Optimal Parameter Upscaling for Partial Differential Equation Models in Mathematical Biology

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Introduction

- 2 Related Work
- Mathematics of the Method
- 4 Numerical Results
- **5** Work in Progress

Let

$$u = \mathcal{G}(x; \theta)$$

and consider the problem of finding θ , an input to a mathematical model, given u an observation of solution to the model at point x.



• Consider the most basic coupled epidemic model

$$\begin{aligned} \frac{ds(t)}{dt} &= -\alpha \, s(t) \, i(t), \qquad s(0) = s_0 \\ \frac{di(t)}{dt} &= -\alpha \, s(t) \, i(t) - \beta \, i(t), \qquad i(0) = i_0 \\ \frac{dr(t)}{dt} &= -\beta \, i(t), \qquad r(0) = r_0 \end{aligned}$$

where

- s = susceptible population density,
- i = infected population density,
- r = recovered population density,
- $\theta = (\alpha, \beta)$ is a random field that may depend on t, i, r and s.

- We are interested in estimating the model parameters given some realizations {u_i}^m_{i=1}.
- There are uncertainties in the governing equation due to incomplete knowledge of the underlying physics and/or inevitable errors in measurements.
- These uncertainties are encapsulated in the model parameters α , β and initial and boundary conditions, for some problems

Finite element requires high resolution to capture stochastic information



Naive Approach

- First, we explore the forward model.
- A naive approach is to convert the problem into a deterministic one by replacing θ with a statistic and solving analytically or numerically using finite difference, finite element or finite volume method. Simple example:

$$\frac{du(t)}{dt} = -\theta u(t); \qquad u(0) = u_0$$



Naive solution: $u(t) = u_0 e^{-\bar{\theta}t}$

Stochastic solution: $u(t) = u_0 \int_{\Omega} e^{-\theta t} d\mu(\theta)$

• Perturbation and homogenization methods have also been considered.

Monte Carlo Sampling (MCS) Methods

- The discovery of MCS algorithms was a huge breakthrough in better explaining the effects of uncertainties in models.
- Metropolis-Hastings update step for the inverse case:

$$\theta^{(k+1)} = \begin{cases} \theta^* & \text{with probability} & \min\left\{1, \frac{\mathcal{L}(u|\theta^*)\pi(\theta^*)}{\mathcal{L}(u|\theta^k)\pi(\theta^k)}\right\}\\ \theta^k & \text{otherwise} \end{cases}$$

assuming a symmetric proposal.

- MCMC sampling has a very slow convergence rate, it is mostly not suitable for practical purposes.
- MCMC samples are often correlated

Karhunen-Loeve Expansions I

- A very successful approach also considered stem from the Kosambi–Karhunen–Loève theorem
- For a random field θ(t), let μ_θ(t) be the mean of the process and let C(t, s) = cov(θ_t, θ_s) be its covariance function. The Karhunen-Loève expansion of θ_t is

$$heta_t(\omega) = \mu_ heta(t) + \sum_{i=1}^\infty \sqrt{\lambda_i} \phi_i(t) heta_i(\omega)$$

where ϕ_i 's are the orthogonal eigenfunctions and λ_i 's are the corresponding eigenvalues of the eigenvalue problem

$$\int_{\mathcal{T}} C(t,s)\phi_i(s)ds = \lambda_i\phi_i(t), \qquad t \in \mathcal{T}$$

and

$$heta_i(\omega) = rac{1}{\sqrt{\lambda}}\int_{\mathcal{T}}(heta_t(\omega)-\mu_ heta(t))\phi_i(t)dt$$

are mutually uncorrelated random variables with zero mean and unit variance.

- KKL modes are the principal components of the covariance kernel and are expensive to compute
- KKL requires prior knowledge of the covariance kernel C(t, s) and the underlying distribution.

- Naive approach does not carry forward any stochastic information
- Monte Carlo Simulation is very slow
- Karhunen-Loeve expansions require prior knowledge of the underlying distribution, which is not known for many problems.

What then is a way forward?

Polynomial Chaos

 Generalized polynomial chaos, in many cases, is arguably the only feasible method for stochastic simulations of complex systems [Xiu, 2010] [1]

Theorem (Cameron & Martin, 1947)

Let $L^2(\Omega, \mathcal{A}, P)$ be a Hilbert space of real-valued random variables and $\mathfrak{D} \subset L^2(\Omega, \mathcal{A}, P)$, a complete subspace. Suppose for any $n \in \mathbb{N}$, $\mathcal{B}_n(\mathfrak{D}) = \{f(\xi_1, \cdots, \xi_m) : f \text{ an } m\text{-variate polynomial of degree} \leq m$, $\xi_i \in \mathfrak{D}, i \in [m], m \in \mathbb{N}\}$ and $\{\overline{\mathcal{B}}_n\}_{n \in \mathbb{N}} \subset L^2(\Omega, \mathcal{A}, P)$ is a strictly increasing complete subspace and Gaussian for n = 1. Then, there exists $\{\mathfrak{D}_n\}_{n \in \mathbb{N}}$ such that

$$\bigoplus_{n=0}^{\infty} \mathfrak{D}_n = L^2(\Omega, \sigma(\mathfrak{D}), P)$$

Specifically, for $\sigma(\mathfrak{D}) = \mathcal{A}$, $\bigoplus_{n=0}^{\infty} \mathfrak{D}_n = L^2(\Omega, \mathcal{A}, P)$

Generalized Polynomial Chaos

- The work of [Xiu & Karniadakis, 2002] is pivotal in understanding, applying and generalizing polynomial chaos method. They coined the term *generalized polynomial chaos expansion*
- Let {ξ_n}_{n∈ℕ} be a sequence of (not necessarily identically distributed) basic random variables satisfying conditions

•
$$\mathbb{E}[|\xi_n|^m] < \infty$$
 for all $n, m \in \mathbb{N}$

② $F_{\xi_n}(x) := P(\xi_n \le x)$ is continuous for each *n* ∈ \mathbb{N}

Definition

A distribution is said to be *determinate*, in the Hamburger sense (aka *solvability of the moment problem* if the distribution function is *uniquely* defined by the sequence of its moments

$$\mathbb{E}[\xi^m] = \int_{\mathbb{R}} x^m dF_{\xi}(x)$$

With conditions (1) and (2) and the preceding definition, it follows that:

Theorem

The sequence of orthogonal polynomials associated with a real random variable ξ satisfying the two conditions above is dense in $L^2(\mathbb{R}, \mathfrak{B}(\mathbb{R}), dF_{\xi})$ if and only if the moment problem is uniquely solvable for its distribution.

Following this development and other equivalent formulations, [Ernst etal, 2012] proved conditions under which a random vector admits generalized polynomial chaos expansion

Theorem

If the distribution function F_{ξ} of a random vector $\xi = (\xi_1, \dots, \xi_n)$ with continuous distribution and finite moments of all orders satisfies one of the following conditions, then the multivariate polynomials in ξ_1, \dots, ξ_n are dense in $L^2(\Omega, \sigma(\xi), P)$. In this case any random variable $\zeta \in L^2(\Omega, \sigma(\xi), P)$ is the limit of its generalized polynomial chaos expansion, which converges in quadratic mean.

- The distributiion function F_{ξ} has compact support, i.e., there exists a compact set $S \subset \mathbb{R}^n$ such that $P(\xi \in S) = 1$
- The random vector is exponentially integrable, i.e., there exists c > 0 such that

 $\mathbb{E}(e^{c\|\xi\|}) < \infty$

Courtesy: [Ernst etal, 2010]

Distribution	polynomials	density
Gaussian	Hermite	$ \rho(\xi) = \frac{1}{\sqrt{2\pi}} e^{-\xi^2/2} $
$Gamma(\alpha,\lambda)$	Laguerre	$ \rho(\xi) = \frac{\lambda}{\xi(\alpha)} (\lambda \xi)^{\alpha - 1} e^{-\lambda \xi} $
$Beta(\alpha,\beta)$	Jacobi	$\rho(\xi) = \frac{(1-\xi)^{\alpha}(1+\xi)^{\beta}}{2^{\alpha+\beta+1}B(\alpha+1,\beta+1)}$
$Uniform(\alpha,\beta)$	Legendre	$ \rho(\xi) = \frac{1}{\beta - \alpha} $
Arcsin	Chebyshev	$\rho(\xi) = \frac{1}{\sqrt{1-\xi^2}}$

Transport Maps

- Polynomial chaos expansion method is not without faults
- Its computational complexity increases fast with increase in the number of parameters θ [Moselhy & Marzouk, 2012]
- Transport maps were introduced to mitigate the computational burden involved modeling with polynomial chaos
- A transport map, $T : \mathbb{R}^n \to \mathbb{R}^n$ is a deterministic transformation that pushes forward μ to ν , yielding [Parno & Marzouk, 2012]

$$\nu(B) = \mu(T^{-1}(B))$$

• Uniqueness of *T* is ensured by assuming the *triangular formulation*

$$T(x_1, x_2, \dots, x_d) = \begin{bmatrix} T_1(x_1) \\ T_2(x_1, x_2) \\ \vdots \\ T_d(x_1, x_2, \dots, x_d) \end{bmatrix}$$

First, we consider the case where $\boldsymbol{\theta}$ is a scalar random variable

$$\theta \sim Uniform(0,1)$$

$$\theta \sim exp(0.5)$$





Numerical Experiments

Consider the exponential decay model given by the differential equation

$$rac{du(t; heta)}{dt} = - heta u(t, heta); \qquad u_0 = c$$

where $\theta > 0$ is a random variable.

C



- Most, if not all biology inspired mathematical models depend on certain random parameter(s)
- Successes in making inference from or validating these models depend on how well the stochastic information from these parameters are propagated into the state variables
- We demonstrated that transport maps are powerful and handy in this regard
- In progress, we are looking to leverage the expressive power of Deep Neural Networks in constructing transport maps

[1] Dongbin Xiu. *Numerical methods for stochastic computations*. Princeton university press, 2010.