# Sequential Monte Carlo (SMC) Approaches for Ocean Biology\*: Our Decade of Experimentation

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\* Lower trophic level biology – the planktonic ecosystem.

An Acknowledgement to Weather Forecasting

### Numerical Weather Prediction pioneered large-scale estimation for time dependent systems based on dynamic / numerical models.

**1960s:** Optimal Interpolation (Lev Gandin):

The data assimilation cycle – sequential estimation via approximate
 Kalman filters

**1980s:** Variational Data Assimilation (Olivier Talagrand) - online optimization using numerical weather models as an IVP, needs adjoints need for gradient

**2000s:** Ensemble Kalman filter (Geir Evensen)

- Modular, sample based, and incorporates dynamical model error

#### Clear Performance Metric: Forecast Skill

# Statistical Framework for Biological Data Assimilation: State Space Model

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**Main Goals:** Given observations,  $y_t$ , we estimate:

- $x_t$ : the state (filtering for online prediction, smoothing for retrospective analysis)
- θ's: parameters (for scientific understanding)

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 x<sub>t</sub>: the state (filtering for online prediction) smoothing for retrospective analysis) Often done separately

• θ's: parameters (for scientific understanding)

### Features of Biological Data Assimilation

#### **Dynamics are highly informative:**

 Numerical ocean models considered a good representation of reality. Encapsulate accumulated scientific knowledge
 True for the physics aspects, less so for biology

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#### Data Poor:

- Relies on opportunistic sampling
- A partially observable system

But this is changing .. An observing revolution is well underway

### State Space Model: Dynamics

$$x_{t} = d(x_{t-1}, \theta, e_{t}) \quad \text{or} \quad x_{t} \sim p(x_{t}, \theta \mid x_{t-1}) \quad \longleftarrow \text{DYNAMICS}$$
$$t = 1, \dots, T$$



### Basic Biology: The PZND Model Class



P: phytoplankton
Z: zooplankton
N: nutrients
D: detritus
chl: chlorophyll

Systems of ODEs: (complex dynamics for biology)

$$\frac{dX}{dt} = f(X,\theta) + e(t)$$

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*Role of these simple models?*: Test beds for prototyping methods. But Little utility in mimicking and understanding reality (why? physical control of fluid system)

### Coupling Biology to Physics

**Biology:** 

$$\frac{\partial X_i}{\partial t} + \vec{u} \cdot \nabla X_i - K \nabla^2 X_i = f(x_1, \dots, x_m) + e_i$$

Tracer Equations: Biology in a fluid flow

# Coupling Biology to Physics

Biology:  $\frac{\partial}{\partial t}$ 

$$\frac{\partial X_i}{\partial t} + \vec{u} \cdot \nabla X_i - K \nabla^2 X_i = f(x_1, \dots, x_m) + e_i$$
$$\frac{\partial \vec{u}_h}{\partial t} + \vec{u}_h \cdot \nabla \vec{u}_h + \vec{\Omega} \times \vec{u}_h = \frac{1}{\rho_0} \nabla P + K_h \nabla^2 \vec{u}_h + e_m$$
$$\nabla \cdot \vec{u}_h + \frac{\partial w}{\partial z} = 0; \quad \frac{\partial \rho}{\partial z} = -\rho g; \quad \rho = \rho(T, S, z)$$
$$\frac{\partial T}{\partial t} + \vec{u} \cdot \nabla T - K \nabla^2 T = e_T$$
$$\frac{\partial S}{\partial t} + \vec{u} \cdot \nabla S - K \nabla^2 S = e_S$$

Physics:

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- Numerical integration requires short time steps and fine spatial resolution (= computationally costly).
- Community ocean models available ...





- Vertical: 36 depth levels
- 7 prognostic variables (P,Z,chl,2Ds,N03)
- = state dimension of  $x_t$ : 2,686,320\*
- ~30 biological parameters (many with quite informative prior knowledge)



-500

-1000

\*effective d.o.f. a lot smaller – spatial correlation, and variable inter-dependence

#### A snapshot of model predicted ocean chlorophyll



### Approaches to Stochastic Simulation

... for dynamical models that are essentially formulated as deterministic ones

1.Add Dynamical noise:

2.Random Initial and Boundary Conditions:

3.Use Multi-model Ensembles

4. Stochastic parameters:

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### State Space Model: Observations



### OBSERVATIONS



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*New observations*: complex spatio-temporal multivariable sampling via autonomous robotic sampling platforms



#### **ARGO** Floats





# Keep in Mind ... the General Probabilistic Solution

The Hierarchical Bayesian Model:

$$p(x_{1:T}, \theta \mid y_{1:T}) \propto p(y_{1:T} \mid x_{1:T}, \theta) \cdot p(x_{1:T} \mid \theta) \cdot p(\theta)$$

where:  $x_{1:T} = (x_1, ..., x_T)$  is the system state  $y_{1:T} = (y_1, ..., y_T)$  are the observations  $\theta$  are the dynamical model parameters

- $p(x_{1:T}, \theta | y_{1:T})$  is our target distribution
- $p(y_{1:T} | x_{1:T}, \theta)$  is the conditional measurement distribution
- $p(x_{1:T} \mid \theta)$  is indentified with the numerical ocean model
- $p(\theta)$  is any prior information (expert knowledge)

#### Rely on sampling based solutions in practice $\rightarrow$

A General Computational Solution: Particle MCMC for k = 1 to m

Generate candidate  $\theta^* = \theta^{(k)} + \varepsilon$ 

Run particle filter to determine  $\{x_{t|t}^*\}$  for  $\theta^*$ Evaluate likelihood  $L(\theta \mid y_{1:T}) \propto \prod_{t=1}^T \left(\sum_{k=1}^n p(y_t \mid x_{t|t}^*)\right)$ 

Do Metropolis-Hastings accept/reject step Compute the acceptance probability:  $\alpha = \frac{L(\theta^* | y_{1:T})}{L(\theta^{k-1} | y_{1:T})}$ 

Draw  $u \sim U(0,1)$ 

If  $\min(1,\alpha) \ge u$  then  $\theta^k = \theta^*$ ,

else  $\theta^k = \theta^{(k-1)}$ 

end (for k)

→ yields sample drawn from target  $p(x_{1:T}, \theta | y_{1:T})$ 

Computational Solution: Particle MCMC for k = 1 to m Generate candidate  $\theta^* = \theta^{(k)} + \epsilon$ Run particle filter to determine  $\{x_{t|t}^*\}$  for  $\theta^*$ Evaluate likelihood  $L(\theta \mid y_{1:T}) \propto \prod_{t=1}^{T} \left[ \sum_{i=1}^{n} p(y_t \mid x_{t|t}^*) \right]$ Do Metropolis-Hastings accept/reject step Compute the acceptance probability:  $\alpha = \frac{L(\theta^* | y_{1:T})}{L(\theta^{k-1} | y_{1:T})}$ Draw  $u \sim U(0,1)$ If  $\min(1,\alpha) \ge u$  then  $\theta^k = \theta^*$ , else  $\theta^k = \theta^{(k-1)}$ 

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end (for k)

# State estimation\* via the particle filter (or SMC) is the "engine" for sample based estimation

\* Focus here on estimating state, but can use to estimate parameters via sample based likelihoods or state augmentation

### Particle Filter Schematic: Sequential State Estimation

Single stage transition of system from time *t*-1 to time *t* 



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### Basic Particle Filter:Sequential Importance Resampling

for t = 1 to T

(a) Prediction: generate sample  $\{x_{t|t-1}^{(i)}\}$  following  $p(x_t | y_{1:t-1}, \theta)$  $x_{t|t-1}^{(i)} = d(x_{t-1|t-1}^{(i)}, \theta, e_t^{(i)})$  for i = 1, ..., n

(b) Observation update: Using newly available observation  $y_t$ 

• 
$$w_t^{(i)} \propto p(y_t \mid x_{t|t-1}^{(i)}, \theta)$$
 for  $i = 1, ..., n$ 

- resample with replacement from  $\{x_{t|t-1}^{(i)}\}$  using weights  $w_t^{(i)}$
- $\rightarrow$  yields  $\{x_{t|t}^{(i)}\}$  from  $p(x_t \mid y_{1:t}, \theta)$

end (for t)

Note: there are lots of other (better) particle filtering algorithms!

### Bottleneck for High Dimensional Applications

#### NUMERICAL MODELS ARE COMPUTATIONALLY EXPENSIVE

#### **Practical Consequence:**

*small* ensembles must represent a *large* state space

TWO STRATEGIES:

1. Approximate the **Observation update** 

(so small ensembles work better)

2. Approximate the **Prediction step** 

(so we can generate bigger ensembles)

### Approximating the Observation Update Step

**IDEA**: Use Kalman filter updating for observation step

$$\tilde{x}_{t|t}^{(i)} = x_{t|t-1}^{(i)} + K(y_t^{(i)} - Hx_{t|t-1}^{(i)}), \quad i = 1,...,n$$
  
where:  $y_t^{(i)} = y_t + v_t^{(i)}, \quad i = 1,...,n$  and  $K = PH^T (HPH^T + R)^{-1}$ 

#### The most common approximation for inference in large-scale dynamical systems

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- "Works" for large systems (needs localization, variance inflation).
- Easy to implement (modularity)
- "Breaks" under strong nonlinear, non-Gaussianity.
- Messes up dynamical balances

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CAN ALSO USE AS A PROPOSAL FOR SMC ALGORITHM (lookahead filters / conditional simulation / manifold method)

### Results EnKF: Ensemble Mean

Particulate Organic Nitrogen



time

\* observed state variables: particulate organic N, dissolved inorganic N, chloropyhll, oxygen

### Results EnKF: Ensemble Std Dev

Particulate Organic Nitrogen



\* assimilated state variables: particulate organic N, dissolved inorganic N,, chloropyhll, oxygen

# (2) An Alternative Observation Update: Approximate Bayesian Computation

#### Problems:

(1) Likelihood intractable for some data types,

(2) Difficult for proposals to populate area near posterior mode in high dimensional distribution

**Approach:** Replace likelihood with scalar distance metric in a particle filter. Use this to compute weights

**Benefit:** Allows for use of small sample sizes (no weight collapse)

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#### EXAMPLE

For image comparison, we used *Adaptive Grey Block Distance to* measure discrepancy between *model predicted spatial field* and the *observed one*.



*Issues addressed:* missing values, mis-alignment/registration errors, multiscale dependence

#### Application:

- State estimation using 3-D ocean model. AGBD replaced likelihood in particle filter
- Application fairly successful with small ensembles (<100). Also estimated parameters via state augmentation

# Approximating the Prediction Step

### (1) An Emulator for Parameter Estimation

#### (for Deterministic Dynamics)

GOAL: Estimate biological ocean state in the mid-Atlantic Bight using:
(1) *Data*: Satellite chlorophyll observations,
(2) *Model*: Deterministic 3-D ocean biogeochemical model

*Input*: two key selected 'independent' biological parameters.

**Output:** discrepancy metric, i.e. the AGB distance between model predicted surface field and satellite observations.

Used a Polynomical Chaos emulator **→** 

### Polynomial Chaos Emulator

$$f(x,t,\theta) = \sum_{k=0}^{k_{\max}} a_k(x,t) \phi_k(\theta) + \varepsilon_{trunc}(\theta)$$

where:

 $\theta$  : inputs

 $f(x,t,\theta)$  : outputs

 $a_k(x,t)$  : expansion coefficients

 $\phi_k(\theta)$ : basis functions

 $\varepsilon_{trunc}(\theta)$  : truncation error

#### Note:

- Assumptions about  $p(\theta)$  determine which polynomial basis to use
- The polynomial basis and order determines the *n* design points.
- Mean and Variance of output are given by:

$$E\{f(x,t,\theta)\} = a_0(x,t), \quad \operatorname{var}\{f(x,t,\theta)\} = \sum_{k=1}^n a_k^2(x,t)$$

### Results: Seasonal co-evolution of the 2 parameters



Parameter 1: Phytoplankton Growth

# (2) An Emulator for Particle Filtering (Stochastic Dynamics)

•We want:  $x_t \sim p(x_t | x_{t-1}, \theta)$  - predictive/transition density

•We have:  $x_t = d(x_{t-1}, \theta, e_t)$  - a numerical model to generate samples

Idea: create multivariate distributions for the transition density using copulas ...

$$p(x_t \mid x_{t-1}) = c_K(v_1, \dots, v_K) \prod_{k=1}^K p(x_{t,k} \mid x_{t-1,1:K})$$

- Used elliptical copulas (normal and t) to build the transition density
- Numerical simulations yield CORRELATIONS and MARGINALS used to build the desired distribution →



### Results Sequential MC: Ensemble Mean



time

### Results Sequential MC: Ensemble Std Dev



# Concluding Remarks

- Presented an application-specific tailoring of Sequential Monte Carlo approaches for large scale dynamic systems
- Good approximations are key for workable inference in realistic (high dimension, spatio-temporal) applications for Data Assimilation.
- Ongoing Work: better stochastic simulations (conditional simulation, optimal proposals), and new filtering methods (e.g. a spatial dimension particle filter)

### **Questions/Comments/Concerns?**