ABF-IR. Consider a limit with a growing number of replicates,  $\mathcal{I} \to \infty$ . Suppose regularity assumptions listed in the paper. There are quantities  $\epsilon(U,N) = O(1)$  and  $V(U,N) = O(U^2N^2)$  such that

$$\mathcal{I}^{1/2} [\ell^{MC} - \ell - \epsilon UN] \xrightarrow[\tau \to \infty]{d} \mathcal{N}[0, V],$$

where  $\xrightarrow[\mathcal{I} \to \infty]{d}$  denotes convergence in distribution and  $\mathcal{N}[\mu, \Sigma]$  is the normal distribution with mean  $\mu$  and variance  $\Sigma$ . If an additional spatiotemporal mixing assumption holds, we obtain an improved variance bound

$$V(U,N) = O(UN)$$

### Future work

- We are getting close to the point where we can carry out likelihood-based inference for a flexible class of SpatPOMP models for measles. Flexibility supports generation and testing of scientific hypotheses.
- Measles was previously a motivating model system for POMP methods for single populations.
- Many systems in ecology, epidemiology and elsewhere could be studied in a SpatPOMP framework (Bjørnstad and Grenfell, 2001).

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