

# Bayesian Quantification of Uncertainty in Systems with Intrinsic Noise

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*B. Subtilis* endospore stain, by A. Schenkel, P. Justice and E. Suchman, Colorado State University.

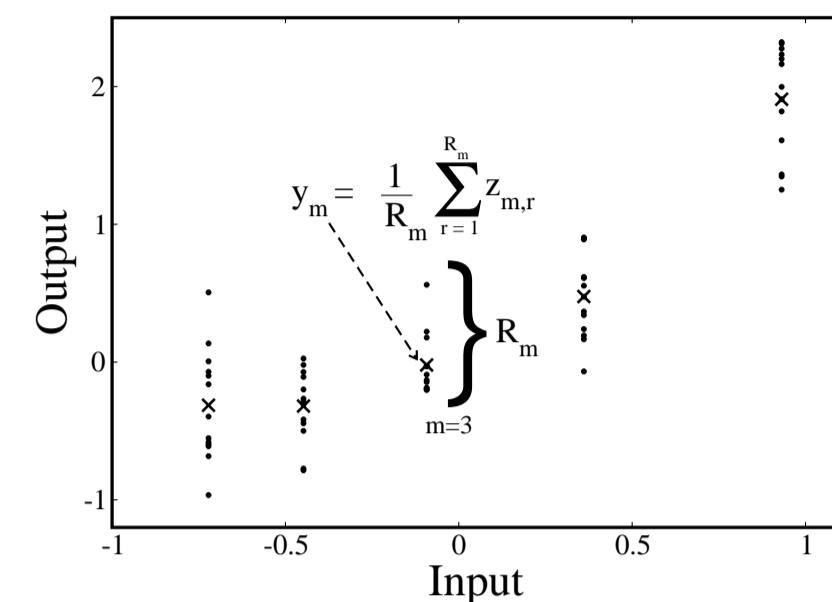
## Stochasticity plays an important role in many phenomena

- In stochastic reaction networks, intrinsic stochasticity is due to reactions between small number of molecules
- Applications
  - Gene regulatory networks, bioenergy and bioremediation
  - Interfacial reaction processes, fuel cells and batteries
  - Cellular signaling, immunology

- Uncertainty sources include intrinsic stochasticity, parametric uncertainty, sparsity of the available data, experimental noise.
- Questions that uncertainty quantification helps to answer
  - How predictive is the model?
  - If the model is good enough, what is the mismatch with experiments due to?
  - Does the system work in spite of the noise or because of it?

## Problem formulation

- Stochastic model  $Y(\lambda)$  with a  $d$ -dimensional input parameter vector  $\lambda = (\lambda_1, \dots, \lambda_d)$
- Observable of interest  $y = \mathbb{E}[Y]$
- Training runs at  $M$  input parameter values
- $R_m$  replica runs for  $m$ -th input parameter
- A total of  $N = \sum_{m=1}^M R_m$  model evaluations,  $\{z_{m,r}\}$



## Polynomial chaos spectral representation

To build a representation for input-output relationship, Polynomial Chaos (PC) spectral expansions are used; see Ghanem and Spanos, "Stochastic Finite Elements: A Spectral Approach", 1991.

- Interprets input parameters as random variables
- Allows propagation of input parameter uncertainties to outputs of interest
- Serves as a computationally inexpensive surrogate for calibration or optimization

Input parameters are represented via their cumulative distribution function (CDF)

$$\eta_i = 2F_{\lambda_i}(\lambda_i) - 1, \quad \text{for } i = 1, 2, \dots, d.$$

If input parameters are uniform  $\lambda_i \sim \text{Uniform}[a_i, b_i]$ , then

$$\eta_i = \frac{2}{b_i - a_i} \left( \lambda_i - \frac{a_i + b_i}{2} \right).$$

Output is represented with respect to Legendre polynomials

$$y(\eta) \approx y_C(\eta) \equiv \sum_{k=0}^K c_k \Psi_k(\eta).$$

## Sparse quadrature intergation fails with noisy data

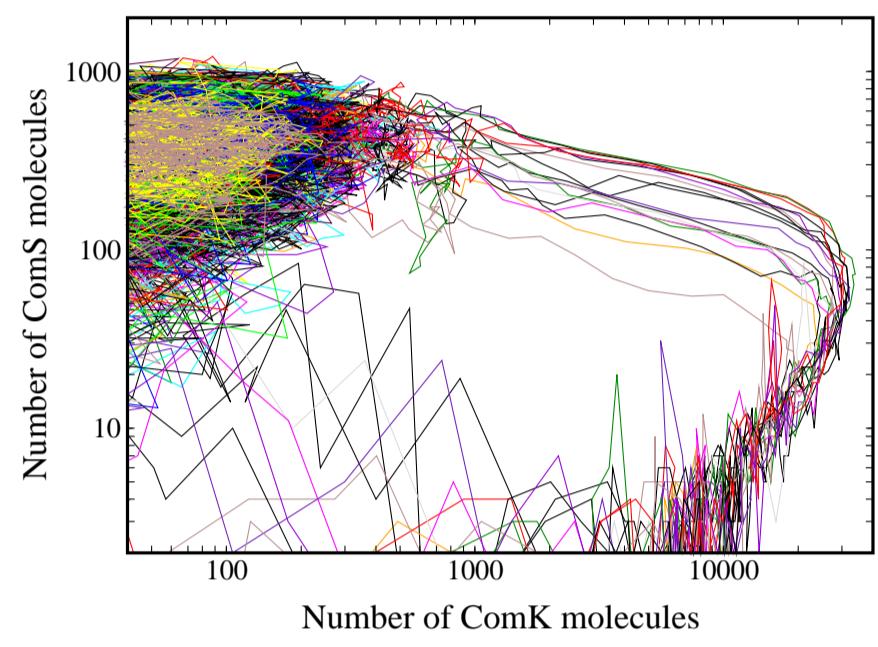
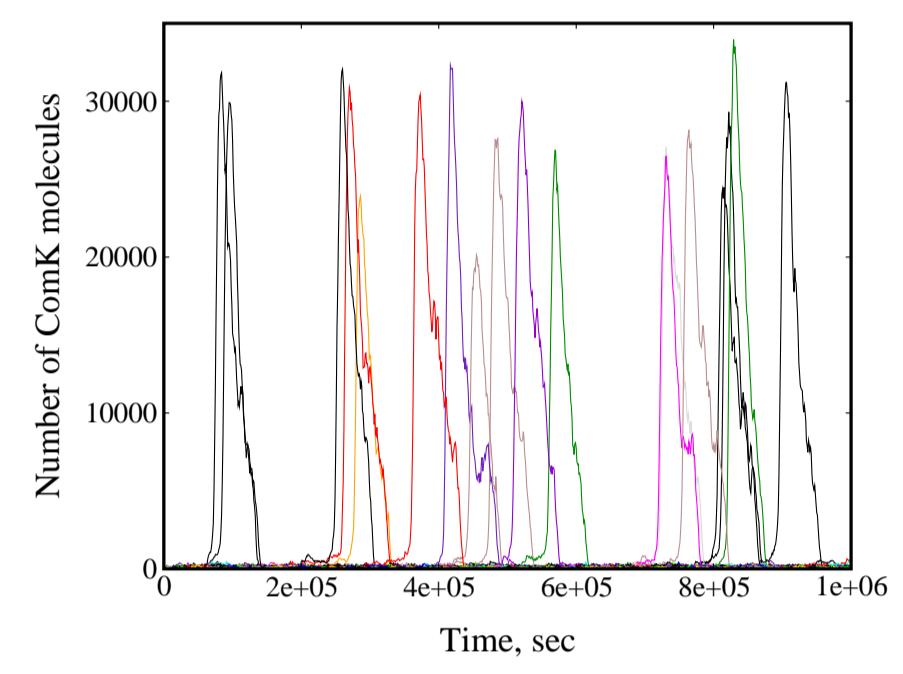
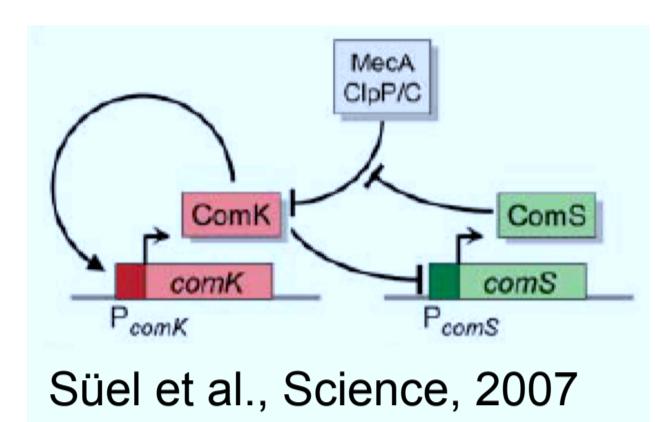
Quadrature approaches fail as well.

$$\sum_{q=1}^Q y_q \Psi_k(\eta_q) w_q$$

- *Tensor product* quadrature suffers from the curse of dimensionality
- *Sparse grid* quadrature is infeasible for noisy systems due to negative weights. Even a very small error in function evaluation is amplified by a factor that increases with dimensionality!

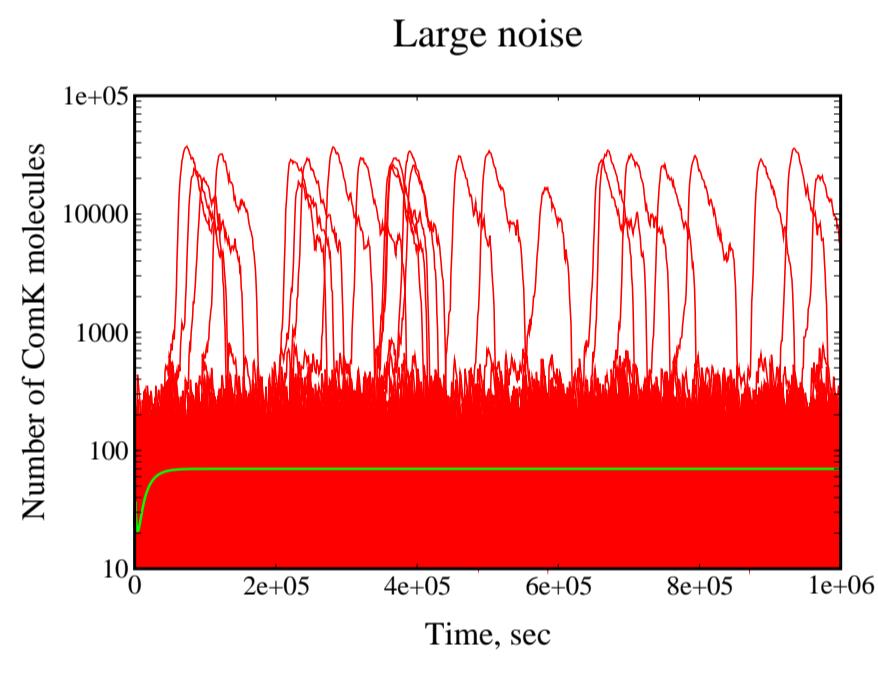
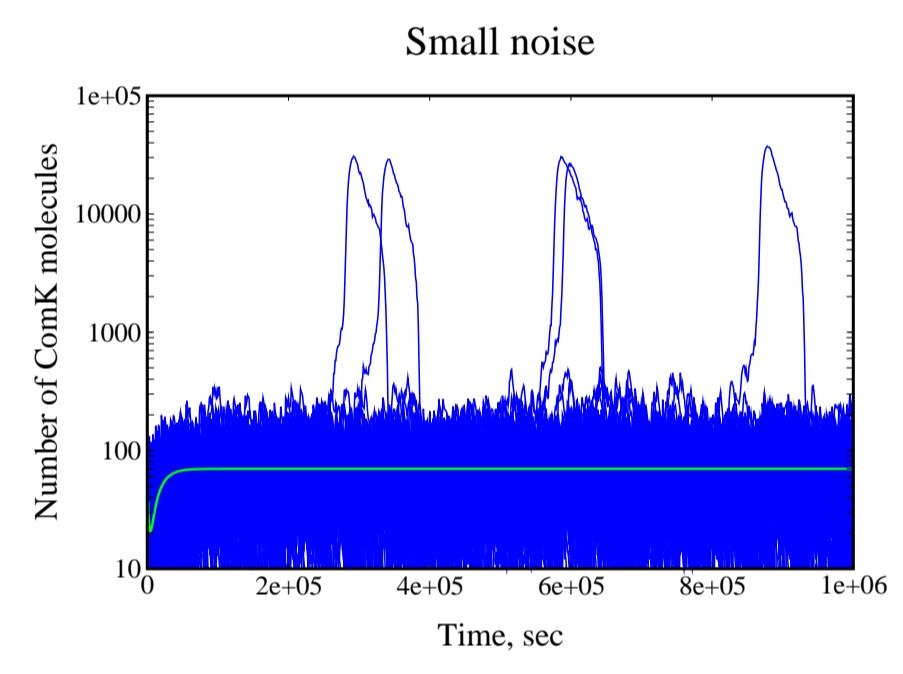
## *Bacillus Subtilis* is a gram positive soil bacterium

- Competence in *B. Subtilis* is a state that allows uptake of external DNA
- It is characterized by a sporadic jump in the number of comK molecules
- Stochastic reaction network of competence dynamics consists of 11 species and 16 reactions, see Suel *et al.*, Science, 2007
- Input parameters are reaction rate parameters in logarithmic scale,  $\eta = \log \tilde{k} \pm \log f$ , i.e. the range is  $[\tilde{k}/f, \tilde{k}f]$  with a range factor  $f > 1$  and a nominal parameter value  $\tilde{k}$ .



## ODE limit and noise-induced transition to competence

- Competence events, i.e. sporadic jumps in the number of comK molecules, are driven by noise
- In the limit of large volume, the system is described by a system of ODEs, called rate equations
- By tuning reaction network parameters in a special way, one can keep the corresponding ODE limit unchanged, focusing on pure noise dependence



## Bayesian inference of PC modes

Bayesian framework allows quantifying different sources of uncertainties - parametric, intrinsic, or uncertainties associated with lack-of-sampling.

Bayes formula

$$p(c|D) \propto L_D(c)p(c)$$

relates prior distribution  $p(c)$  of PC modes to the posterior  $p(c|D)$ , where the data  $D$  is the set of all training runs  $\{z_{m,r}\}$ ,  $m = 1 : M$ ,  $r = 1 : R_m$ .

Estimates of the mean of the data  $z_{m,r}$  and its variance at the  $m$ -th parameter location are, respectively,

$$y_m = \frac{1}{R_m} \sum_{r=1}^{R_m} z_{m,r},$$

$$s_m^2 = \frac{1}{R_m - 1} \sum_{r=1}^{R_m} (z_{m,r} - y_m)^2.$$

Prior distribution on  $c$  is uniform,  $p(c) = \text{const.}$

The likelihood accounts for the discrepancy between the averaged data and the model,

$$L_D(c) = L_D(c; s^2) = \frac{1}{(2\pi)^{M/2} \prod_{m=1}^M (s_m/\sqrt{R_m})} \exp \left( -\sum_{m=1}^M \frac{(y_m - y_C(\eta_m))^2}{2s_m^2/R_m} \right)$$

The posterior is analytically tractable, it is a multivariate normal distribution,

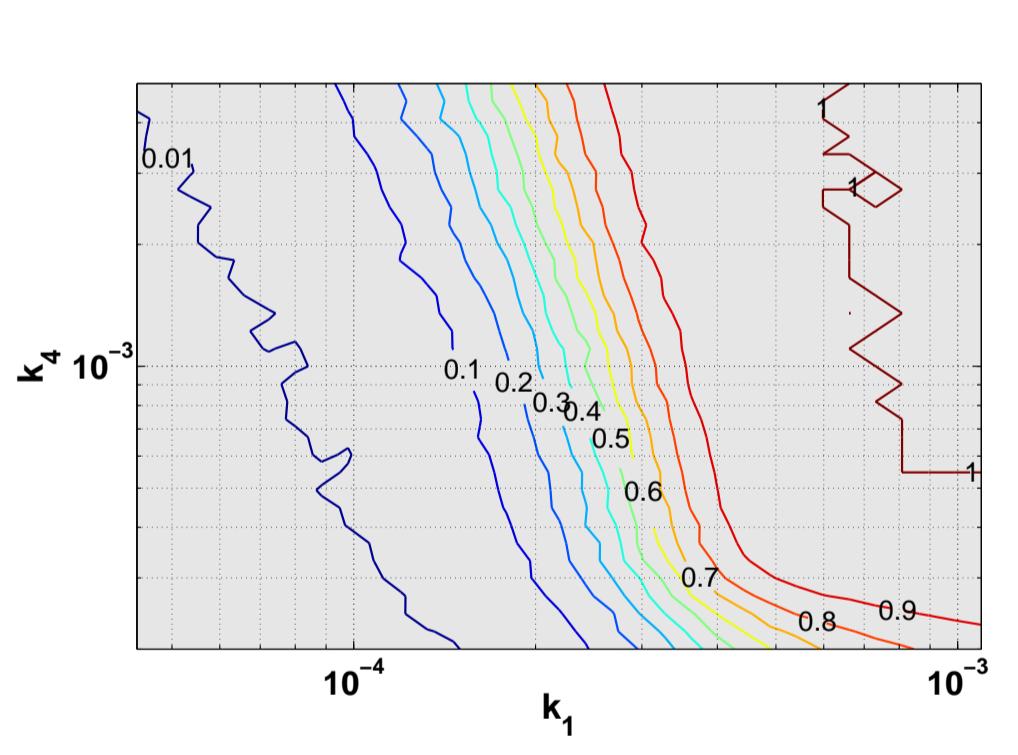
$$c \in \mathcal{MVN}(\underbrace{\Psi^T Q^{-1} \Psi^{-1} \Psi^T Q^{-1} y}_{\text{mean}}, \underbrace{\Psi^T Q^{-1} \Psi^{-1}}_{\text{covariance}}),$$

where  $\Psi$  is a  $M \times (K+1)$  matrix with elements  $\Psi_{mk} = \Psi_k(\eta_m)$  and  $Q$  is a diagonal weight matrix with entries  $Q_{mm'} = \delta_{m,m'} R_m / (2s_m^2)$ .

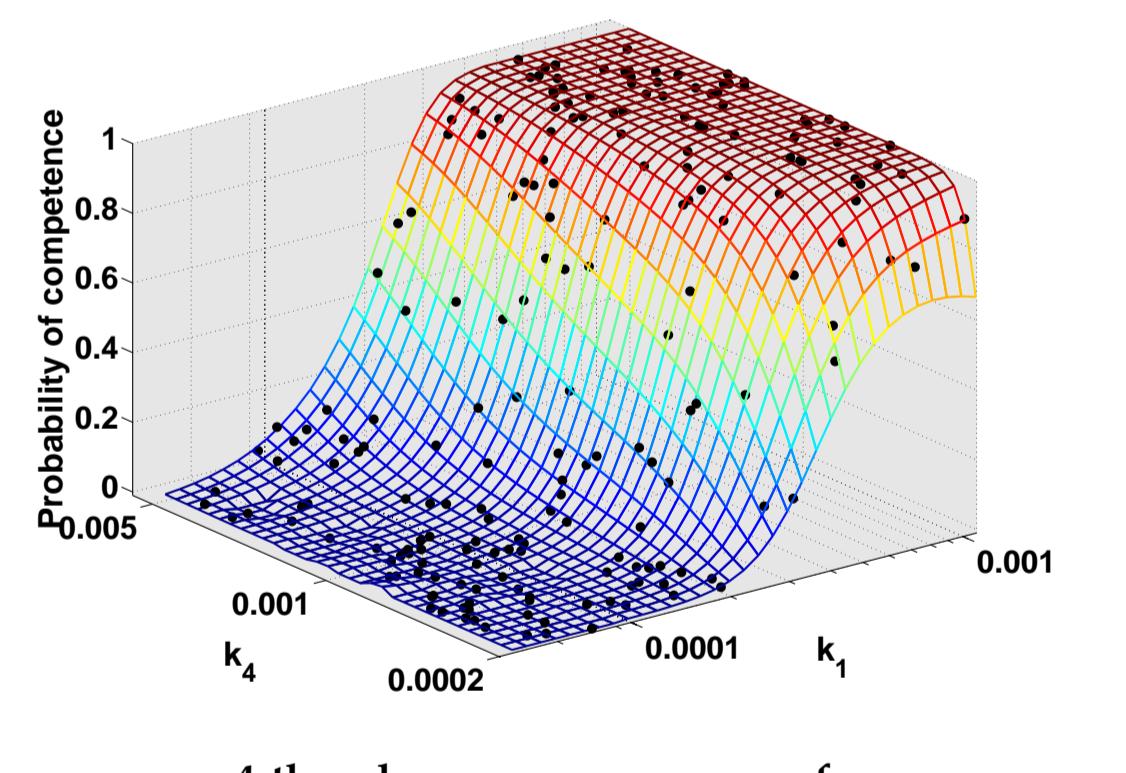
## Mixture PC expansion based on nearest neighbor classification

- If data has quantitatively different behavior in different regions, global polynomial fit is inaccurate
- A mixture PC formulation is developed based on a nearest neighbor classification
  - The input set of points is clustered according to the corresponding output values
  - For each cluster, a separate PC expansion is obtained
  - The resulting expansion is a weighted sum of PC expansions for a certain number of nearest neighbors
- If the output values are bounded, a map to  $(-\infty; +\infty)$  is utilized before PC representation to keep the approximation from exceeding physical bounds
  - For example, if  $y \in [0, 1]$ , the effective output is  $\bar{y} = \log \frac{y}{1-y}$

Sargsyan *et al.*, "Multiparameter spectral representation of noise-induced competence in *Bacillus Subtilis*", to be submitted to *Biophys J*, 2011.

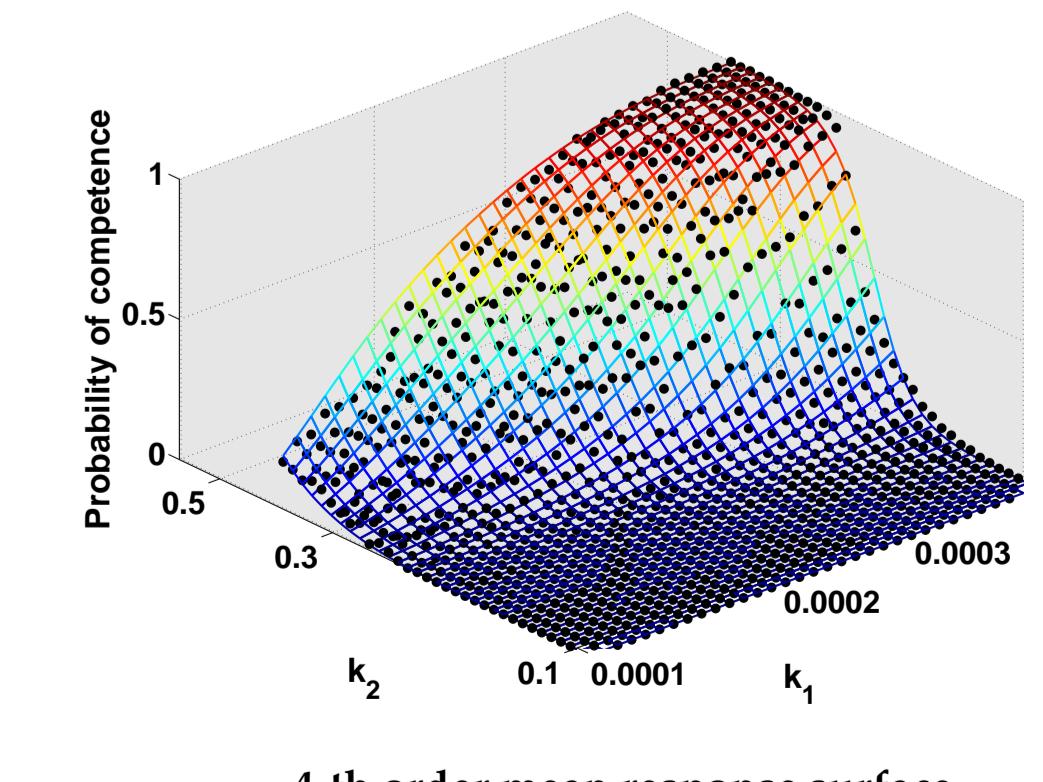
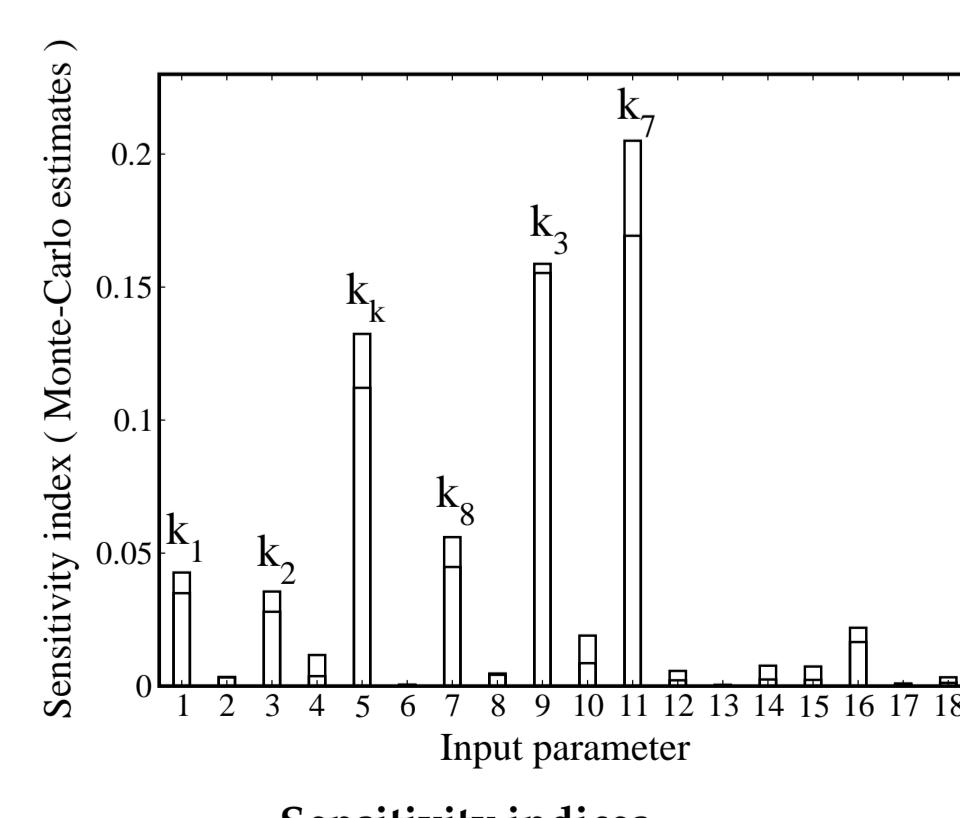


Contour plots of the probability of competence



## Dimensionality reduction using sensitivity indices

- All 18 reaction rate parameters are taken
- Due to sparsity of the data ( $M = 1000$  points in 18-dimensional parameter space) the global PC expansion is more reliable than the clustering-based mixture PC
- Variance-based sensitivity indices  $S_i = \frac{\text{Var}[E(y_C(\eta)|\eta_i)]}{\text{Var}[y_C(\eta)]}$  are computed from the global PC to down-select from 18 dimensions to 6 dimensions
- The comK-related reaction parameters have shown larger sensitivity indices
- For the resulting 6-dimensional problem, a mixture PC is constructed and shown to be more accurate
- For each of the  $M = 1000$  input parameters,  $R_m = 100$  replica simulations are taken
- The resulting *uncertain* response surface has a relative  $L_2$  error of  $\sim 0.08$



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