Calibration of computer models Extension to Stochastic Simulator

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Stochastic Simulator

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Outline

- 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 σ_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

$$y_i^e = y^e(\mathbf{x}_i^e) = f(\mathbf{x}_i^e, \theta) + \delta(\mathbf{x}_i^e) + \epsilon_i,$$
 (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,
- \bullet δ is the discrepancy may be assumed to be stochastic.

Remarks: If reality is stochastic, δ has to be stochastic and may be heteroskedastic as the simulator.

 y_S is the simulator with additional unknown, non-measurable, inputs u_C , $y_F(x)$ (with variance σ_ϵ^2), and $\delta_{\rm MD}(x)$ is an important term that accounts for the simulator not being a perfect representation of reality. y_F "observes" reality with error ϵ ; reality = $y_S + \delta_{\rm MD}$.

[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.

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Stochastic Kriging

[Ankenman et al., 2010]

Observation model:

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

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

From a design with replications:

- 'full-N' dataset, n of unique x_i -values in X_N with n << N, a_i 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*:

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

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

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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: ξ_1, \dots, ξ_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_{ξ} 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.



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homoskedastic GP

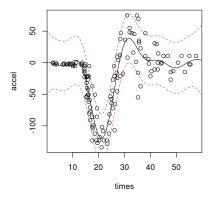


Figure: Homoskedastic GP fit to the motorcycle data via mean (solid-black) and 90% error-bars (dashed-red).

from [Gramacy, 2020]



heteroskedastic GP

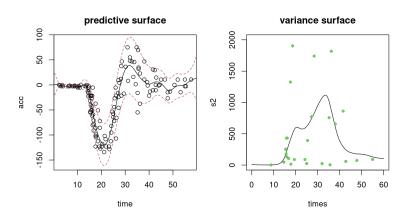


Figure: Heteroskedastic GP fit to the motorcycle data. Left panel shows the predictive distribution via mean (solid-black) and 90% error-bars (dashed-red). Right panel shows the estimated variance surface and moment-based estimates of variance (green dots).

from [Gramacy, 2020]



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KOH

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KOH

Calibration of Stochastic Simulators

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

 $\delta(\cdot)$ 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, δ should be considered as deterministic or Stochastic and then modeled as a standard GP...



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KOH

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.



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ABC

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Approximate Bayesian Computation produces samples from a posterior distribution $\pi(\theta|\mathbf{y}^{\theta})$ 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 θ 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 θ will be obtained, or an overly high value of *tol* will be required.



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ABC

Fish example

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



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

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Andrianakis, I., Vernon, I. R., McCreesh, N., McKinley, T. J., Oakley, J. E., Nsubuga, R. N., Goldstein, M., and White, R. G. (2015).

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.



Ankenman, B., Nelson, B. L., and Staum, J. (2010). Stochastic kriging for simulation metamodeling. Operations Research, 58(2):371–382.



Binois, M., Gramacy, R. B., and Ludkovski, M. (2018). Practical heteroscedastic Gaussian process modeling for large simulation experiments.

Journal of Computational and Graphical Statistics, 27(4):808-821.



Goldberg, P. W., Williams, C. K., and Bishop, C. M. (1997).

Regression with input-dependent noise: a Gaussian process treatment.

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



Gramacy, R. B. (2020).

 $\underline{\text{Surrogates: Gaussian process modeling, design, and optimization for the applied } \underline{\text{sciences}}.$

CRC press.



Oakley, J. E. and Youngman, B. D. (2017).

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.



Wilkinson, R. D. (2013).

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

Statistical Applications in Genetics and Molecular Biology, 12(2):129-141.

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