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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- ocean salinity and ocean temp at different depths in the ocean
- 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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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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$$h(\mathbf{x}) = (1, \mathbf{x})^T$$

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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:

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

Mahalanobis distance

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



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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:

- $D^{PC}(\mathbf{y}^*)^T D^{PC}(\mathbf{y}^*) = D_{MD}(\mathbf{y}^*)$
- $Var(D^{FG}(n(X)) = T$
- 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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Pivoted Cholesky errors

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

Properties:

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$$D^{PC}(\mathbf{y}^*)^T D^{PC}(\mathbf{y}^*) = D_{MD}(\mathbf{y}^*)$$

- $Var(D^{PC}(\eta(\mathbf{X})) = \mathcal{I})$
- 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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- 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.
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17 / 29

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
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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.



24 / 29

Graphical Diagnostics - Input 5





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- "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)



28 / 29

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- Our diagnostics are useful tools inside the validation process.

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