

Data driven inference of model discrepancies in Zika virus dynamics

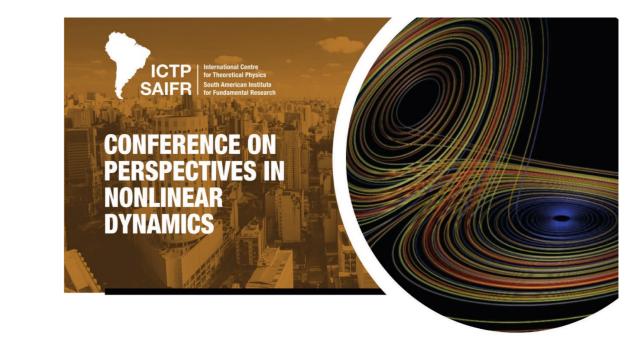
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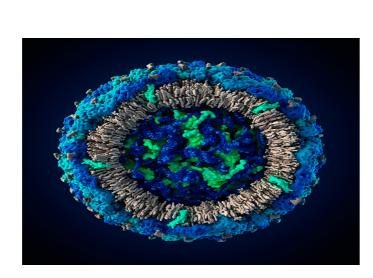
NUMERICO – Nucleus of Modeling and Experimentation with Computers

Introduction

- Zika virus: global widespread and connection with congenital diseases
- 2016: Zika becomes a public health emergency of international concern
- Main vector: Aedes mosquitoes
- A validated model can reveal new characteristics of the disease
- Prediction results are affected by model errors and lack of data
- Relations of model parameters are also of interest



Aedes aegypti



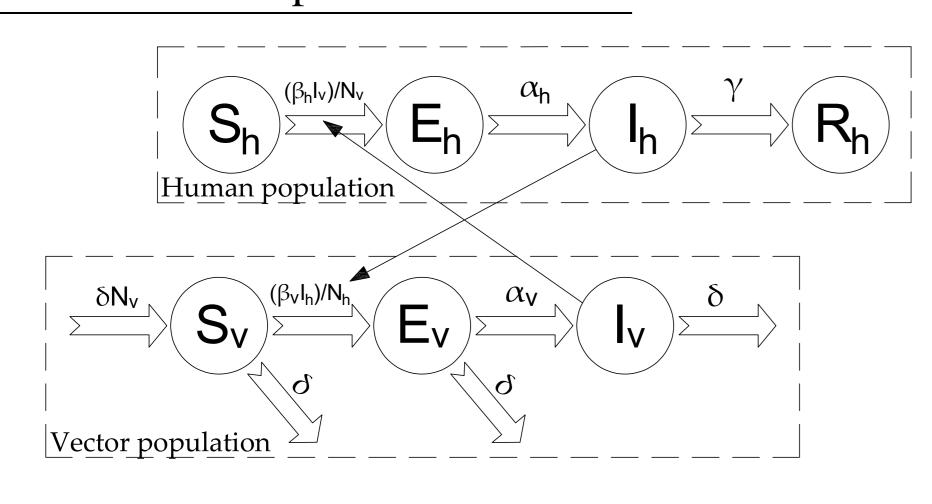
Zika virus

Objectives

- Perform sensitivity analysis to compare the parameters' global effect for a set of hierarchical models
- Improve a calibration result using a bigger data set obtained from a hierarchical superior model
- Develop a statistical framework using Bayesian Inference and Polynomial Chaos Expansion to quantify epidemic model discrepancies

Computational Model

SEIR-SEI Compartmental Model



Dynamical System

$$\frac{\mathrm{d}S_h}{\mathrm{d}t} = -\beta_h \, S_h \, \frac{I_v}{N_v} \,,$$

$$\frac{\mathrm{d}S_h}{\mathrm{d}t} = -\beta_h S_h \frac{I_v}{N_v}, \qquad \qquad \frac{\mathrm{d}S_v}{\mathrm{d}t} = \delta N_v - \beta_v S_v \frac{I_h}{N_h} - \delta S_v,$$

$$\frac{\mathrm{d}E_h}{\mathrm{d}t} = \beta_h \, S_h \, \frac{I_v}{N_v} - \alpha_h \, E$$

$$\frac{\mathrm{d}E_h}{\mathrm{d}t} = \beta_h S_h \frac{I_v}{N_v} - \alpha_h E_h , \qquad \frac{\mathrm{d}E_v}{\mathrm{d}t} = \beta_v S_v \frac{I_h}{N_h} - (\alpha_v + \delta) E_v ,$$

$$\frac{1}{\mathrm{d}t} = \alpha_h \, E_h$$

$$\frac{\mathrm{d}I_h}{\mathrm{d}t} = \alpha_h E_h - \gamma I_h , \qquad \qquad \frac{\mathrm{d}I_v}{\mathrm{d}t} = \alpha_v E_v - \delta I_v ,$$

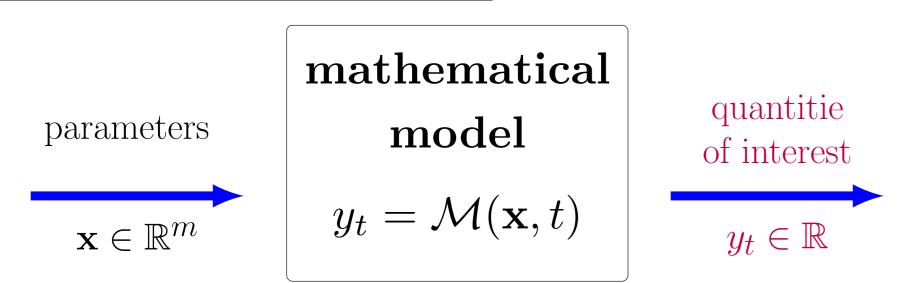
$$\frac{\mathrm{d}R_h}{\mathrm{d}t} = \gamma I_h \,,$$

$$\frac{\mathrm{d}C}{\mathrm{d}t} = \alpha_h E_h$$

+ Initial Conditions

$\frac{\mathrm{d}C}{\mathrm{d}t} = \alpha_h E_h .$

Quantities of Interest (QoI)



- Cumulative cases of infectious: $C(t) = \int_{\tau=0}^{t} \alpha_h E_h(\tau) d\tau$
- New cases per week: $\mathcal{N}_w = C_w C_{w-1}$, $w = 1 \dots 52$, $\mathcal{N}_1 = C_1$

Sensitivity Analysis

The Hoeffding-Sobol' decomposition for n iid inputs $X_i \sim \mathcal{U}(0,1)$ gives $Y_t = \mathcal{M}_0 + \sum \mathcal{M}_i(X_i) + \sum \mathcal{M}_{ij}(X_i, X_j) + \dots + \mathcal{M}_{1 \dots n}(X_1 \dots X_n),$ $1 \le i \le n$

 $\mathcal{M}_0 = \mathbb{E}[Y_t], \ \mathcal{M}_i(X_i) = \mathbb{E}[Y_t|X_i] - \mathcal{M}_0, \ \mathcal{M}_{ij}(X_i,X_j) = \mathbb{E}[Y_t|X_i,X_j] - \mathcal{M}_0 - \mathcal{M}_i - \mathcal{M}_j.$

Sobol' Indices: interaction effect of inputs in u

$$S_{\mathbf{u}} = \operatorname{Var} \left[\mathcal{M}_{\mathbf{u}}(X_{\mathbf{u}}) \right] / \operatorname{Var} \left[\mathcal{M}(\mathbf{X}) \right]$$

Metamodelling: Polynomial Chaos

The Polynomial Chaos Expansion of model $Y = \mathcal{M}(\mathbf{X})$, for a multivariate orthonormal polynomial family Φ_{α} with coefficients y_{α} ,

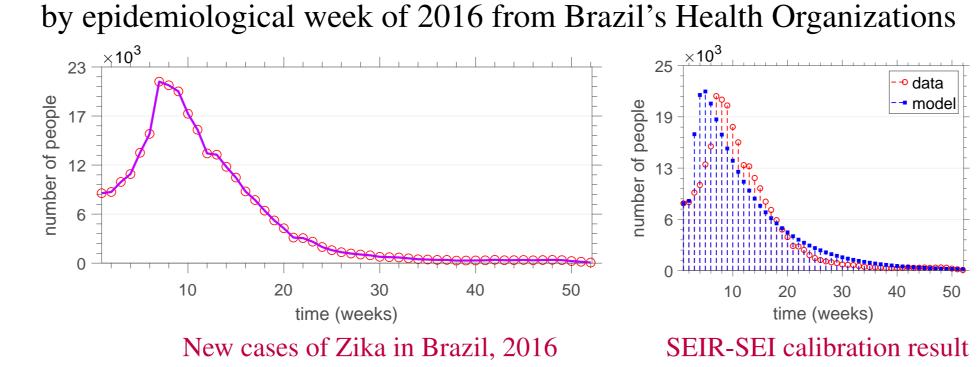
$$Y_t = \sum_{\boldsymbol{\alpha} \in \mathbb{N}^k} y_{\boldsymbol{\alpha}}(t) \, \Phi_{\boldsymbol{\alpha}}(\mathbf{X}) \,,$$

enables analytic computation of Sobol Indices:

$$S_{\mathbf{u}} = \sum_{\alpha \in \mathcal{A}_{\mathbf{u}}} y_{\alpha}^2 / \sum_{\alpha \in \mathcal{A} \setminus 0} y_{\alpha}^2, \quad \mathcal{A}_{\mathbf{u}} = \{ \alpha \in \mathcal{A} : i \in \mathbf{u} \iff \alpha_i \neq 0 \}$$

Hierarchical Calibration

⇒ Real data set: 52 values of New Cases of infectious humans for Zika



⇒ New data set: 358 values of Cumulative Cases from SEIR-SEI model

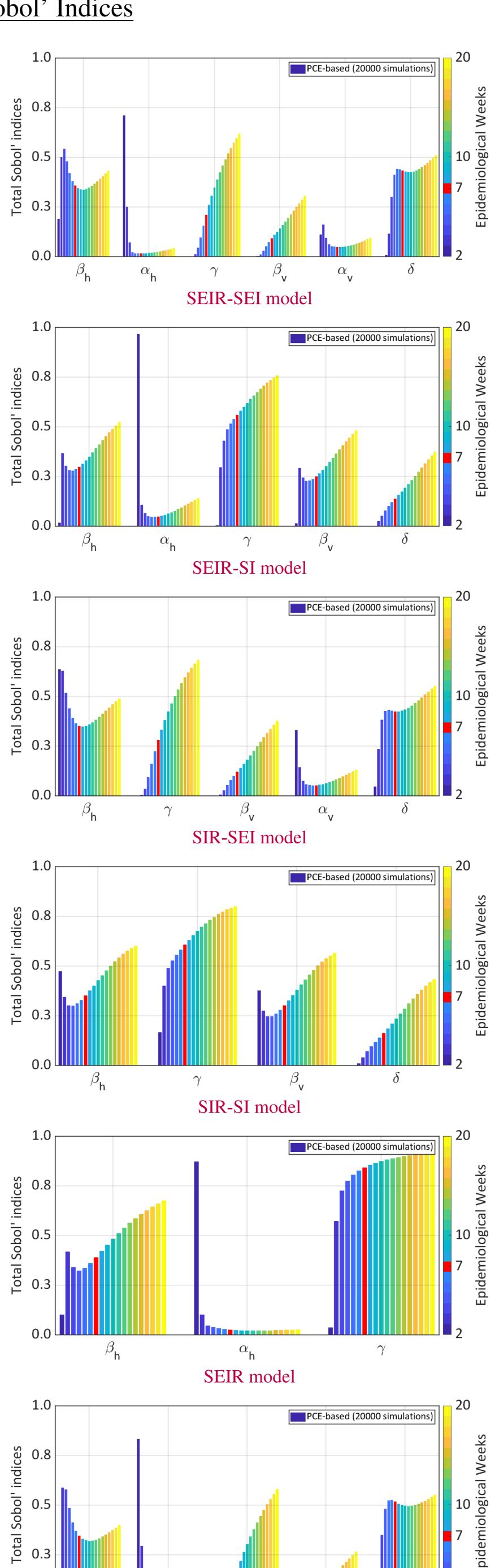
Use bigger data set from a calibrated model to improve the calibration results for other models and test some hypotheses of modeling

Models of Interest

- SIR-SEI model
- SIR-SI model
- SEIR-SI model
- SEIR model

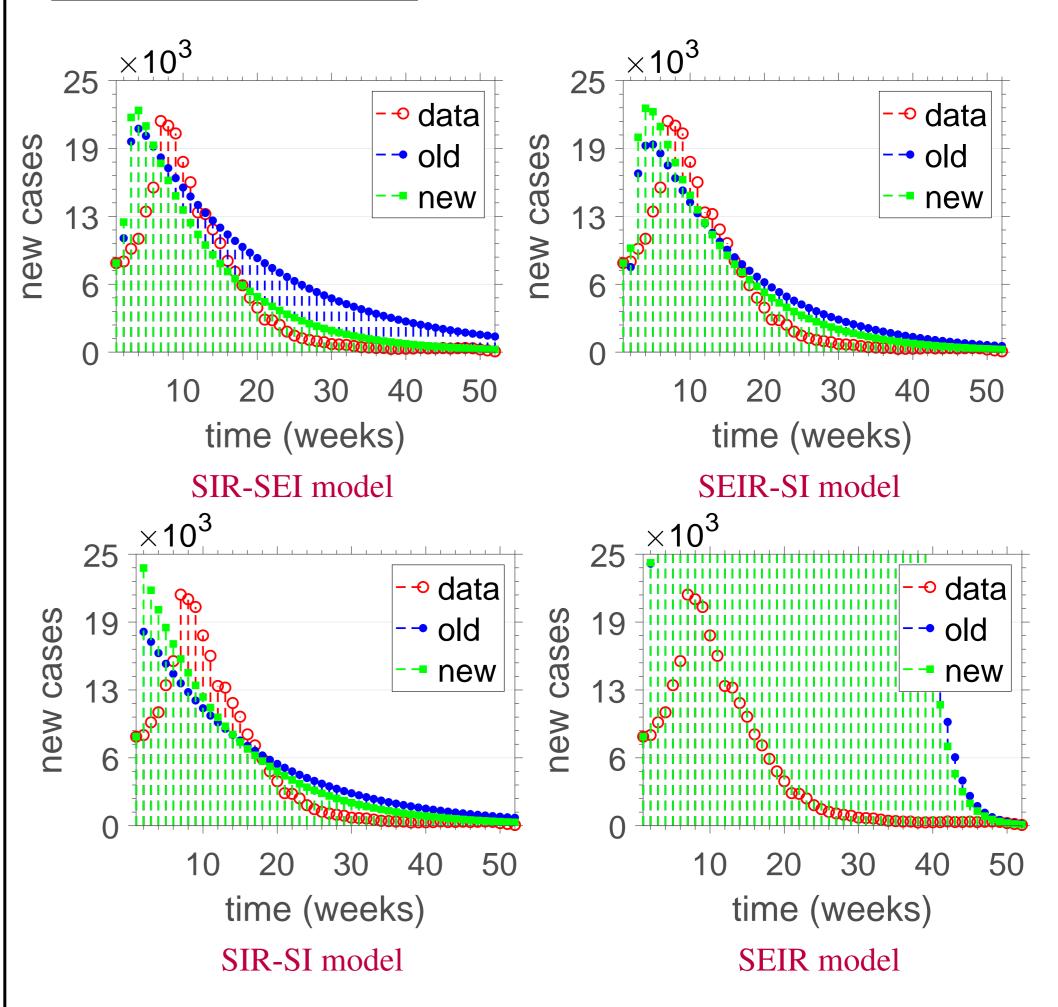
Results

Sobol' Indices



SEIR-SEI model with α_v constant

Calibration Results



Statistical Inference (ongoing research)

Discrepancy Calculation

Suppose a data set $\mathcal{D} = (t_1, y_1^{dat}), (t_2, y_2^{dat}), \dots, (t_{N_d}, y_{N_d}^{dat})$ of measures of the y_t . The *i*-th observation is given by

$$\underbrace{y_i^{dat}}_{reality} = \underbrace{\mathcal{M}(\mathbf{x}, t_i)}_{model} + \underbrace{\varepsilon_i}_{error}$$

Sargsyan, Najm and Ghanem's [4] novel approach to deal with the model discrepancies is to adopt a metamodel structure which lumps the error into the parameters

$$Y^{dat} \approx \mathcal{M}(\mathbf{X}_{\varepsilon}, t) , \ \mathbf{X}_{\epsilon} = \sum_{\alpha \in \mathcal{I}} \mathbf{X}_{\alpha}(t) \Psi_{\alpha}(\boldsymbol{\xi}),$$

where X_{α} coefficients are defined as random to be able to be identified by using Bayesian Inference

Bayesian Inference

- Inference problem become use data information to update the *prior* probability density function(PDF), defined for X_{α} . The solution corresponds posterior PDF
- From Bayes' rule,

$$\pi(\mathbf{X}_{\boldsymbol{\alpha}}|\mathcal{D}) = \frac{\pi(\mathcal{D}|\mathbf{X}_{\boldsymbol{\alpha}})\pi(\mathbf{X}_{\boldsymbol{\alpha}})}{\pi(\mathcal{D})} \,.$$

 $\rightarrow \pi(\mathbf{X}_{\alpha}|\mathcal{D})$: posterior distribution

 $\rightarrow \pi(\mathbf{X}_{\alpha})$: prior distribution

 $\rightarrow \pi(\mathcal{D}|\mathbf{X}_{\alpha})$: likelihood function

 $\rightarrow \pi(\mathcal{D})$: evidence

To define a good point of start, the Maximum Entropy Principle is applied to construct the most informative prior distribution

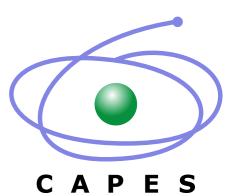
Final Remarks

- Comparative results of global Sobol' Indices show how the lack of some parameters can change the sensibility effect of the others
- With a bigger data set, the limitations in the prediction capacity of the hierarchically inferior models become more evident
- A framework for statistical inference exploring Polynomial Chaos to measure the model discrepancies was presented
- In future works, the authors intend explore this new framework to quantify model discrepancy and then improve its predictions

Acknowledgements







References

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