

# *Predictability in Stochastic Reaction Networks*

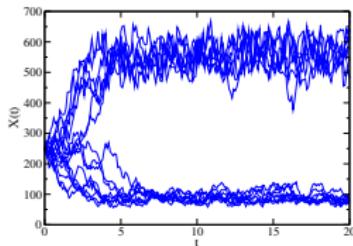
Khachik Sargsyan, Bert Debusschere, Habib Najm

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Livermore, CA

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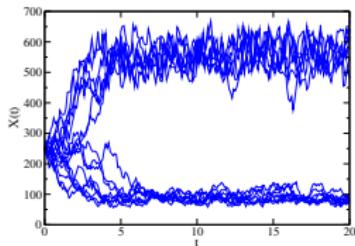
# Stochastic Reaction Networks

- Reaction networks involving small number of molecules necessitate the use of *stochastic* modeling instead of the *deterministic* one. E.g.
  - Microbial processes  
(bioenergy, bioremediation)
  - Surface catalytic reactions  
(fuel cells, batteries)
  - Immune system signaling reactions
- SRNs are modeled as Jump Markov Processes
  - Governed by Chemical Master Equation
$$\dot{P}(X(t) = n) = \sum_m A_{nm}P(X(t) = n)$$
  - Reduces to deterministic Rate Equations in the large volume limit
  - Trajectories simulated by Gillespie's Stochastic Simulation Algorithm (SSA, Gillespie, 1977)



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# Objective: predictability in high-d

$$X(t, \theta, \lambda)$$

- Develop tools for *predictability*( $\lambda$ ) and *dynamical analysis*( $t$ ) of SRNs accounting for
  - Inherent stochasticity ( $\theta$ )
  - Model/parameter uncertainty ( $\lambda$ )
  - Limited data

$$\mathcal{D} = \{X_i\}_{i=1}^N$$

- Predictability assessment
  - Fix  $t$ , focus on  $\lambda$  dependence
  - Statistical properties  $Y(\lambda) = \langle f(X(\theta, \lambda)) \rangle$  have sampling noise
  - How uncertainty in  $\lambda$  affects uncertainty in  $Y(\lambda)$  given limited data
- High dimensionality of  $\lambda$

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# High-dimensional parametric uncertainty in stochastic systems

- Statistical property  $Y(\lambda) = \langle f(X(\theta, \lambda)) \rangle$  of interest.
  - High-dimensional parametric uncertainty ( $\lambda$ )
  - Sampling noise due to limited data  $\{X_i\}$
- Expectation  $\langle \cdot \rangle$  filters intrinsic noise.
  - Averaging over sample realizations of  $X$
  - Still leftover noise, width  $\sim 1/\sqrt{N}$
- Polynomial Chaos expansion to represent input-output relationship
  - Sensitivity analysis
  - Surrogate model for optimization or inverse problems
  - Identify key reaction mechanisms

# Polynomial Chaos expansion represents a random variable as a polynomial of a standard random variable

- Truncated PCE: finite dimension  $n$  and order  $p$

$$Y \simeq \sum_{k=0}^P c_k \Psi_k(\boldsymbol{\eta})$$

with the number of terms  $P + 1 = \frac{(n+p)!}{n!p!}$ .

- $\boldsymbol{\eta} = (\eta_1, \dots, \eta_n)$  standard i.i.d. r.v.  
 $\Psi_k$  standard orthogonal polynomials  
 $c_k$  spectral modes.
- Most common standard Polynomial-Variable pairs:  
(continuous) Gauss-Hermite, Legendre-Uniform,  
(discrete) Poisson-Charlier.

[Wiener, 1938; Ghanem & Spanos, 1991; Xiu & Karniadakis, 2002]

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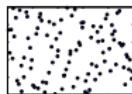
# Alternative methods to obtain PC coefficients

$$Y \simeq \sum_{k=0}^P c_k \Psi_k(\boldsymbol{\eta}) \quad c_k = \frac{\langle Y(\boldsymbol{\eta}) \Psi_k(\boldsymbol{\eta}) \rangle}{\langle \Psi_k^2(\boldsymbol{\eta}) \rangle}$$

The integral  $\langle Y(\boldsymbol{\eta}) \Psi_k(\boldsymbol{\eta}) \rangle = \int Y(\boldsymbol{\eta}) \Psi_k(\boldsymbol{\eta}) \pi(\boldsymbol{\eta}) d\boldsymbol{\eta}$  can be estimated by

- Monte-Carlo

$$\frac{1}{K} \sum_{j=1}^K Y(\boldsymbol{\eta}_j) \Psi_k(\boldsymbol{\eta}_j)$$



many samples from  $\pi(\boldsymbol{\eta})$

- Quadrature

$$\sum_{j=1}^Q Y(\boldsymbol{\eta}_j) \Psi_k(\boldsymbol{\eta}_j) w_j$$

samples at quadrature

- Bayesian inference

$$P(c_k | Y(\boldsymbol{\eta}_j)) \propto P(Y(\boldsymbol{\eta}_j) | c_k) P(c_k)$$

any (number of) samples

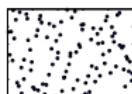
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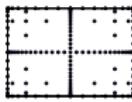
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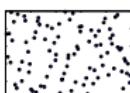
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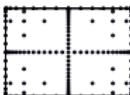
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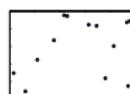
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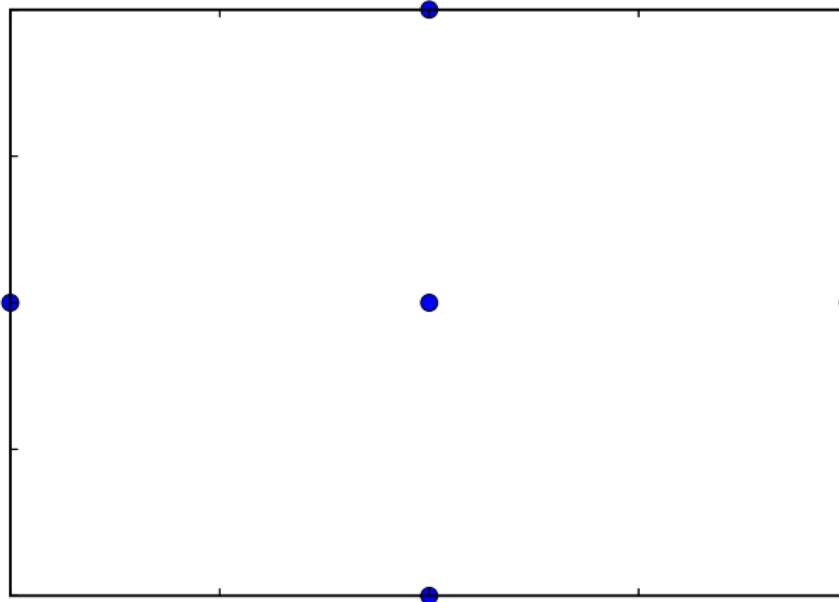
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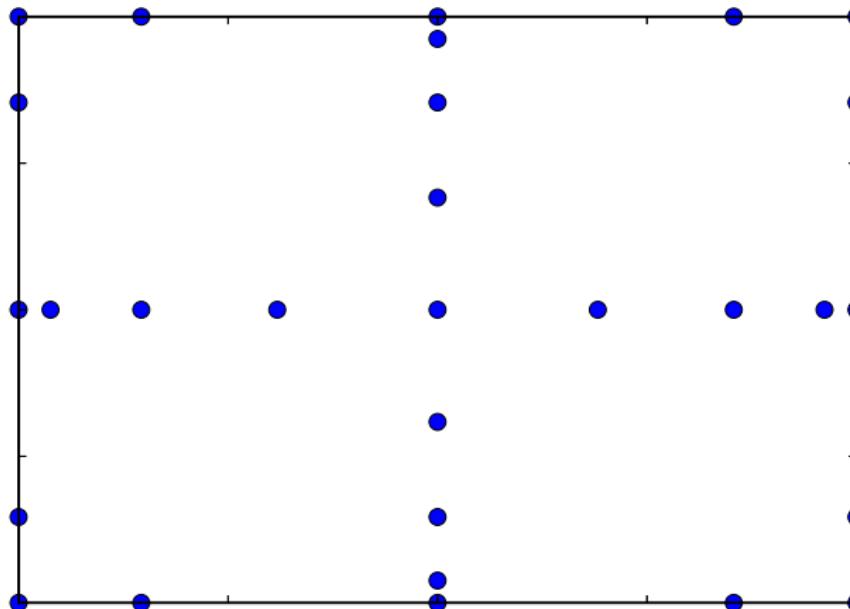
# Sparse quadrature integration well-suited for high-dimensional *smooth* integrands

Clenshaw-Curtis sparse grid, Level = 1



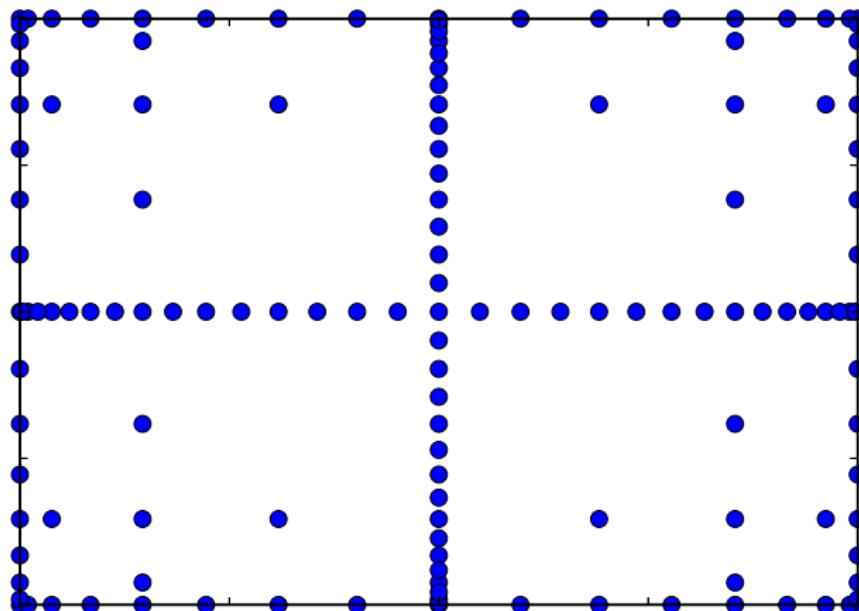
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Clenshaw-Curtis sparse grid, Level = 3



# Sparse quadrature integration well-suited for high-dimensional *smooth* integrands

Clenshaw-Curtis sparse grid, Level = 5



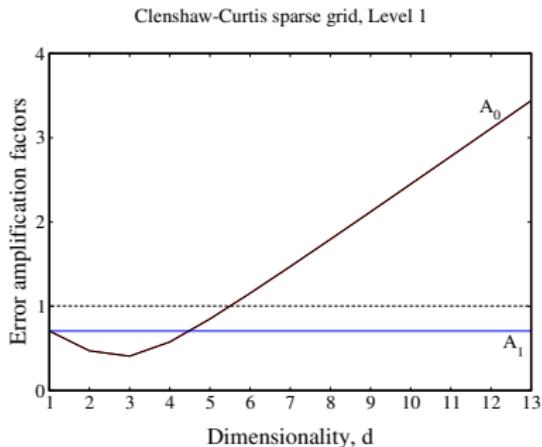
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$$c_k \approx \frac{1}{\langle \Psi_k^2(\boldsymbol{\eta}) \rangle} \sum_{j=1}^Q Y(\boldsymbol{\eta}_j) \Psi_k(\boldsymbol{\eta}_j) w_j$$

Noise  $Y \sim \sigma \implies$  Error  $c_k \sim A_k \sigma$



- amplification factor  $A_k$  grows with dimensionality
  - CC, level 1:  $A_0 = \frac{1}{3} \sqrt{(d-3)^2 + \frac{d}{2}}$ ,  $A_1 = \frac{1}{\sqrt{2}}$ .
- blame the negative weights.
- for full quadrature,  $\frac{1}{n^{d/2}} \leq A_0 \leq 1$ , no amplification!

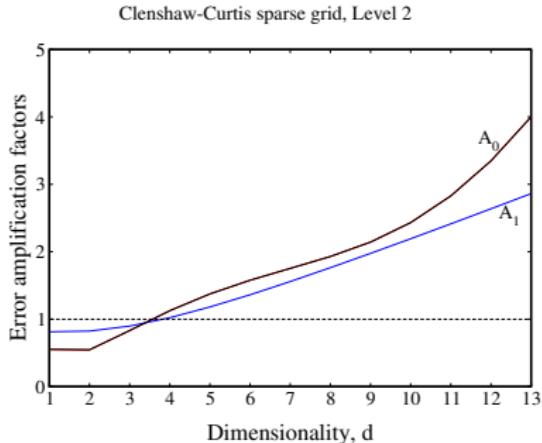
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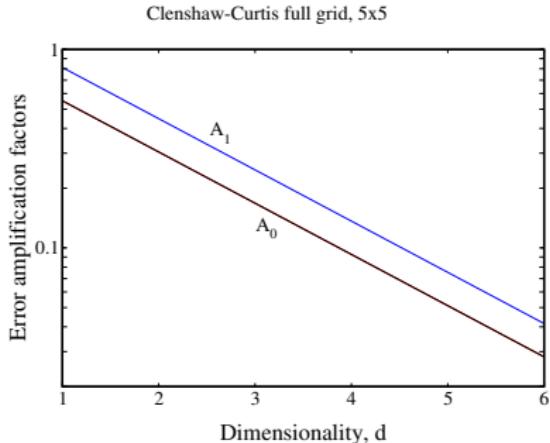
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## Bayesian inference handles the intrinsic stochasticity well

$$Y = \langle X \rangle \simeq \sum_{k=0}^P c_k \Psi_k(\boldsymbol{\eta})$$

$$\overbrace{P(\mathbf{c}|\mathcal{D})}^{\text{Posterior}} \propto \overbrace{P(\mathcal{D}|\mathbf{c})}^{\text{Likelihood}} \overbrace{P(\mathbf{c})}^{\text{Prior}}$$

$$L(\mathbf{c}) = P(\mathcal{D}|\mathbf{c}) = \left( \frac{1}{\sigma \sqrt{2\pi}} \right)^N \prod_{i=1}^N \exp \left( -\frac{(X_i - \sum_{k=0}^P c_k \Psi_k(\boldsymbol{\eta}_i))^2}{2\sigma^2} \right).$$

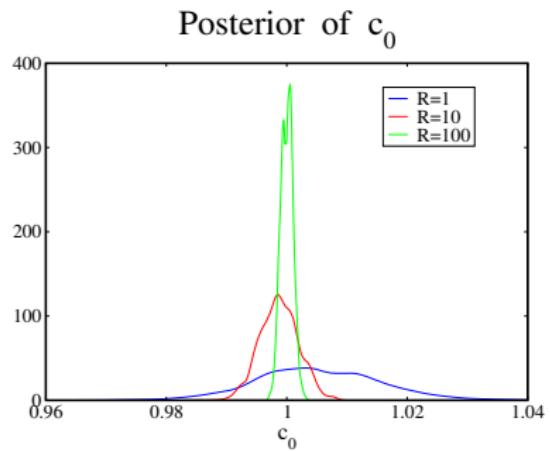
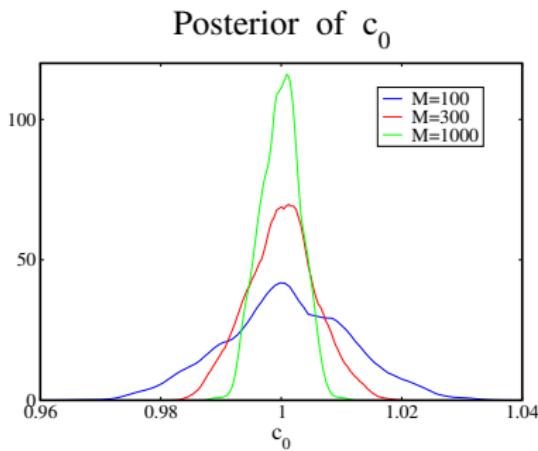
- Noise model is assumed gaussian with  $\sigma$  from CLT or inferred
- Uniformly distributed priors
- Posterior exploration using Markov Chain Monte Carlo (MCMC)
- The whole posterior distribution is accessible,  
*i.e.* uncertain response surface
- Input parameters can have arbitrary values

# Posterior narrows around the true value as more samples are taken

- M parameter locations
- R replicas per parameter
- Second order Legendre polynomial expansion with unit coefficients.

No noise in function evaluations,  $R = 1$

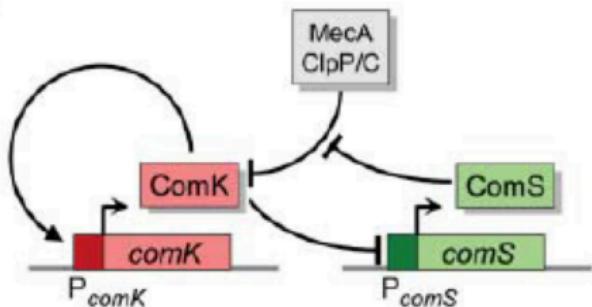
Noisy function evaluations,  $M = 100$



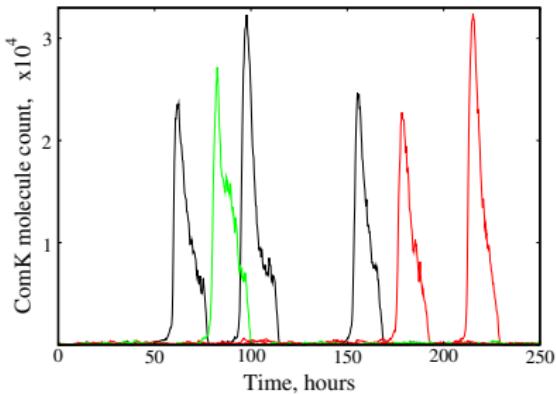
# *Bacillus subtilis* is a soil bacterium

relevant to bioenergy and bioremediation

- 16 reactions, 11 species
- Competence in *B. Subtilis* allows uptake of external DNA
- Rapid rise in transcription factor comK molecules
- Vegetative → Competent state transition is driven by stochasticity
- Input parameters: rate constants of underlying reactions (high-d)
- Output observable: probability of competence  $P_c = P(X_\infty > 5000)$



Süel et al., Science, 2007



# Intrinsic stochasticity induces transition to competence

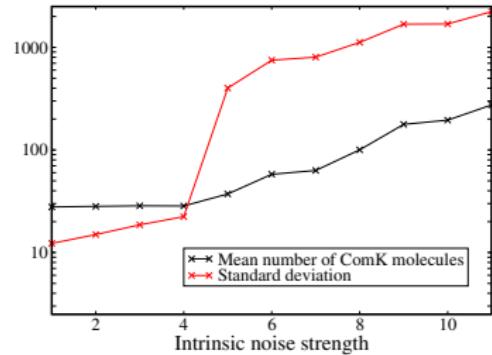
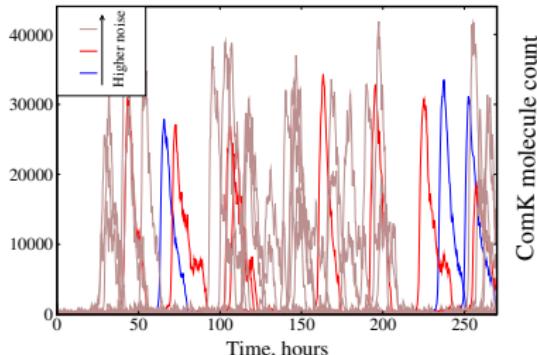
Chemical Master Equation (CME):

$$\frac{d}{dt}P(X(t) = n) = \sum_m A_{nm}P(X(t) = n)$$

Rate equation (ODE):

$$\frac{d}{dt}X(t) = \tilde{A}(X)$$

- No-noise or large volume limit (ODE) does not produce competence
- Many parameter combinations lead to the same ODE limit, but correspond to different effective volumes, i.e. intrinsic noise strength
- Increasing intrinsic noise leads to more frequent transitions



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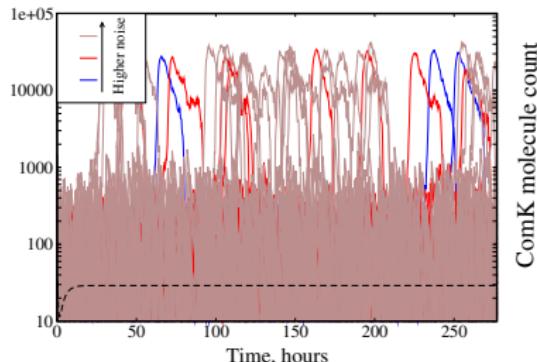
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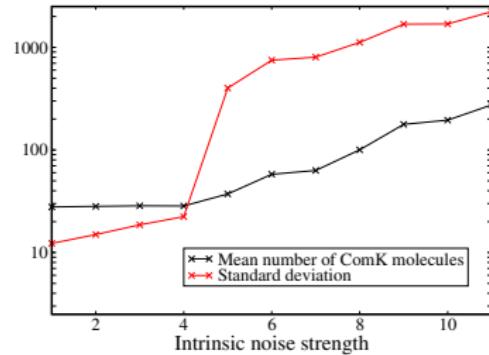
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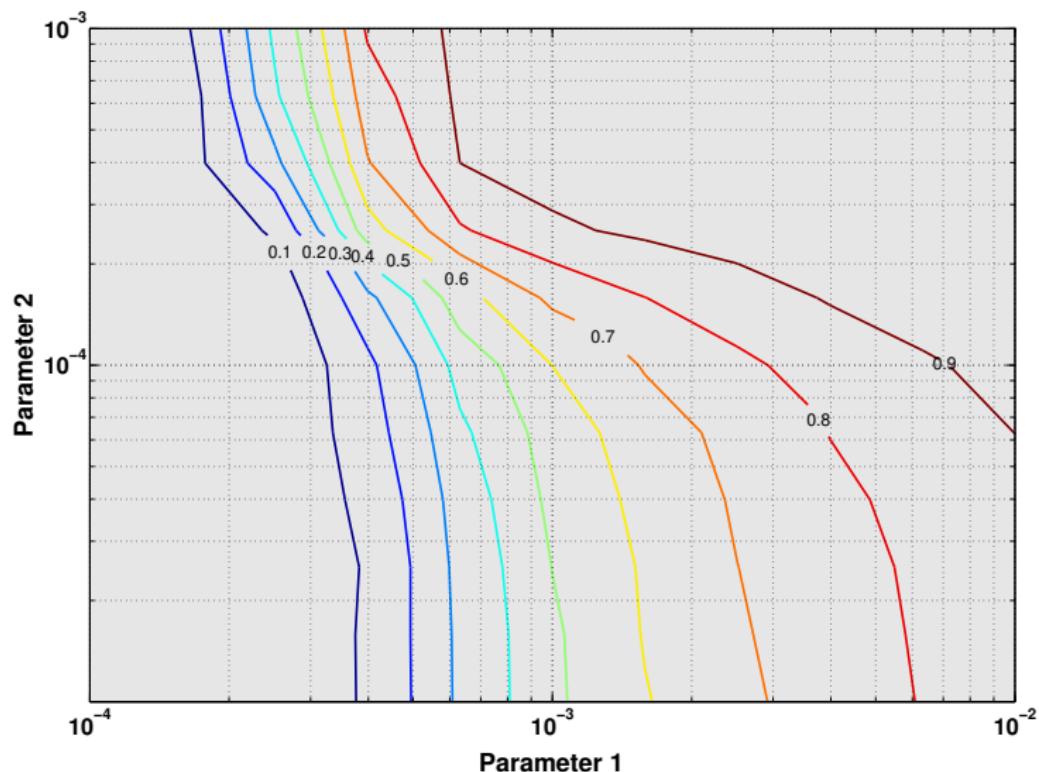
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ComK molecule count

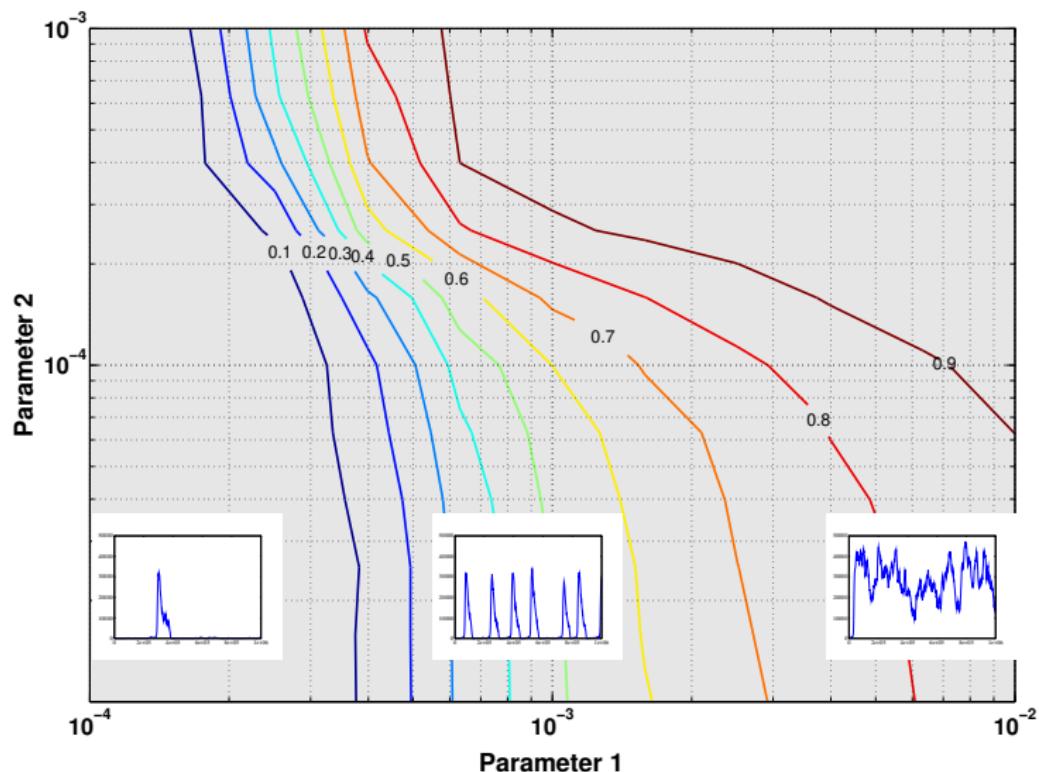


### Contours of probability of competence $P_c$



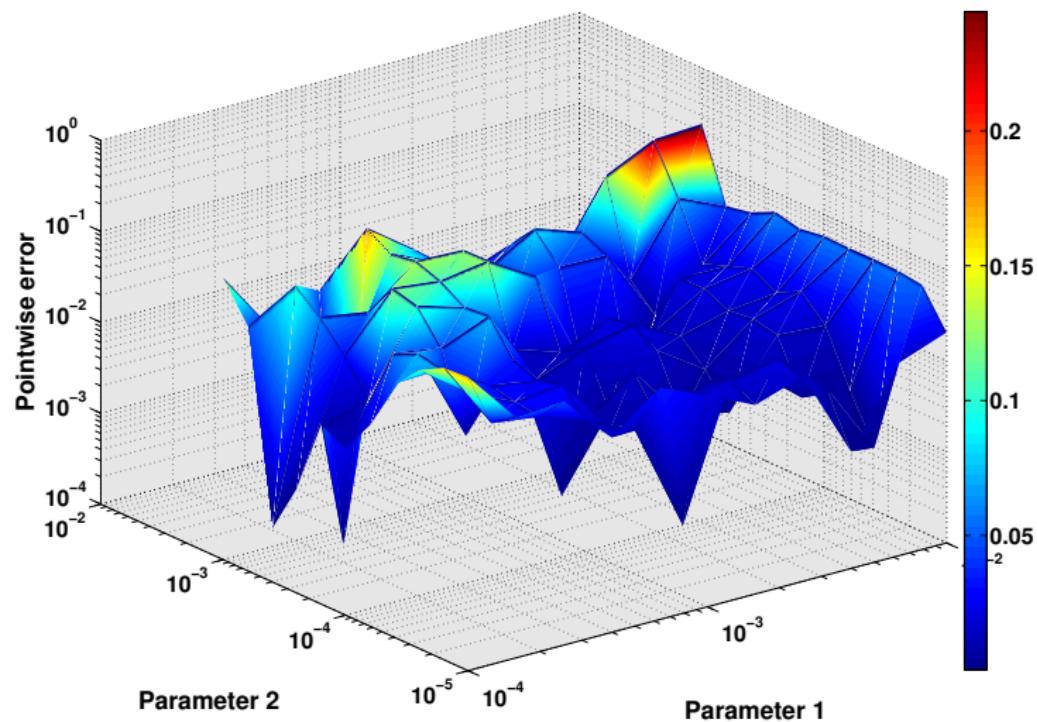
# Various dynamical regimes revealed by exploring parameter space

## Contours of probability of competence $P_c$



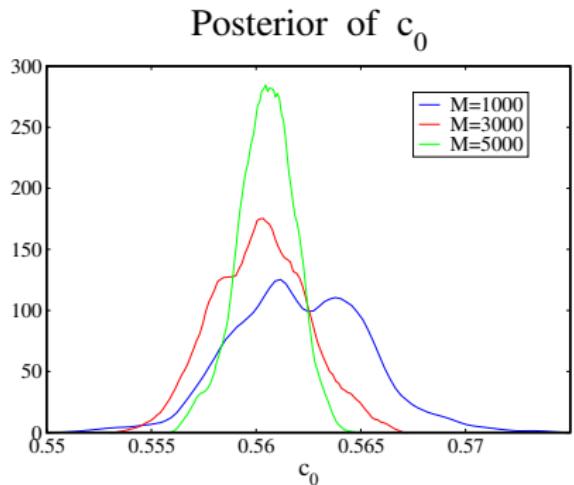
## Uncertain response surface in two-parameter case, 4th order PC

# Pointwise error in MAP estimate of 4th order PC



# Convergence both in posterior width and order

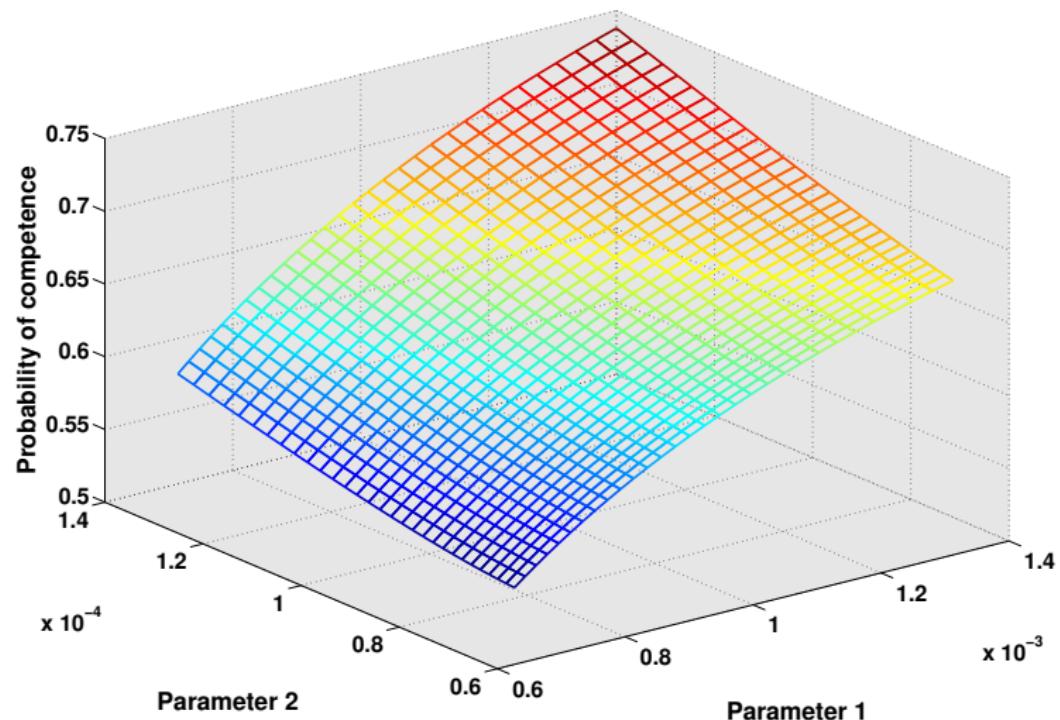
- With more input parameter samples, posterior narrows around the true value
- Convergence in PC order is established



Relative  $L_2$  error



## MAP estimate of 2-nd order response surface in 10-d case



# High Dimensional Model Representation (HDMR)

breaks the function into group-wise contributions of input variables

$$f(\boldsymbol{\lambda}) = f(\lambda_1, \dots, \lambda_d) = f_0 + \sum_i f_i(\lambda_i) + \sum_{i < j} f_{ij}(\lambda_i, \lambda_j) + \sum_{i < j < k} f_{ijk}(\lambda_i, \lambda_j, \lambda_k) + \dots$$

Component functions are found by

$$f_0 = \int_{\mathbb{R}^d} f(\boldsymbol{\lambda}) d\boldsymbol{\lambda}, \quad f_i(\lambda_i) = \int_{\mathbb{R}^{d-1}} f(\lambda_i, \boldsymbol{\lambda}_{\bar{i}}) d\boldsymbol{\lambda}_{\bar{i}} - f_0$$

$$f_{ij}(\lambda_i, \lambda_j) = \int_{\mathbb{R}^{d-2}} f(\lambda_i, \lambda_j, \boldsymbol{\lambda}_{\bar{j}}) d\boldsymbol{\lambda}_{\bar{j}} - f_i(\lambda_i) - f_j(\lambda_j) - f_0$$

- Component function  $f_{i_1 \dots i_s}(\lambda_{i_1}, \dots, \lambda_{i_s})$  is found by a  $(d - s)$ -dimensional integral. Still too high-dimensional.
- Otherwise called ANOVA decomposition (analysis of variance)
- Exact in the limit, but not unique.

# High Dimensional Model Representation (HDMR)

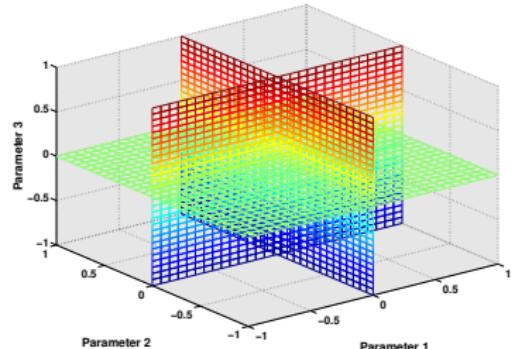
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- Two major variants:
  - Cut-HDMR
  - RS-HDMR



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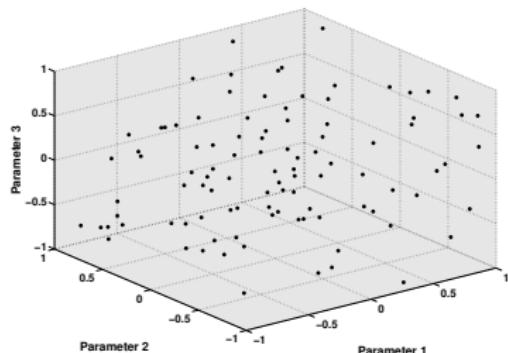
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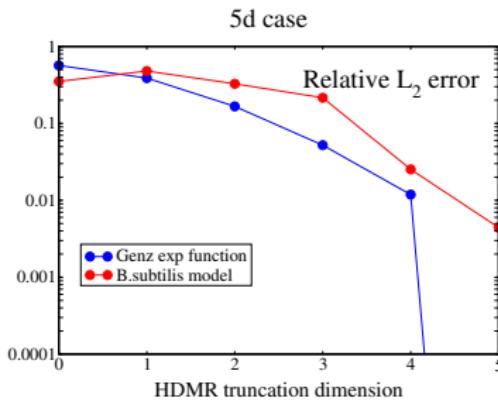
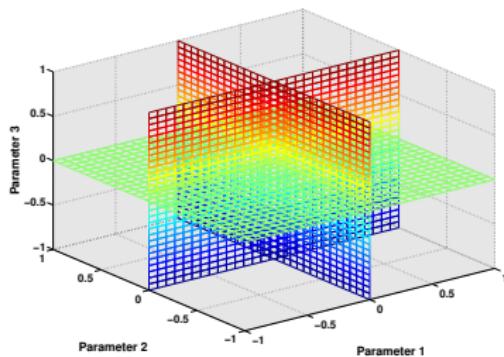


cut-HDMR disregards corners in the parameter space  
and does not guarantee accuracy in general

- Component functions are

$$f_0 = f(\boldsymbol{\lambda}^a), \quad f_i(\lambda_i) = f(\lambda_i, \boldsymbol{\lambda}_{\bar{i}}^a) - f_0 \\ f_{ij}(\lambda_i, \lambda_j) = f(\lambda_i, \lambda_j, \boldsymbol{\lambda}_{\bar{ij}}^a) - f_i(\lambda_i) - f_j(\lambda_j) - f_0$$

- Relies on values at lower-dimensional hyperplanes
- Depends on the anchor point  $\boldsymbol{\lambda}^a$
- Does not account for ‘corners’

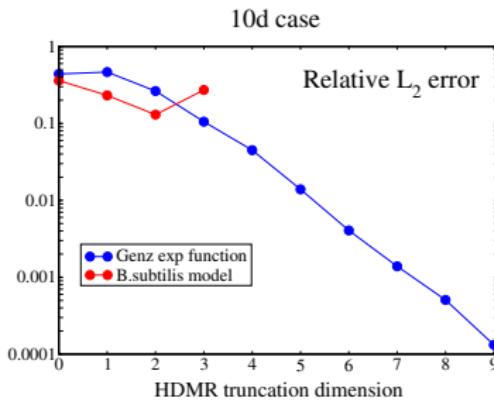
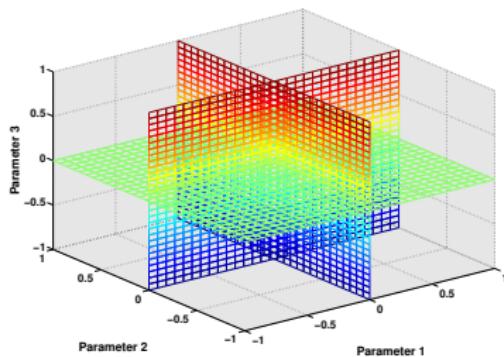


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## Random Sampling (RS) HDMR in principle equivalent to PC expansion with Monte-Carlo integration

$$f(\boldsymbol{\lambda}) = f(\lambda_1, \dots, \lambda_d) = f_0 + \sum_i f_i(\lambda_i) + \sum_{i < j} f_{ij}(\lambda_i, \lambda_j) + \sum_{i < j < k} f_{ijk}(\lambda_i, \lambda_j, \lambda_k) + \dots$$

- Component functions are found by

$$f_0 = \int_{\mathbb{R}^d} f(\boldsymbol{\lambda}) d\boldsymbol{\lambda}, \quad f_i(\lambda_i) = \int_{\mathbb{R}^{d-1}} f(\lambda_i, \boldsymbol{\lambda}_{\bar{i}}) d\boldsymbol{\lambda}_{\bar{i}} - f_0$$
$$f_{ij}(\lambda_i, \lambda_j) = \int_{\mathbb{R}^{d-2}} f(\lambda_i, \lambda_j, \boldsymbol{\lambda}_{\bar{ij}}) d\boldsymbol{\lambda}_{\bar{ij}} - f_i(\lambda_i) - f_j(\lambda_j) - f_0$$

- MC integrals still too expensive (new random samples needed for each hyperplane)
- Represent component functions with a polynomial expansion and use same set of samples
- Equivalent to Monte-Carlo PC with reordered multiindices!
  - PC (total order):  $f(\xi_1, \xi_2) = 1 + [\xi_1 + \xi_2] + [(\xi_1^2 - 1) + \xi_1 \xi_2 + (\xi_2^2 - 1)] + \dots$
  - RS-HDMR:  $f(\xi_1, \xi_2) = 1 + [\xi_1 + (\xi_1^2 - 1)] + [\xi_2 + (\xi_2^2 - 1)] + \xi_1 \xi_2 \dots$
- In future: employ Bayesian inference on component functions.

# Summary

- Polynomial Chaos expansions represent effects of uncertainties of input parameters to output statistical properties
  - Sensitivity analysis
  - Uncertainty quantification
  - Response surface construction
- Noise in function evaluations hampers quadrature methods
  - Sparse integration of noisy functions useless in high-d !
- HDMR constructions do not always guarantee accuracy with small computational effort
  - Generally still require high-d integrals
  - cut-HDMR overcomes this requirement but is not accurate enough
- Bayesian inference well-suited to handle noisy data

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