## Calibration of computer models Extension to Stochastic Simulator

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- Statistical Models
- Heteroskedastic GP
- Calibration
  - KOH
  - ABC

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# Stochasticity in Computer Experiments

Basic model for a stochastic simulator:

$$f(\mathbf{x}) = m(\mathbf{x}) + v, \ v \sim N(0, \sigma_v^2(x)), \tag{1}$$

#### where

- $m(\mathbf{x})$  is the expected value:  $\mathbb{E}_f[f(\mathbf{x})]$ ,
- v independent variability representing randomness of the simulator,
- variance  $\sigma_v^2$  may depend on **x**, be constant.

#### Remarks

- If  $\sigma_v^2 = 0$  deterministic simulator,
- What does stochasticity acknowledge for? numerical approximation (Monte Carlo), aleatory experiment.



#### Extension of KOH

$$y_i^{\theta} = y^{\theta}(\mathbf{x}_i^{\theta}) = f(\mathbf{x}_i^{\theta}, \theta) + \delta(\mathbf{x}_i^{\theta}) + \epsilon_i,$$
 (2)

#### where

- $\mathbf{y}^e = \{y_1^e, \dots, y_{n_e}^e\}$  are real-world field observations at controllable (or measurable) inputs  $(\mathbf{x}_i)_{1 \leq \dots \leq n_e}$ ,
- f is a stochastic simulator.
- $\bullet$  is measurement error for the observations,
- ullet  $\delta$  is the discrepancy may be assumed to be stochastic.

#### Remarks:

If reality is stochastic,  $\delta$  has to be stochastic and may be heteroskedastic as the simulator.

[Sung et al., 2019] use a hetGP for the discrepancy (but with a deterministic simulator), estimating parameters via maximum likelihood and following [Tuo et al., 2015] to avoid confounding.



5/19

- Statistical Models
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# Stochastic Kriging

### [Ankenman et al., 2010]

Observation model:

$$y_i^e = f(\mathbf{x}_i^e) = m(\mathbf{x}_i^e) + v_i$$
, with  $v_i \stackrel{ind}{\sim} \mathcal{N}(0, r(\mathbf{x}_i^e))$ .

In homoskedastic cases  $r(\mathbf{x}_i^e) = \tau^2$  which is called the nugget.

From a design with replications:

- 'full-N" dataset, n of unique x<sub>i</sub>-values in X<sub>N</sub> with n << N, a<sub>i</sub> replicates at unique locations,
- compute

$$\bar{y}_i = \frac{1}{a_i} \sum_{j=1}^{a_i} y_i^{(j)}$$
 and  $\hat{\sigma}_i^2 = \frac{1}{a_i - 1} \sum_{j=1}^{a_i} (y_i^{(j)} - \bar{y}_i)^2$ .

predictions with BLUP for *M* when GP is assumed on *M*:

$$\begin{split} \mu_n^{\text{SK}}(\mathbf{x}) &= k_n^{\top}(\mathbf{x})(C_n + S_n)^{-1} \, \overline{Y}_n \\ \sigma_n^{\text{SK}}(\mathbf{x})^2 &= c_{S,\psi}(\mathbf{x},\mathbf{x}) - k_n^{\top}(\mathbf{x})(C_n + S_n)^{-1} k_n(\mathbf{x}), \end{split}$$

$$k_n(\mathbf{x}) = (c_{S,\psi}(\mathbf{x},\bar{\mathbf{x}}_1),\dots,c_{S,\psi}(\mathbf{x},\bar{\mathbf{x}}_n))^{\top} S_n = [\hat{\sigma}_{1:n}^2]A_n^{-1} = \operatorname{Diag}(\hat{\sigma}_1^2/a_1,\dots,\hat{\sigma}_n^2/a_n), \text{ and } C_n = \{c_{S,\psi}(\bar{\mathbf{x}}_i,\bar{\mathbf{x}}_i)\}_{1 \le i,i \le n}.$$

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7/19

# Modeling the variance

- In [Ankenman et al., 2010], no specific model for the variance,
- [Goldberg et al., 1997] assumes log(r(x)) ~ GP for modeling heteroskedasticity, and they estimate the combined parameters of the two GPs with an MCMC scheme.

[Binois et al., 2018] make use of Stochastic Kriging with GP model for  $log(r(\mathbf{x}))$ .

- consider latent variances:  $\xi_1, \dots, \xi_n$  for the *n* unique locations,
- GP prior on this matrix  $\Xi_n \sim \mathcal{N}_n(0, \nu(C_{\xi} + g_{\xi}A_n^{-1}))$  where  $g_{\xi}$  regularizes the behavior of the variance process,
- Estimate parameters by MLE using Woodbury trick which put all the computation in O(n³),
- implementation in hetGP package.



## homoskedastic GP

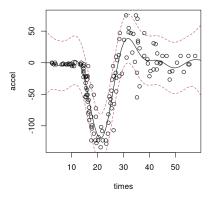


Figure: Homoskedastic GP fit to the motorcycle data via mean (solid-black) and 90% error-bars (dashed-red).

from [Gramacy, 2020]



### heteroskedastic GP

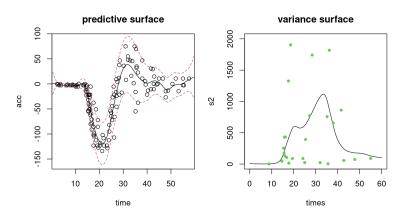


Figure: Heteroskedastic GP fit to the motorcycle data. Left panel shows the predictive distribution via mean (solid-black) and 90% error-bars (dashed-red). Right panel shows the estimated variance surface and moment-based estimates of variance (green dots).

from [Gramacy, 2020]



10/19

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#### Calibration of Stochastic Simulators

$$y_i^{\theta} = f(\mathbf{x}_i^{\theta}, \boldsymbol{\theta}^*) + \delta(\mathbf{x}_i^{\theta}) + \epsilon(\mathbf{x}_i^{\theta}).$$
 (3)

 $\delta(\cdot)$  models the difference between the simulator and the physical system:

$$\delta(\mathbf{x}) = \zeta(\mathbf{x}) - f(\mathbf{x}, \theta^*).$$

Here f is Stochastic but its link with reality is questionable. Is reality  $\mathbb{E}(f)$  or f? Depending on that,  $\delta$  should be considered as deterministic or Stochastic and then modeled as a standard GP...



13/19

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# Ocean Example

see https://github.com/Demiperimetre/Ocean



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## **History Matching**

[Andrianakis et al., 2015] contains a thorough description of HM whilst applying it to a complex epidemiology model of HIV.



15/19

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  - KOH
  - ABC

#### **Basics**

Approximate Bayesian Computation produces samples from a posterior distribution  $\pi(\theta|\mathbf{y}^e)$  by

- ullet generating samples for heta from the prior
- and outputs **y** from the generating model  $\pi(\cdot|\mathbf{y})$  (this implies runs of the simulator),
- samples are kept provided that  $\mathbf{y} = \mathbf{y}^e$  or  $|h(\mathbf{y}) h(\mathbf{y}^e)| < tol$ ,
- accepted  $\theta$ s produce an approximated posterior sample.

#### Remark:

- For calibration, tol can be interpreted as a bound on the observational error and model discrepancy, leading to a "correct" posterior rather than an approximation [Wilkinson, 2013]. This is then similar to HM with the subjective choice of bounds.
- ABC can be done without the use of a surrogate, but this will require many runs of the simulator itself. Otherwise, very few accepted  $\theta$  will be obtained, or an overly high value of tol will be required.



17/19

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## Fish example

See https://github.com/jhuang672/fish/blob/master/fish\_fits.md



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# Other papers

[Oakley and Youngman, 2017] removes  $\delta$  but compensates by inflating the variability in the simulator output.



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Bayesian history matching of complex infectious disease models using emulation: a tutorial and a case study on hiv in uganda.

PLoS computational biology, 11(1):e1003968.



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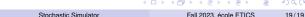
In Proceedings of the 10th International Conference on Neural Information Processing Systems, pages 493–499.



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Calibration of stochastic computer simulators using likelihood emulation. Technometrics, 59(1):80–92.



Sung, C.-L., Barber, B. D., and Walker, B. J. (2019).

Calibration of computer models with heteroscedastic errors and application to plant relative growth rates.

arXiv preprint arXiv:1910.11518.



Tuo, R., Wu, C. J., et al. (2015).

Efficient calibration for imperfect computer models.

The Annals of Statistics, 43(6):2331–2352.



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Approximate Bayesian computation (abc) gives exact results under the assumption of model error.

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