Validating Gaussian Process Emulators

Leo Bastos

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Joint work: Jeremy Oakley and Tony O'Hagan



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Outline

Computer model

Definition

2 Emulation

- Gaussian Process Emulator
- Toy Example

3 Diagnostics and Validation

- Numerical diagnostics
- Graphical diagnostics
- Examples

Conclusions



- **Computer model** is a mathematical representation $\eta(\cdot)$ of a complex physical system implemented in a computer.
- We need Computer models when real experiments are very expensive or even implossible to be "done" (e.g. Nuclear experiments)
- Computer models have an important role in almost all fields of science and technology
 - System Biology models (Rotavirus outbreaks).
 - Cosmological models (Galaxy formation)
 - Climate models* (Global warming).



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- area of sea ice
- thickness of sea ice
- atmospheric CO2 concentrations
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- Large number of outputs (Both time series and field data)
- Several inputs (e.g. model resolution, initial conditions)
- Each run takes about an hour on the Linux Boxes at NOC



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Computer models can be very expensive



IBM supercomputers used for climate and weather forecasts

• One single run of the computer model can take a lot of time





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IBM supercomputers used for climate and weather forecasts

- One single run of the computer model can take a lot of time
- Most of analyses need several runs



Emulating a computer model

• $\eta(\cdot)$ is considered an unknown function



- Emulator is a predictive function for the computer model outputs
- Assumptions for the computer model:
 - Deterministic single-output model $g(\cdot) = g : \mathcal{X} \in \Re^{p} \to \Re$
 - Relatively "Smooth" function
- Statistical Emulator is an stochastic representation of our judgements about the computer model $\eta(\cdot)$.



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Gaussian process emulator:

$$\eta(\cdot)|\beta,\sigma^2,\psi\sim GP(m_0(\cdot),V_0(\cdot,\cdot)),$$

where

$$m_0(\mathbf{x}) = h(\mathbf{x})^T \beta$$

$$V_0(\mathbf{x}, \mathbf{x}') = \sigma^2 C(\mathbf{x}, \mathbf{x}'; \psi)$$

• Prior distribution for (β, σ^2, ψ)

Conditioning on some training data

$$y_k = \eta(\mathbf{x_k}), \quad k = 1, \dots, n$$



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Predictive Gaussian Process Emulator

 $\eta(\cdot)|\mathbf{y}, \mathbf{X}, \psi \sim \text{Student-Process}\left(n-q, m_1(\cdot), V_1(\cdot, \cdot)\right),$

where

$$\begin{aligned} m_1(x) &= h(x)^T \widehat{\beta} + t(x)^T \mathbf{A}^{-1} (\mathbf{y} - H \widehat{\beta}), \\ V_1(x, x') &= \widehat{\sigma}^2 \left[C(x, x'; \psi) - t(x)^T \mathbf{A}^{-1} t(x') + \left(h(x) - t(x)^T \mathbf{A}^{-1} H \right) \right. \\ &\times \left. (H^T \mathbf{A}^{-1} H)^{-1} \left(h(x') - t(x')^T \mathbf{A}^{-1} H \right)^T \right]. \end{aligned}$$



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• $\eta(\cdot)$ is a two-dimensional known function

• GP emulator:

• $h(\mathbf{x}) = (1, \mathbf{x})^{7}$ • $O(\mathbf{x}, \mathbf{x}) = \exp\left[-\sum_{k} \left(\frac{\mathbf{x}_{k} - \mathbf{x}_{k}}{\sigma_{k}}\right)^{2}\right]$ • $p(\beta, \sigma^{2}, \psi) \propto \sigma^{-2}$



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Toy example





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Design for Computer models

- Emulation (Multiple output emulation, Dynamic emulation)
- UA/SA Uncertainty and Sensitivity Analyses
- Calibration (Bayes Linear and Full Bayesian approaches)
- Diagnostics and Validation



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Diagnostics and Validation

• Every emulator should be validated

- Non-valid emulators can induce wrong conclusions
- There is little research into validating emulators
- Validation generally means: "the emulator predictions are close enough to the simulator outputs".
- We want to take account all the uncertainty associated with the emulator.
- "Do the choices that I have made, based on my knowledge of this simulator, appear to be consistent with the observations?"
- Choices for the Gaussian process emulator:
 - Normality
 - Stationarity
 - Correlation parameters



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Our diagnostics should be based on a set of new runs of the simulator

Why? Because predictions at observed input points are perfect.
 Validation data (y*, X*) : y_k* = η(x_k*), k = 1,..., m

Simulator and the predictive emulator outputs are compared

- Numerical diagnostics
- Graphical diagnostics



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Individual predictive errors

$$D_i^l(\mathbf{y}^*) = rac{(y_i^* - m_1(\mathbf{x}_i^*))}{\sqrt{V_1(\mathbf{x}_i^*, \mathbf{x}_i^*)}}$$

However, the $D^{l}(\mathbf{y}^{*})$ s are correlated:

 $D^{I}(\eta(\mathbf{X}^{*})) \sim \text{Student-t}_{m}(n-q, \mathbf{0}, C_{1}(\mathbf{X}^{*}))$

Mahalanobis distance

 $D_{MD}(\mathbf{y}^*) = (\mathbf{y}^* - m_1(\mathbf{X}^*))^T V_1(\mathbf{X}^*)^{-1} (\mathbf{y}^* - m_1(\mathbf{X}^*))$



Individual predictive errors

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$$D^{PC}(\mathbf{y}^*) = (\mathbf{G}^{-1})^T (\mathbf{y}^* - m_1(\mathbf{X}^*))$$

where $V_1(\mathbf{X}^*) = \mathbf{G}^T \mathbf{G}$, and $\mathbf{G} = \mathbf{P} \mathbf{R}^T$.

Properties:

- $D^{PC}(\mathbf{y}^*)^T D^{PC}(\mathbf{y}^*) = D_{MD}(\mathbf{y}^*)$
- $Var(D^{PG}(n(X)) = T$
- Invariant to the data order
- Pivoting order given by P has an intuitive explanation
- Each D^{PC}(y*) associated with a validation element



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- $Var(D^{PC}(\eta(\mathbf{X})) = \mathcal{I}$
- Invariant to the data order
- Pivoting order given by P has an intuitive explanation
- Each *D^{PC}*(**y**^{*}) associated with a validation element



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Pivoted Cholesky errors

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$$D^{PC}(\mathbf{y}^*) = (\mathbf{G}^{-1})^T (\mathbf{y}^* - m_1(\mathbf{X}^*))$$

where
$$V_1(\mathbf{X}^*) = \mathbf{G}^{\mathsf{T}}\mathbf{G}$$
, and $\mathbf{G} = \mathbf{P}\mathbf{R}^{\mathsf{T}}$.

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- Individual errors against emulator's predictions Problems on mean function, non-stationarity
- Errors againts the pivoting order Poor estimation of the variance, correlation parameters
- QQ-plots of the uncorrelated standardized errors
 Non-normality, Local fitting problems or non-stationarity
- Individual or (pivoted) Cholesky errors against inputs Non-stationarity, pattern not included in the mean function



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Example: Nuclear Waste Repository



Source: http://web.ead.anl.gov/resrad/

- RESRAD is a computer model designed to estimate radiation doses and risks from RESidual RADioactive materials.
- Output: 10,000 year time series of the release of contamination in the series of the release of contamination in the series of the release of contamination is the series of the series of the release of contamination is the series of the release of the series of the series

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Example: Nuclear Waste Repository



- Output Log of maximal dose of radiation in drinking water
- 27 inputs
- Training data: *n* = 190*(900)
- Validation data: $m = 69^*(300)$



Graphical Diagnostics: Individual errors





Graphical Diagnostics: Individual errors



Leo Bastos (University of Sheffield)

Diagnostics

UFRJ, December 2008

Graphical Diagnostics: Correlated errors



 $D_{MD}(\mathbf{y}^*) = 58.96$ and the 95% Cl is (47.13; 104.70)



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For interpretation of remote sensoring data

For determination of agronomical and phytometric parameters

- The Nilson-Kuusk model is a single output model with 5 inputs
- The training data contains 150 points.
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Graphical Diagnostics - Individual Errors





Graphical Diagnostics - Uncorrelated Errors



 $D_{MD}(\mathbf{y}^*) = 750.237$ and the 95% CI is (69.0, 142.6) Indicating a conflict between emulator and simulator.



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Graphical Diagnostics - Input 5





• The mean function $h(\cdot) = (1, \mathbf{x}, x_5^2, x_5^3, x_5^4)$

- Log transformation on outputs
- "new" dataset for validation



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Individual errors





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Uncorrelated Errors



 $D_{MD}(\mathbf{y}^*) = 63.873$ and the 95% CI is (32.582, 79.508)



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