# Bayesian Calibration of Simulators with Structured Discretization Uncertainty

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### Probabilistic Numerics

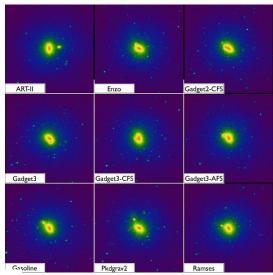
This is an active new field that challenges historical perspectives on numerical analysis.

It is important for this community to develop new methods with an eye to overcoming challenges that lay ahead.

This talk focuses on calibration for forward problems defined by the solution of ordinary and partial differential equations. If you're not already convinced that probabilistic numerics is useful in this setting ...

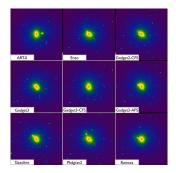


# Example - galaxy simulation (Kim et al., 2016)





# Example - galaxy simulation (Kim et al., 2016)



- These are not realizations of a field (the model is deterministic)
- The initial conditions and inputs are held fixed
- How do we evaluate these numerical solver outputs?

### Perspectives on Probabilistic Numerics

Probability measures on numerical solutions via randomization

- Conrad *et al* (2015/16), Lie *et al* (2017)
- defined outside of the Bayesian framework, but resulting algorithms overlap

Bayesian uncertainty quantification for differential equations

- Skilling (1991), Chkrebtii et al (2013/16), Arnold et al (2013)
- defined outside of the numerical analysis framework, but resulting methods can be analogous in some sense

Bayesian numerical methods

- Hennig & Hauberg (2013/14), Schober et al (2014).
- computationally efficient probabilistic GP based methods; can recover numerical solvers in the mean

## Calibrating stochastic computer models

Regardless of the perspective, the deterministic but unknown forward model is replaced by a stochastic process: for fixed inputs, the output is a random variable with (often) unknown distribution.



Pratola & Chkrebtii (2017+) describe a hierarchical framework to calibrate stochastic simulators with highly structured output uncertainty/variability.

### Calibration problem

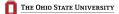
We wish to estimate the unknowns,  $\theta \in \Theta$ , given observations,

$$y(x_t) = A \{ u(x_t, \theta) \} + \varepsilon(x_t), \quad x_t \in \mathcal{X}, \ t = 1, \dots, T,$$

of the deterministic state  $u_t = u(x_t, \theta)$  transformed via an observation process A, and contaminated with stochastic noise  $\varepsilon$ .

The likelihood defines a discrepancy between the model and the data:

$$f(y_{1:T} \mid \theta) \propto \rho \{y_{1:T} - A(u_{1:T})\}$$



### The Bayesian paradigm

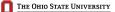
Bayesian inference is concerned with modeling degree of belief about an unknown quantity via probability models.

For example, we may not know  $\theta \in \Theta$  but we may have some prior belief about, e.g., its range, most probable values,

 $\theta \sim \pi(\theta).$ 

We seek to update our prior belief by conditioning on new information,  $y_{1:T} \in \mathcal{Y}$ , e.g., data, model evaluations, via Bayes' Rule:

$$p(\theta \mid y_{1:T}) = \frac{p(y_{1:T} \mid \theta) \pi(\theta)}{\int p(y_{1:T} \mid \theta) \pi(\theta) d\theta} \propto p(y_{1:T} \mid \theta) \pi(\theta).$$



# A Hierarchical model representation

Hierarchical modelling enables inference over the parameters of a stochastic state,

$$\begin{aligned} [y_{1:T} \mid u_{1:T}, \theta] &\propto \rho \{ y_{1:T} - A(u_{1:T}) \} \\ [u_{1:T} \mid \theta] &\sim \rho(u_{1:T} \mid \theta) \\ [\theta] &\sim \pi(\theta). \end{aligned}$$

When  $p(u_{1:T} \mid \theta)$  is not known in closed form, exact inference may still be possible via Monte Carlo, using forward-simulation from the model. However, this is often computationally prohibitive.

## For probabilistic numerics

If the state is deterministic but defined implicitly by a system of differential equations, our uncertainty about the solution can be modelled probabilistically,

$$\begin{array}{lll} [y_{1:\mathcal{T}} \mid u_{1:\mathcal{T}}, \theta] & \propto & \rho \left[ y_{1:\mathcal{T}} - A \left( u_{1:\mathcal{T}} \right) \right] \\ \\ [u_{1:\mathcal{T}} \mid \theta] & \sim & \text{a probability measure representing uncertainty} \\ & \text{ in the solution given discretization of size } N \\ \\ [\theta] & \sim & \pi(\theta). \end{array}$$

We use the Bayesian uncertainty quantification approach to model this middle layer.

### Bayesian UQ for differential equations

Given  $\theta$  and for linear operator D consider the initial value problem,

$$\begin{cases} Du = f(x, u), & x \in \mathcal{X}, \\ u = u_0 & x \in \partial \mathcal{X}. \end{cases}$$

We may have some prior knowledge about smoothness, boundary conditions, etc., described by a prior measure,

$$u \sim \pi, \quad x \in \mathcal{X}$$

We seek to update our prior knowledge by conditioning on model interrogations,  $f_{1:N}$  via Bayes' Rule,

$$p(u(x) \mid f_{1:N}) = \frac{p(f_{1:N} \mid u(x)) \pi(u(x))}{\int p(f_{1:N} \mid u(x)) \pi(u(x)) du(x)} \propto p(f_{1:N} \mid u(x)) \pi(u(x))$$

### Prior uncertainty in the unknown solution

The exact solution function u is deterministic, but unknown. We may describe our prior uncertainty via a probability model defined on the space of suitably smooth derivatives, e.g.,

$$u \sim \mathcal{GP}(m^0, C^0), \qquad m^0: \mathcal{X} \to \mathbb{R}, \ C^0: \mathcal{X} imes \mathcal{X} \to \mathbb{R}$$

with the constraint  $m^0 = u^0, x \in \partial X$ .

This yields a joint prior on the fixed but unknown state and its derivative(s)

$$\begin{bmatrix} u \\ Du \end{bmatrix} \sim \mathcal{GP}\left( \begin{bmatrix} m^0 \\ Dm^0 \end{bmatrix}, \begin{bmatrix} C^0 & C^0 D^* \\ DC^0 & DC^0 D^* \end{bmatrix} \right)$$



### Interrogating the model recursively

1 Draw a sample from the marginal predictive distribution on the state at the next discretization grid point  $s_{n+1} \in \mathcal{X}$ ,  $1 \le n < N$ 

$$u(s_{n+1}) \sim p(u(s_{n+1}) \mid f_{1:n})$$

**2** Evaluate the RHS at  $u(s_{n+1})$  to obtain a model interrogation,

$$f_{n+1} = f(s_{n+1}, u(s_{n+1}))$$

**3** Model interrogations as "noisy" measurements of *Du*:

$$f_{n+1} \mid Du, f_{1:n} \sim \mathcal{N}(Du(s_{n+1}), \Lambda(s_n))$$

### Sequential Bayesian updating

Updating our knowledge about the true but unknown solution given the new interrogation trajectory  $f_{n+1}$ 

$$\begin{bmatrix} u \\ Du \end{bmatrix} f_{n+1} \sim \mathcal{GP}\left( \begin{bmatrix} m^{n+1} \\ Dm^{n+1} \end{bmatrix}, \begin{bmatrix} C^{n+1} & C^{n+1}D^* \\ DC^{n+1} & DC^{n+1}D^* \end{bmatrix} \right)$$

where,

$$m^{n+1} = m^{n} + K^{n} (f_{n+1} - m^{n}(s_{n+1}))$$
  

$$C^{n+1} = C^{n} - K^{n} D C^{n*}$$
  

$$K^{n} = C^{n} D^{*} (D C^{n} + \Lambda(s_{n}))^{-1}$$

This becomes the prior for the next update.

## Bayesian UQ for differential equations

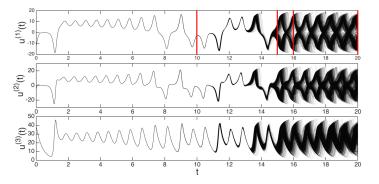
Due to the Markov property, we cannot condition the solution on multiple trajectories  $f_{1:N}^{j}$ ,  $j = 1, \ldots, J$  simultaneously. In fact, the posterior over the unknown solution turns out to be a continuous mixture of Gaussian processes,

$$[u \mid \theta, N] = \int \int [u, Du \mid \mathsf{f}_{1:N}, \theta, N] \ d(Du) \ d\mathsf{f}_{1:N}.$$

Samples from this posterior can be obtained via Monte Carlo.

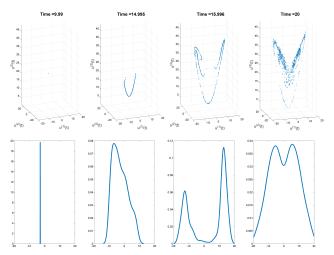
#### Example - Lorenz63 forward model

A probability statement over probable trajectories given fixed model parameters and initial conditions for the Lorenz63 model:



1000 draws for the probabilistic forward model for the Lorenz63 system given fixed initial states and model parameters in the chaotic regime.

#### Example - Lorenz63 forward model



1000 draws from forward model for Lorenz63 system at four fixed time points.

## For probabilistic numerics

If the state is deterministic but defined implicitly by a system of differential equations, our uncertainty about the solution can be modelled probabilistically,

$$\begin{array}{lll} [y_{1:\mathcal{T}} \mid u_{1:\mathcal{T}}, \theta] & \propto & \rho \left[ y_{1:\mathcal{T}} - A \left( u_{1:\mathcal{T}} \right) \right] \\ \\ [u_{1:\mathcal{T}} \mid \theta] & \sim & \text{a probability measure representing uncertainty} \\ & \text{ in the solution given discretization of size } N \\ \\ [\theta] & \sim & \pi(\theta). \end{array}$$

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### Computer model emulation

We interrogate the model at M regimes,  $\theta_{1:M} = (\theta_1, \dots, \theta_M)^{\top}$ . Each interrogation is comprised of an ensemble of K output samples.

Let  $\tilde{u}_{1:T}^{1:K}(\theta_{1:M})$  denote the ensemble of K output simulations for each of the regimes  $\theta_{1:M}$ .

$$\begin{split} \left[ y_{1:T} \mid \tilde{u}_{1:T}^{1:K}(\theta_{1:M}), u_{1:T}, \delta_{1:T}, \theta \right] &\propto \rho \left[ y_{1:T} - A \left( u_{1:T}(\theta) \right) - \delta_{1:T} \right] \\ \left[ \tilde{u}_{1:T}^{k}(\theta_{m}) \mid u_{1:T}^{k}, \theta \right] &\sim \mathcal{N} \left( u_{1:T}^{k}(\theta_{m}), \Lambda \right) \quad k = 1, \dots, K \\ \left[ u_{1:T}^{k} \mid \theta \right] &\sim \text{generative stochastic model} \\ \left[ \theta, \delta \right] &\sim \pi(\theta, \delta). \end{split}$$

# A Hierarchical model representation

Challenges of MCMC sampling from the posterior include:

- Emulation based on *MK* model evaluations is computationally expensive
- Models are evaluated at multiple spatio-temporal locations and over multiple states

Our approach:

- Dimension reduction over the second (output) layer of the hierarchical model
- We include the dimension reduction specifications within the hierarchical model, resulting in a fully probabilistic approach

#### Example: Exact vs Emulation Based Inference in a Model of Protein Dynamics

#### Inference for a model of protein dynamics

JAK-STAT chemical signaling pathway model describes concentration of 4 STAT factors by a delay differential equation system on  $t \in [0, 60]$ ,

Contract gene

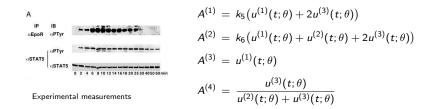
STAT-5

$$\begin{aligned} \frac{d}{dt} u^{(1)}(t,\theta) &= -k_1 u^{(1)}(t,\theta) \operatorname{EpoR}_A(t) + 2 \, k_4 \, u^{(4)}(t-\tau,\theta) \\ \frac{d}{dt} u^{(2)}(t,\theta) &= k_1 \, u^{(1)}(t,\theta) \operatorname{EpoR}_A(t) - k_2 \, \left( u^{(2)}(t,\theta) \right)^2 \\ \frac{d}{dt} u^{(3)}(t,\theta) &= -k_3 \, u^{(3)}(t,\theta) + 0.5 \, k_2 \, \left( u^{(2)}(t,\theta) \right)^2 \\ \frac{d}{dt} u^{(4)}(t,\theta) &= k_3 \, u^{(3)}(t,\theta) - k_4 \, u^{(4)}(t-\tau,\theta) \\ u^{(i)}(t,\theta) &= \phi^{(i)}(t), \qquad t \in [-\tau,0], \ i = 1, \dots, 4 \end{aligned}$$

Illustration of the JAK-STAT mechanism

#### Inference for a model of protein dynamics

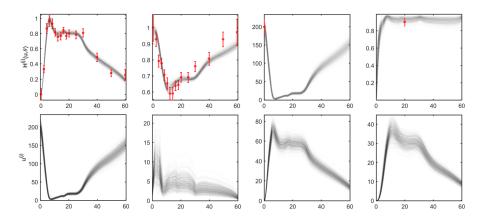
States are observed indirectly through a nonlinear transformation:



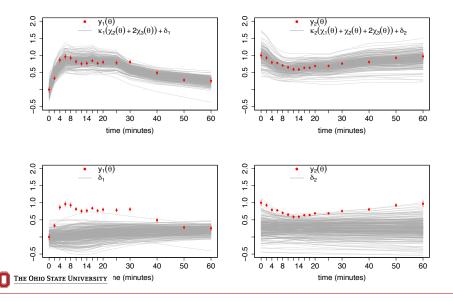
Observations are noisy measurements on the transformed states and forcing function at points  $\mathbf{t} = \{t_{ij}\}_{i=1,...,4;j=1,...,n_i}$ 

$$y(\mathbf{t}) = A_{k_4,k_5}u(\mathbf{t}; k_1, \dots, k_6, \tau, \phi, EpoR_A) + \varepsilon(\mathbf{t})$$

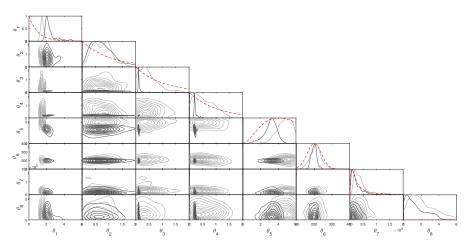
#### Results: exact inference



#### Results: emulation based inference



### Results: emulation based inference



Kernel density estimates of the marginal stochastically calibrated posterior (gray) with M = 100 model runs, and exact posterior (black) for the JAK-STAT system. Marginal prior densities are shown as dotted lines.

Thank you!



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