

Mathematical strategies in the study of epidemiological models based on nonlinear differential equations

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1 Background

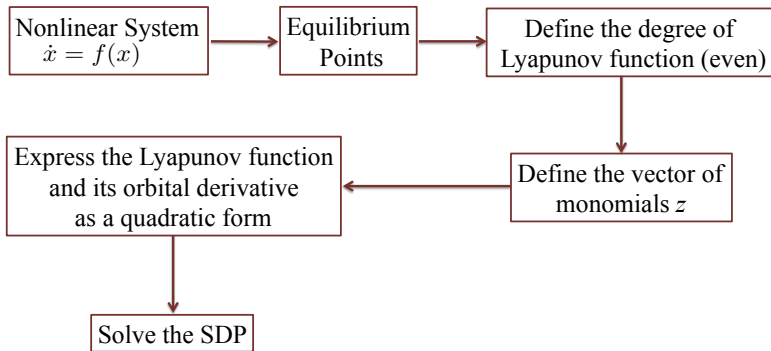
2 Find Lyapunov functions using Picard iterations

3 Control Simulations

4 Uncertainty

5 Results

Perform a simple analysis of model parameters which could be influenced by control strategies. Also we want to establish a framework to formulate the inverse problem associated to estimate interval-valued parameters by considering the uncertainty to obtain robust solutions for epidemiological models.



Theorem (Parrilo, 2000, 2003)

A multivariate polynomial $p(x)$ in n variables and of degree $2d$ is a sum of squares if and only if there exists a positive semidefinite matrix Q such that

$$p(x) = z^T Q z,$$

where z is the vector of monomials of degree up to d

$$z^T = [1, x_1, x_2, \dots, x_n, x_1 x_2, \dots, x_n^d]$$

$$\frac{ds}{dt} = \mu - \beta si - \mu s$$

$$\frac{di}{dt} = \beta si - (\gamma + \mu)i$$

$$\frac{dr}{dt} = \gamma i - \mu r$$

$$\begin{aligned} \frac{ds}{dt} &= \mu - \beta si - \mu s \\ \frac{di}{dt} &= \beta si - (\gamma + \mu)i \end{aligned} \quad (1)$$

Basic Reproductive Number R_0

$$R_0 = \frac{\beta}{\gamma + \mu}$$

Equilibrium Points

- Disease-free point,
 $E_0 = (1, 0)$
- Endemic equilibrium point,
 $E_1 = (s^*, i^*)$, where
 $s^* = \frac{1}{R_0}$, and $i^* = \frac{\mu}{\beta}(R_0 - 1)$

In general, for *sir* model we found $V(s, i) = q_{11}(s - 1)^2 + q_{22}i^2$
 where $q_{11} = \epsilon$ and $q_{22} = \frac{\epsilon(\mu + \gamma)}{(\gamma + 1)}$

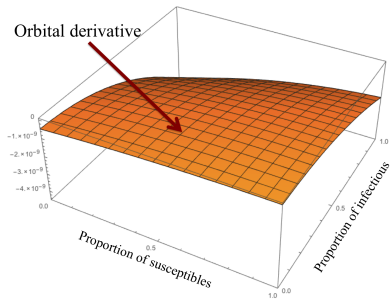
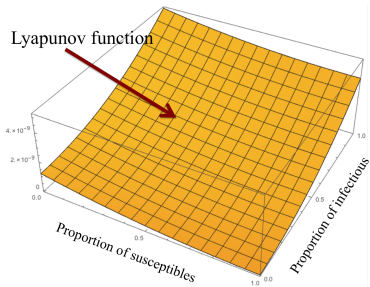


Figure: $\mu = 0.2$, $\beta = 0.5$, $\gamma = 0.8$, $R_0 = 0.5$, $q_{11} = 1.201 \times 10^{-4}$, and $q_{22} = 5.666 \times 10^{-5}$

Dengue transmission model

$$\frac{dm_e}{dt} = b\beta_m h_i (1 - m_e - m_i) - (\theta_m + \mu_m)m_e$$

$$\frac{dm_i}{dt} = \theta_m m_e - \mu_m m_i$$

$$\frac{dh_s}{dt} = \mu_h - b\beta_h m_i h_s - \mu_h h_s$$

$$\frac{dh_e}{dt} = b\beta_h m_i h_s - (\theta_h + \mu_h)h_e$$

$$\frac{dh_i}{dt} = \theta_h h_e - (\gamma_h + \mu_h)h_i$$

The disease-free point, $P_0 = (0, 0, 1, 0, 0)$.

In general, we found

$$V(m_e, m_i, h_s, h_e, h_i) = q_{11}m_e^2 + q_{22}m_i^2 + q_{33}(h_s - 1)^2 + q_{44}h_e^2 + q_{55}h_i^2$$

where

$$q_{11} = \epsilon$$

$$q_{22} = \frac{\lambda}{\sqrt{(\theta_m + \mu_m)}} + \epsilon$$

$$q_{33} \leq \frac{4\mu_h\mu_m}{b^2\beta_h^2}(q_{22} - \epsilon) + \epsilon$$

$$q_{44} \leq \frac{4\mu_m(\theta_h + \mu_h)}{b^2\beta_h^2}(q_{22} - \epsilon) + \epsilon$$

$$q_{55} \leq \frac{4(\theta_h + \mu_h)(\gamma_h + \mu_h)}{\theta_h^2}(q_{44} - \epsilon) + \epsilon$$

with $\epsilon > 0$

Theorem

(Peet and Papachristodoulou, 2012) Suppose that f is a polynomial of degree q and that system

$$\dot{x}(t) = f(x(t)), \quad x(0) = x_0 \quad (2)$$

is exponentially stable on M with

$$\|x(t)\| \leq K \|x_0\| e^{-\lambda t}$$

where M is a bounded nonempty region of radius r . Then, there exist a $\alpha, \beta, \gamma > 0$ and a sum of squares polynomial $V(x)$ such that for any $x \in M$,

$$\begin{aligned} \alpha \|x\|^2 &\leq V(x) \leq \beta \|x\|^2 \\ \nabla V(x)^T f(x) &\leq -\gamma \|x\|^2 \end{aligned} \quad (3)$$

Further, the degree of V will be less than $2q^{(Nk-1)}$, where $k(L, \lambda, K)$ is any integer such that $c(k) < K$ and

$$c(k)^2 + \frac{\log 2K^2}{2\lambda} K \frac{(TL)^k}{T} (1 + c(k))(K + c(k)) < \frac{1}{2}. \quad (4)$$

$$c(k)^2 > \frac{\lambda}{KL \log 2K^2} (1 - (2K^2)^{-\frac{1}{\lambda}}) \quad (5)$$

where $c(k)$ is defined as

$$c(k) = \sum_{i=0}^{N-1} (e^{TL} + K(TL)^k)^i K^2 (TL)^k \quad (6)$$

and $N(L, \lambda, K)$ is any integer such that $NT > (\log 2K^2/2\lambda)$ and $T < (1/2L)$ for some T and where L is a Lipschitz bound on f on B_{4Kr} .

Moving the disease-free point $E_0 = (1, 0)$ to the origin, the system (1) becomes:

$$\begin{aligned}\dot{x}_1 &= \mu - \beta(1 + x_1)x_2 - \mu(1 + x_1) \\ \dot{x}_2 &= \beta(1 + x_1)x_2 - (\mu + \gamma)x_2\end{aligned}\tag{7}$$

where $x_1 = s - 1$, and $x_2 = i$.

The Lipschitz bound for this system is given by:

$$L = \sup_{x \in B_r} \{\beta + \mu, \beta + 1, \beta, \beta + (\mu + \gamma)(1 - R_0)\}$$

To find the converse Lyapunov function we construct the Picard iteration:

$$(Pz)(t, x) = x + \int_0^t f(0) ds = x$$

$$\begin{aligned}(P^2z)(t, x) &= x + \int_0^t f((Pz)(s, x)) ds = x \\ &= x + \int_0^t f(x) ds = x + f(x)t\end{aligned}$$

The converse Lyapunov function is

$$\begin{aligned} V(x) &= \int_0^{\delta} (P^2 z(s, x))^T (P^2 z(s, x)) ds \\ &= \int_0^{\delta} (x + f(x)s)^T (x + f(x)s) ds \\ &= \int_0^{\delta} \begin{bmatrix} x \\ f(x) \end{bmatrix}^T \begin{bmatrix} I \\ sI \end{bmatrix} \begin{bmatrix} I & sI \end{bmatrix} \begin{bmatrix} x \\ f(x) \end{bmatrix} ds \\ &= \begin{bmatrix} x \\ f(x) \end{bmatrix}^T \begin{bmatrix} \delta I & \delta^2/2I \\ \delta^2/2I & \delta^3/3I \end{bmatrix} \begin{bmatrix} x \\ f(x) \end{bmatrix} \end{aligned}$$

If $\delta = \frac{1}{2L}$, for the *sir* model, we get the SOS Lyapunov function

$$\begin{aligned} 24L^3 V(x) &= \begin{bmatrix} x_1 \\ x_2 \\ f_1(x_1, x_2) \\ f_2(x_1, x_2) \end{bmatrix}^T \begin{bmatrix} 12L^2 & 0 & 3L & 0 \\ 0 & 12L^2 & 0 & 3L \\ 3L & 0 & 1 & 0 \\ 0 & 3L & 0 & 1 \end{bmatrix} \begin{bmatrix} x_1 \\ x_2 \\ f_1(x_1, x_2) \\ f_2(x_1, x_2) \end{bmatrix} \\ &= Z^T Q Z \end{aligned}$$

In this case,

$$Q = L^T L, \text{ where } L = \begin{bmatrix} 2\sqrt{3}L & 0 & \frac{3}{2\sqrt{3}} & 0 \\ 0 & 2\sqrt{3}L & 0 & \frac{3}{2\sqrt{3}} \\ 0 & 0 & \frac{1}{2} & 0 \\ 0 & 0 & 0 & \frac{1}{2} \end{bmatrix}$$

And therefore we have the sum of squares decomposition:

$$\begin{aligned}24L^3V(x_1, x_2) &= \left(\left(2\sqrt{3}L - \frac{3}{2\sqrt{3}}\mu \right) x_1 - \frac{3}{2\sqrt{3}}\beta x_2 - \frac{3}{2\sqrt{3}}\beta x_1 x_2 \right)^2 \\ &+ \left(\left(2\sqrt{3}L - \frac{3}{2\sqrt{3}}(\mu + \gamma)(1 - R_0) \right) x_2 + \frac{3}{2\sqrt{3}}\beta x_1 x_2 \right)^2 \\ &+ \frac{1}{4} (-\mu x_1 - \beta x_2 - \beta x_1 x_2)^2 \\ &+ \frac{1}{4} (\beta x_1 x_2 - (\mu + \gamma)(1 - R_0)x_2)^2\end{aligned}$$

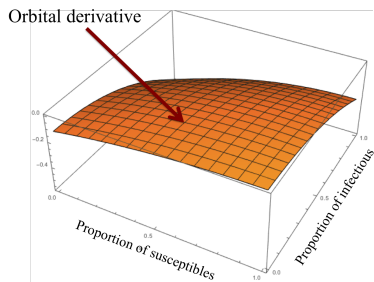
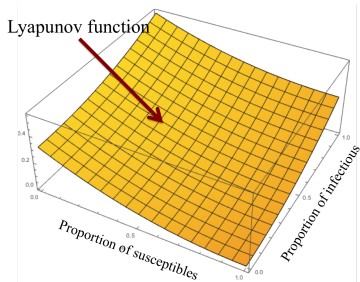


Figure: $\mu = 0.2$, $\beta = 0.5$, $\gamma = 0.8$, $R_0 = 0.5$, $L = \beta + 1 = 1.5$

If the average number of secondary infections caused by an average infective is less than one, a disease will die out, while if it exceeds one there will be an epidemic (Brauer and Castillo-Chavez, 2001).

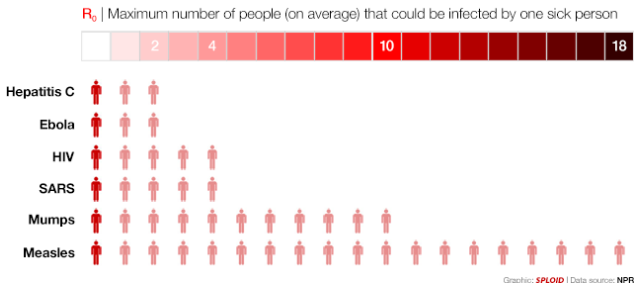


Figure: Basic reproductive number for some infectious disease. Image taken from <https://goo.gl/vDc70u>

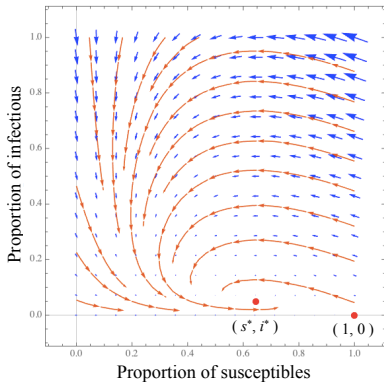
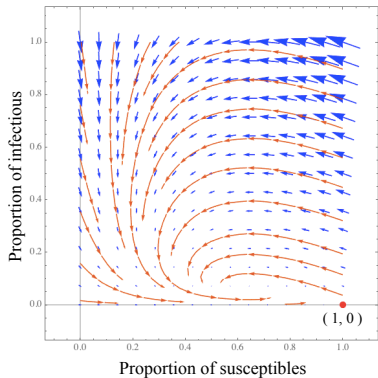


Figure: In (a) $\mu = 0.2$, $\beta = 0.5$, $\gamma = 0.8$, $R_0 = 0.5$, in (b) $\mu = 0.08$, $\beta = 0.9$, $\gamma = 0.5$, $R_0 = 1.55$

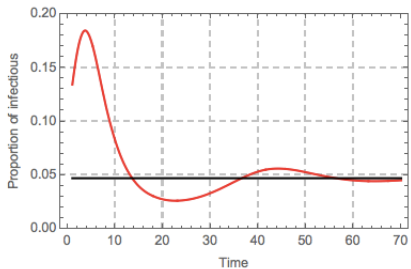
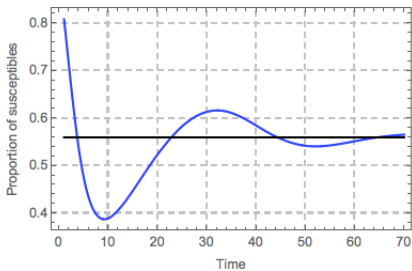


Figure: $\mu = 0.06$, $\beta = 1$, $\gamma = 0.5$, $R_0 = 1.8$, $s^* = 0.42$, and $i^* = 0.028$

For *sir* model (1), the control parameters are: μ , mortality rate.

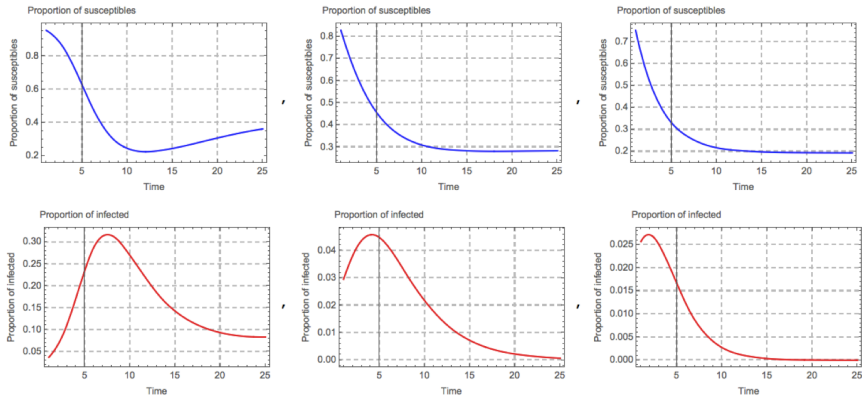


Figure: $\mu = 0.06$, $\beta = 1$, $\gamma = 0.3$, $\mu_c = 0, 0.15, 0.25$ respectively

For *sir* model (1), the control parameters are: β , transmission probability:

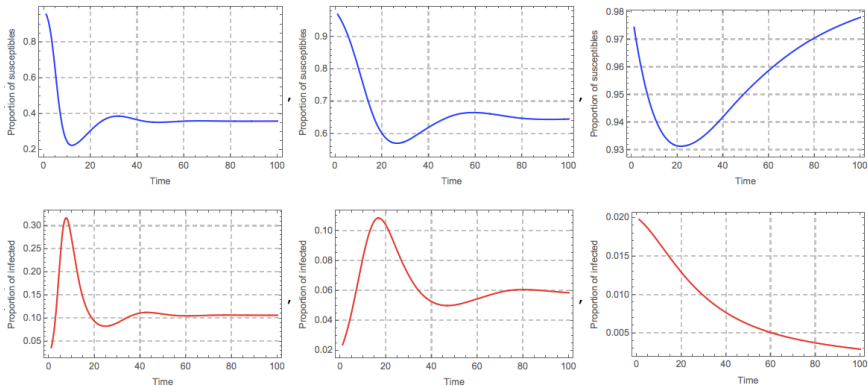


Figure: $\mu = 0.06$, $\beta = 1$, $\gamma = 0.3$, $\beta_c = 1, 0.55, 0.36$,
 $R_0 = 2.78, 1.54, 0.9$ respectively

$$\frac{dA}{dt} = \delta \left(1 - \frac{A}{C} \right) M - (\gamma_m + \mu_a) A$$

$$\frac{dM_s}{dt} = f \gamma_m A - b \beta_m \frac{H_i}{H} M_s - (\mu_m + \mu_c) M_s$$

$$\frac{dM_e}{dt} = b \beta_m \frac{H_i}{H} M_s - (\theta_m + \mu_m + \mu_c) M_e$$

$$\frac{dM_i}{dt} = \theta_m M_e - (\mu_m + \mu_c) M_i$$

$$\frac{dH_s}{dt} = \mu_h H - b \beta_h \frac{M_i}{M} H_s - \mu_h H_s$$

$$\frac{dH_e}{dt} = b \beta_h \frac{M_i}{M} H_s - (\theta_h + \mu_h) H_e$$

$$\frac{dH_i}{dt} = \theta_h H_e - (\gamma_h + \mu_h) H_i$$

$$\frac{dH_r}{dt} = \gamma_h H_i - \mu_h H_r$$

$$R_0 = \frac{b^2 \beta_m \beta_h \theta_h \theta_m}{(\theta_m + \mu_m)(\gamma_h + \mu_h)(\theta_h + \mu_h)\mu_m M} \cdot \frac{f \gamma_m}{\mu_m} \frac{\delta M C}{(\delta M + C(\gamma_m + \mu_a))}$$

$$= \frac{b^2 \beta_m \beta_h \theta_h \theta_m}{(\theta_m + \mu_m)(\gamma_h + \mu_h)(\theta_h + \mu_h)\mu_m} \cdot \frac{M_s^*}{M}$$

Control Parameters

Param.	Meaning
b	Biting rate
μ_a	Mortality rate in the aquatic phase
μ_m	Mortality rate in the adult phase
C	Carrying capacity of the environment

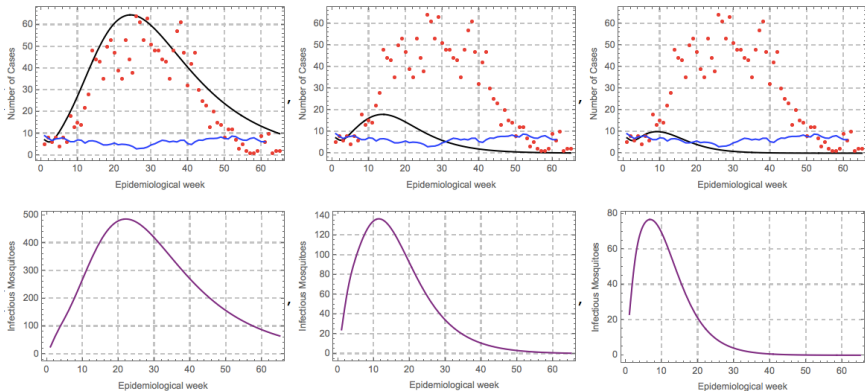


Figure: $\mu_c = 0, 0.05, 0.1$, $\delta = 65$, $\gamma_m = 1.4$, $\mu_a = 0.12$, $b = 4$,
 $\mu_m = 0.12$, $\theta_m = 0.58$, $f = 0.5$, $\theta_h = 0.7$, $C = 10000$, $\gamma_h = 1.2$,
 $\beta_m = 0.75$, $\beta_h = 0.15$, and $\mu_h = 0.0004$, and the initial conditions
 $A(0) = 9000$, $M_s(0) = 1199976$, $M_e(0) = 18$, $M_i(0) = 6$,
 $H_s(0) = 321710$, $H_e(0) = 18$, $H_i(0) = 6$, and $H_r(0) = 81501$.

Estimates of
model parameters

$$\mu = c_1$$

$$\beta = c_2$$

$$\gamma = c_3$$

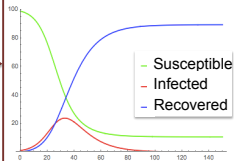
Quantitative model

$$\frac{dS}{dt} = \mu N - \beta SI - \mu S$$

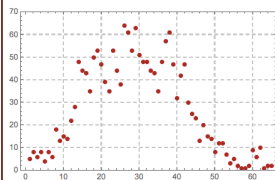
$$\frac{dI}{dt} = \beta SI - (\gamma + \mu)I$$

$$\frac{dR}{dt} = \gamma I - \mu R$$

Predictions of data



Observations of data



Quantitative model

$$\begin{aligned}\frac{dS}{dt} &= \mu N - \beta SI - \mu S \\ \frac{dI}{dt} &= \beta SI - (\gamma + \mu)I \\ \frac{dR}{dt} &= \gamma I - \mu R\end{aligned}$$

Estimates of
model parameters

$$\mu = ?$$

$$\beta = ?$$

$$\gamma = ?$$

Strategies

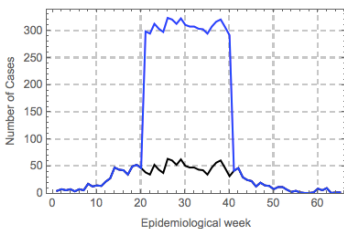
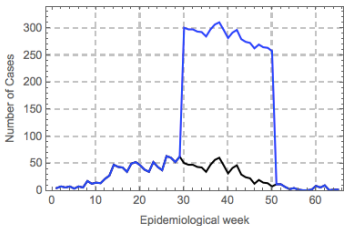
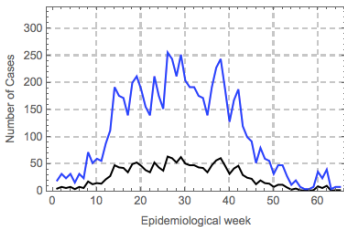
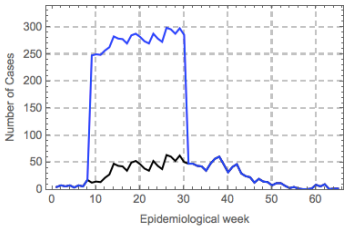
- Least squares
- Heuristic and Metaheuristic algorithms
- Monte Carlo
- Least-Squares Gradient and Hessian

Assumptions

- Independence in database
- Normal distribution
- All initial uncertainties in the problem can be modeled using Gaussian distributions (Tarantola, 2005)



Uncertainty in the dengue cases reported



— Reported Cases 2009 – 2010

— Reported Cases 2009 – 2010 considering a subreport of 75 %

Uncertainty in experimental data



Probability approximation

- Has been widely studied and applied to practical engineering problems.
- This method is based on probability distributions of the parameters with uncertainty.
- Sufficient information on the uncertainty is not always available or sometimes expensive for many practical problems.
- There are researches indicating that even a small deviation of the probability distribution is likely to cause a large error of the reliability analysis (Ben-Haim and Elishakoff, 2013).

Interval-valued approximation

- In the last two decades, the interval method in which *interval* is employed to model the uncertainty has been attracting more and more attentions (Moore, 1979; Braems et al., 2005).
- We only have to establish a bounds of the uncertainty of a parameter
- This approximation can make the uncertainty analysis more convenient and economical
- Interval method has been successfully applied to uncertainty optimization problems (Jiang et al., 2008; Gallego-Posada and Puerta-Yepes, 2017)

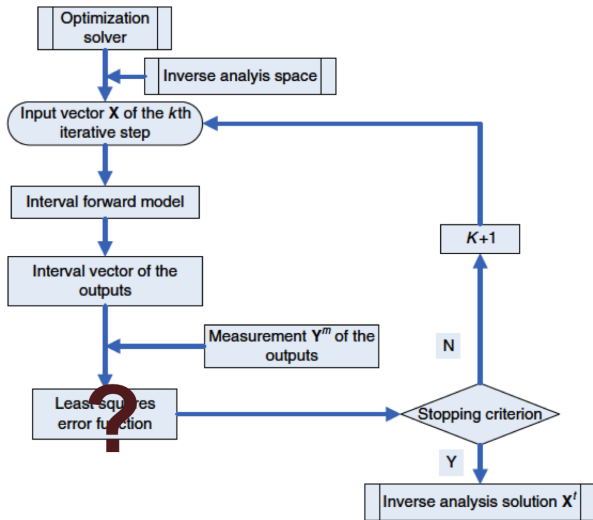


Figure: Inverse analysis process for uncertainty inverse problems. Image taken from (Jiang et al., 2008)

Without Uncertainty

$$\frac{dS}{dt} = -\beta SI$$

$$\frac{dI}{dt} = \beta SI - \gamma I$$

$$\frac{dR}{dt} = \gamma I$$

where,

$$S(0) = S_0$$

$$I(0) = I_0$$

$$R(0) = R_0$$

With Uncertainty

$$\frac{dS}{dt} = -[\beta_1, \beta_2] SI$$

$$\frac{dI}{dt} = [\beta_1, \beta_2] SI - [\gamma_1, \gamma_2] I$$

$$\frac{dR}{dt} = [\gamma_1, \gamma_2] I$$

where,

$$S(0) = [S_{0_1}, S_{0_2}]$$

$$I(0) = [I_{0_1}, I_{0_2}]$$

$$R(0) = [R_{0_1}, R_{0_2}]$$

$$\frac{dA}{dt} = [\delta_1, \delta_2] \left(1 - \frac{A}{[C_1, C_2]} \right) M - ([\gamma_{m_1}, \gamma_{m_2}] + [\mu_{a_1}, \mu_{a_2}]) A$$

$$\frac{dM_s}{dt} = [f_1, f_2][\gamma_{m_1}, \gamma_{m_2}] A - [b_1, b_2][\beta_{m_1}, \beta_{m_2}] \frac{H_i}{H} M_s - [\mu_{m_1}, \mu_{m_2}] M_s$$

$$\frac{dM_e}{dt} = [b_1, b_2][\beta_{m_1}, \beta_{m_2}] \frac{H_i}{H} M_s - ([\theta_{m_1}, \theta_{m_2}] + [\mu_{m_1}, \mu_{m_2}]) M_e$$

$$\frac{dM_i}{dt} = [\theta_{m_1}, \theta_{m_2}] M_e - [\mu_{m_1}, \mu_{m_2}] M_i$$

$$\frac{dH_s}{dt} = [\mu_{h_1}, \mu_{h_2}] H - [b_1, b_2][\beta_{h_1}, \beta_{h_2}] \frac{M_i}{M} H_s - [\mu_{h_1}, \mu_{h_2}] H_s$$

$$\frac{dH_e}{dt} = [b_1, b_2][\beta_{h_1}, \beta_{h_2}] \frac{M_i}{M} H_s - ([\theta_{h_1}, \theta_{h_2}] + [\mu_{h_1}, \mu_{h_2}]) H_e$$

$$\frac{dH_i}{dt} = [\theta_{h_1}, \theta_{h_2}] H_e - ([\gamma_{h_1}, \gamma_{h_2}] + [\mu_{h_1}, \mu_{h_2}]) H_i$$

$$\frac{dH_r}{dt} = [\gamma_{h_1}, \gamma_{h_2}] H_i - [\mu_{h_1}, \mu_{h_2}] H_r$$

Initial Conditions

$$A(0) = [A_0, A'_0]$$

$$M_s(0) = [M_{s_0}, M'_{s_0}]$$

$$M_e(0) = [M_{e_0}, M'_{e_0}]$$

$$M_i(0) = [M_{i_0}, M'_{i_0}]$$

$$H_s(0) = [H_{s_0}, H'_{s_0}]$$

$$H_e(0) = [H_{e_0}, H'_{e_0}]$$

$$H_i(0) = [H_{i_0}, H'_{i_0}]$$

$$H_r(0) = [H_{r_0}, H'_{r_0}]$$

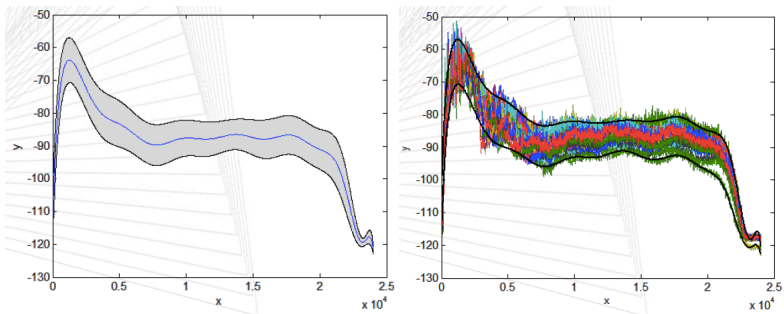


Figure: On the right, interval-valued plot of the estimated Fourier series model, and on the left, Real data vs Model output. Images taken from (Gallego-Posada and Puerta-Yepes, 2017)

We found robust Lyapunov functions to test the asymptotic stability of disease-free equilibrium points in some models simulating the transmission of mosquito-borne infectious diseases.

From the basic reproductive number R_0 it is possible determined how much should change the parameters of the model to satisfy the condition $R_0 \leq 1$

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