



Assessing model mismatch and model selection in a Bayesian uncertainty quantification analysis of a fluid- dynamics model of pulmonary blood circulation

Mihaela Paun
Glasgow University, Scotland

Jointly with ...

Dirk Husmeier



Nick Hill



Mette Olufsen



Mitchel Colebank



Glasgow University

North Carolina State University

Talk based on work presented in our recent publication:

INTERFACE

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Research



Assessing model mismatch and model selection in a Bayesian uncertainty quantification analysis of a fluid-dynamics model of pulmonary blood circulation

L. Mihaela Paun¹, Mitchel J. Colebank², Mette S. Olufsen², Nicholas A. Hill¹
and Dirk Husmeier¹

Outline

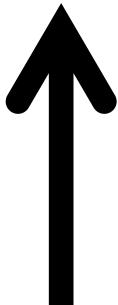
- **Background**
- **Motivation**
- **Aim**
- **Data**
- **Mathematical Model**
- **Parameter estimation**
- **Uncertainty quantification**
- **Numerical results**
- **Conclusions**

Outline

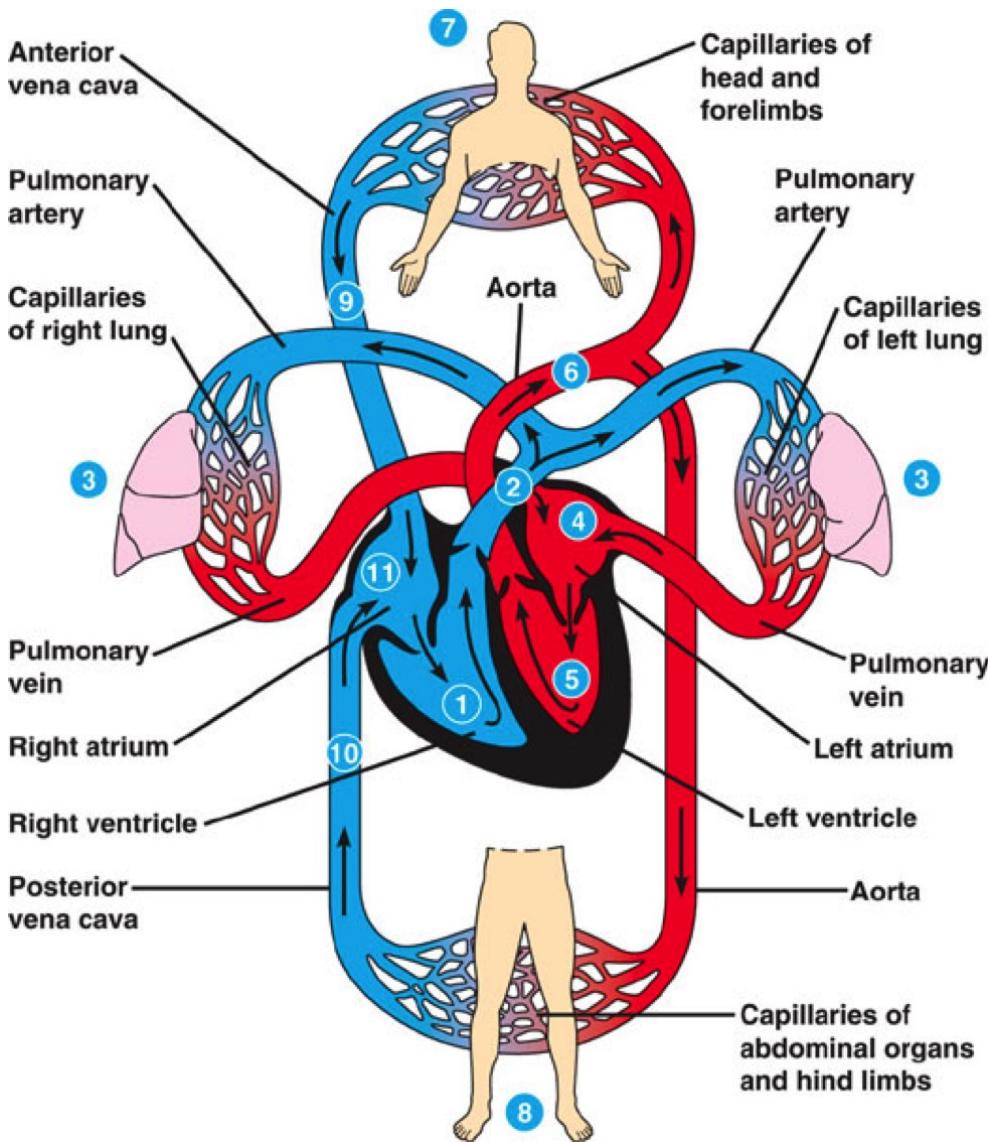
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Blood Circulation

Pulmonary circulation



Systemic circulation

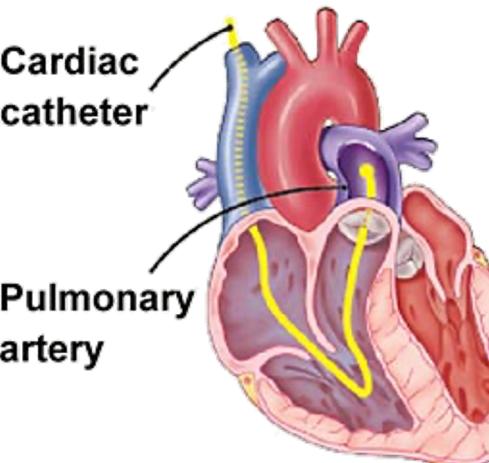
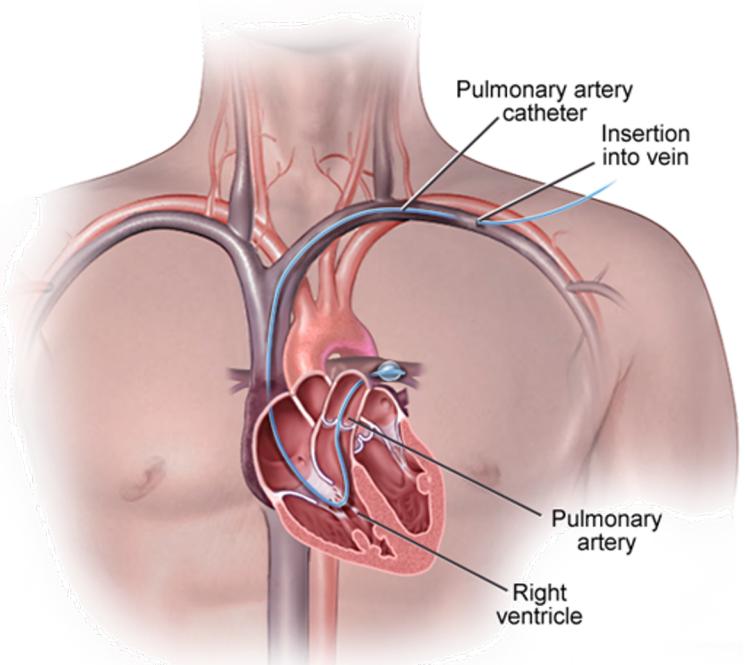
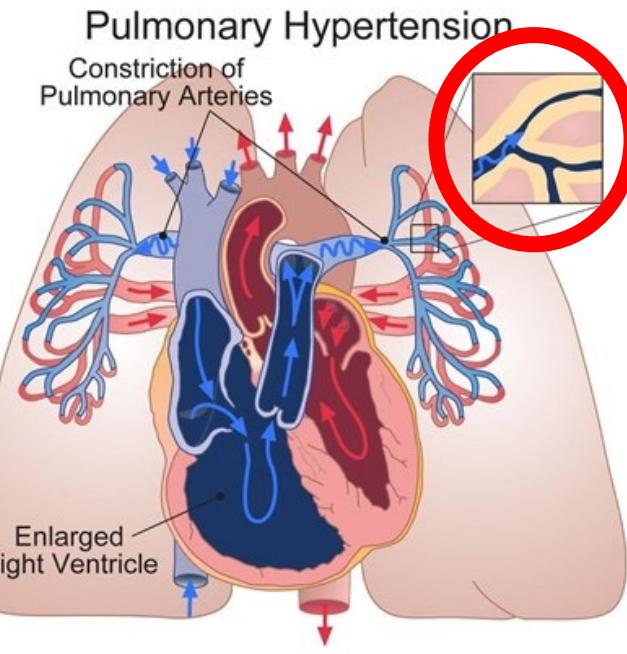
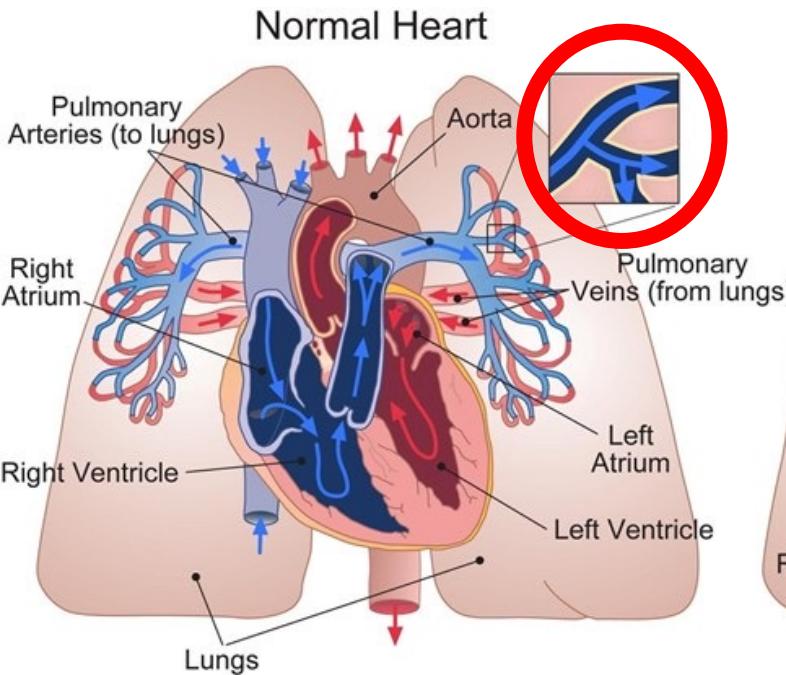
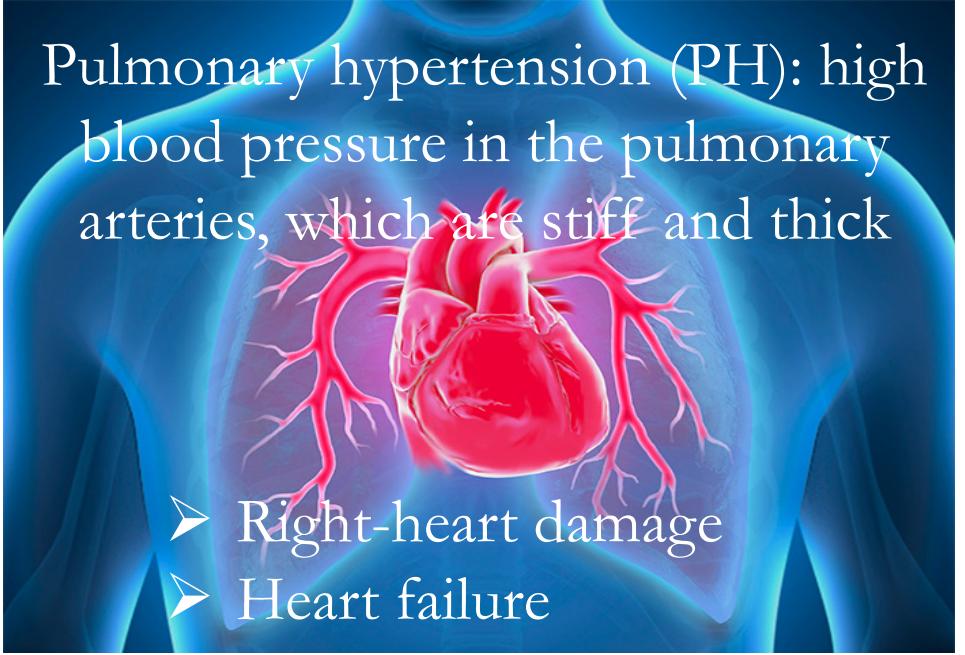


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Pulmonary hypertension (PH): high blood pressure in the pulmonary arteries, which are stiff and thick

- Right-heart damage
- Heart failure



PH: diagnosed by **invasively** measuring pulmonary pressure with right-heart catheterisation

Risks: excessive bleeding;
partial lung collapse
=> **Develop a non-invasive alternative**

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Aim

Combine

- Haemodynamic data (pulmonary blood flow – measured non-invasively with MRI)
- Imaging data (computed tomography scan giving pulmonary vessel network)
- Mathematical modelling
- Statistical modelling

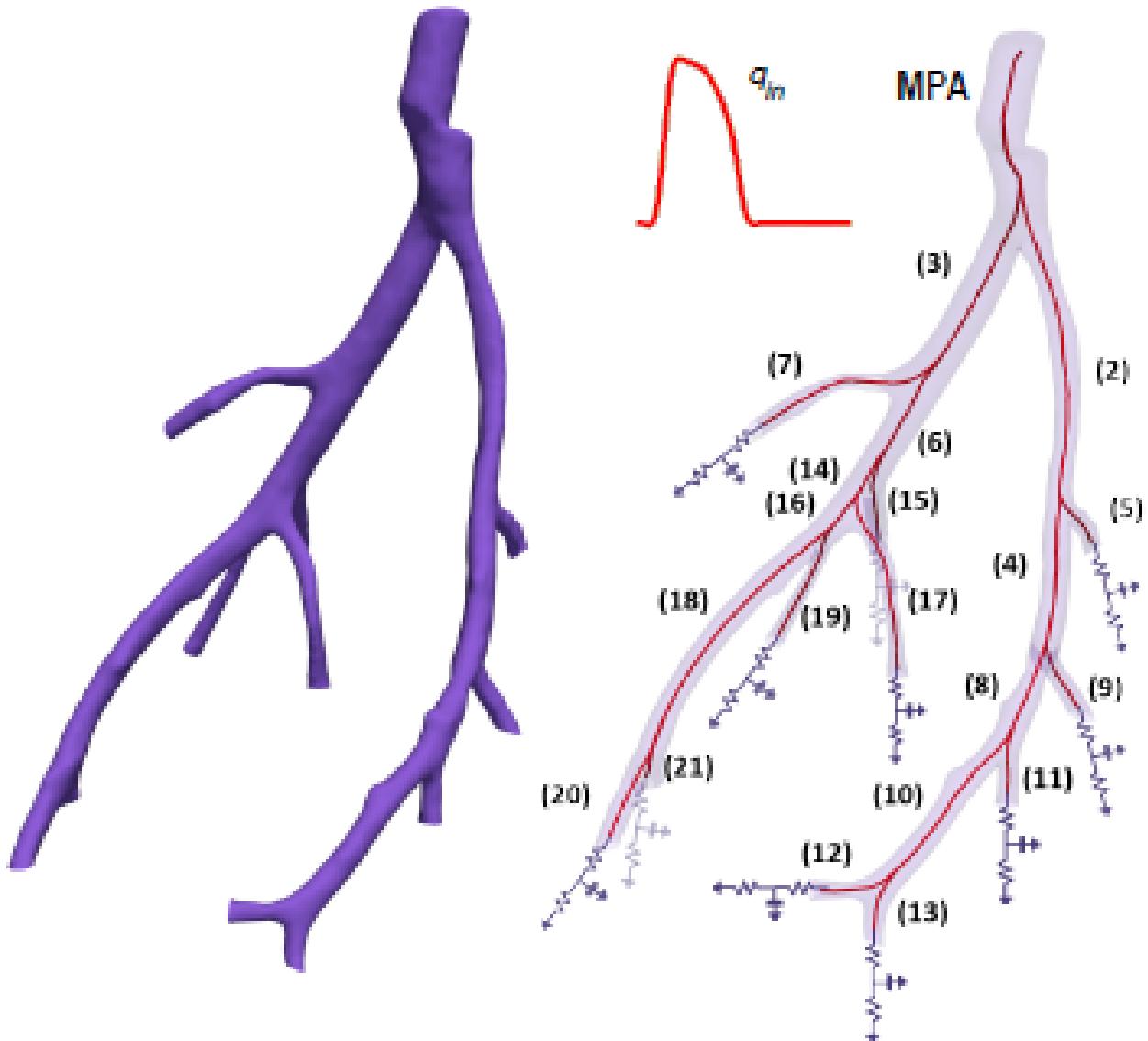
to:

- ❖ Predict pulmonary pressure (rather than measure it)
- ❖ Build a patient-specific model allowing personalised medicine

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Data



Data from a healthy mouse

Imaging data:

- Gives information on pulmonary vessel network and geometry

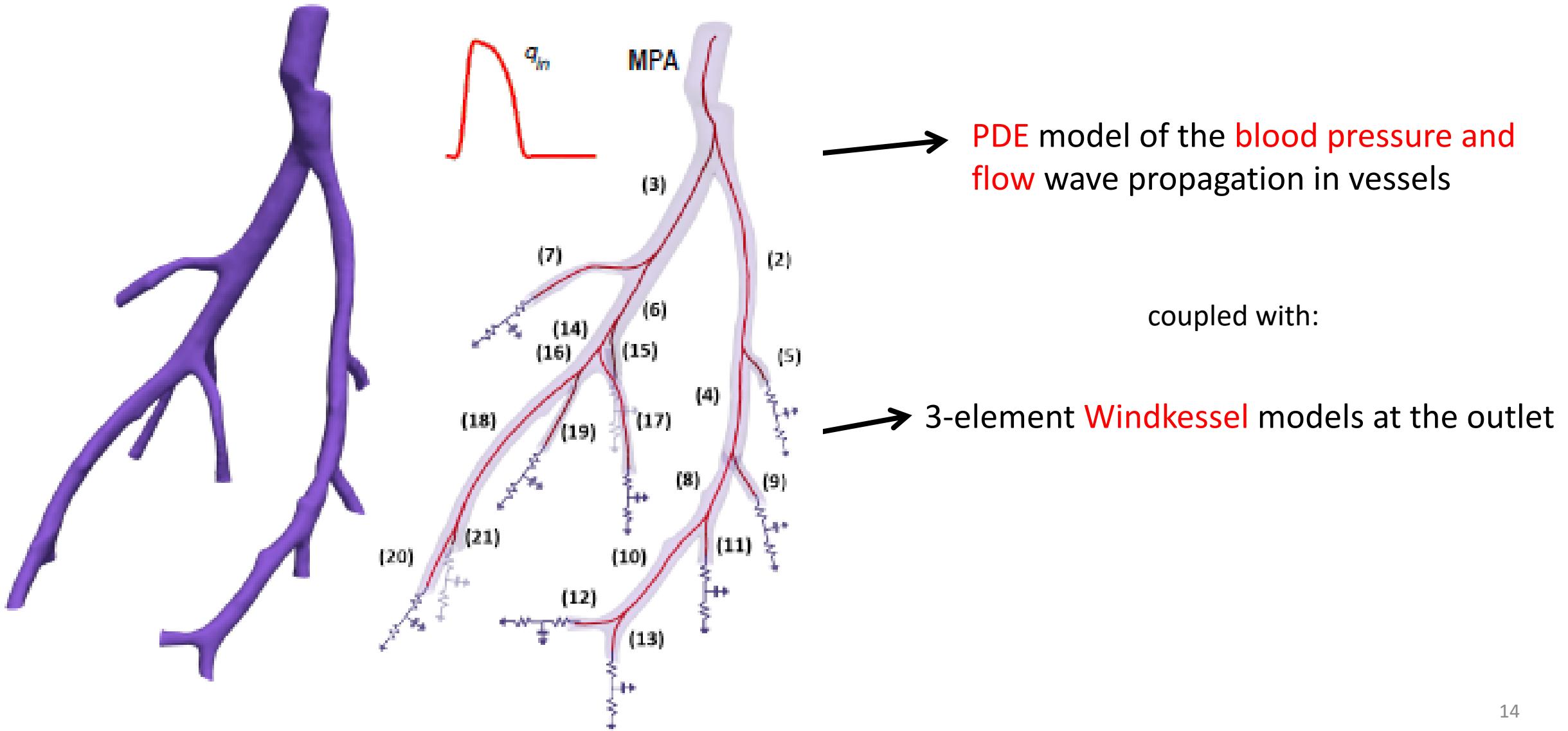
Haemodynamic data:

- Pulmonary blood pressure in the main pulmonary artery (MPA).
- Pulmonary blood flow in MPA.

Outline

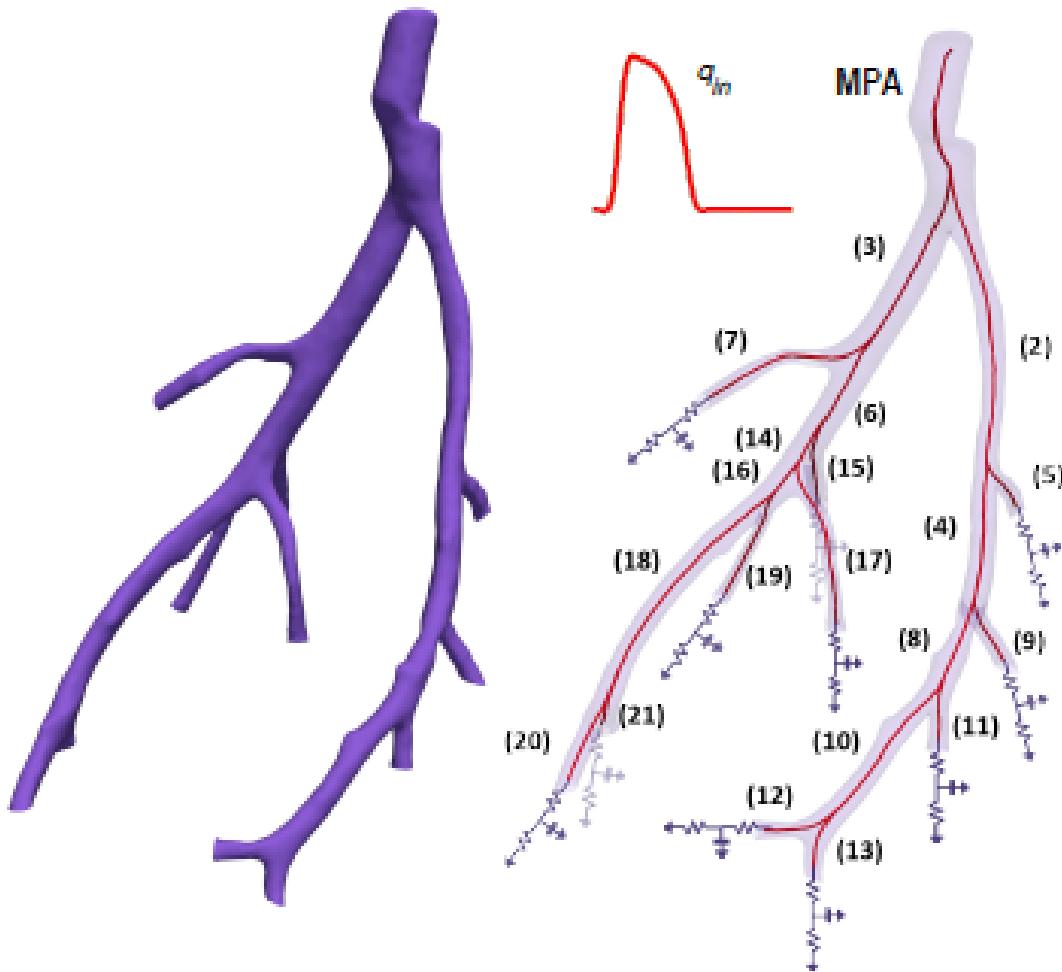
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Mathematical model



Mathematical model

Inlet boundary conditions: Flow in MPA



PDE model of blood pressure and flow

$$\frac{\partial A}{\partial t} + \frac{\partial q}{\partial x} = 0,$$
$$\frac{\partial q}{\partial t} + \frac{\partial}{\partial x} \left(\frac{q^2}{A} \right) + \frac{A}{\rho} \frac{\partial p}{\partial x} = -\frac{2\pi\mu r}{\rho\delta} q,$$

Pressure

$$p = \frac{4Eh}{3r_0} \left(\sqrt{\frac{A}{A_0}} - 1 \right)$$

Flow

Cross-sectional area

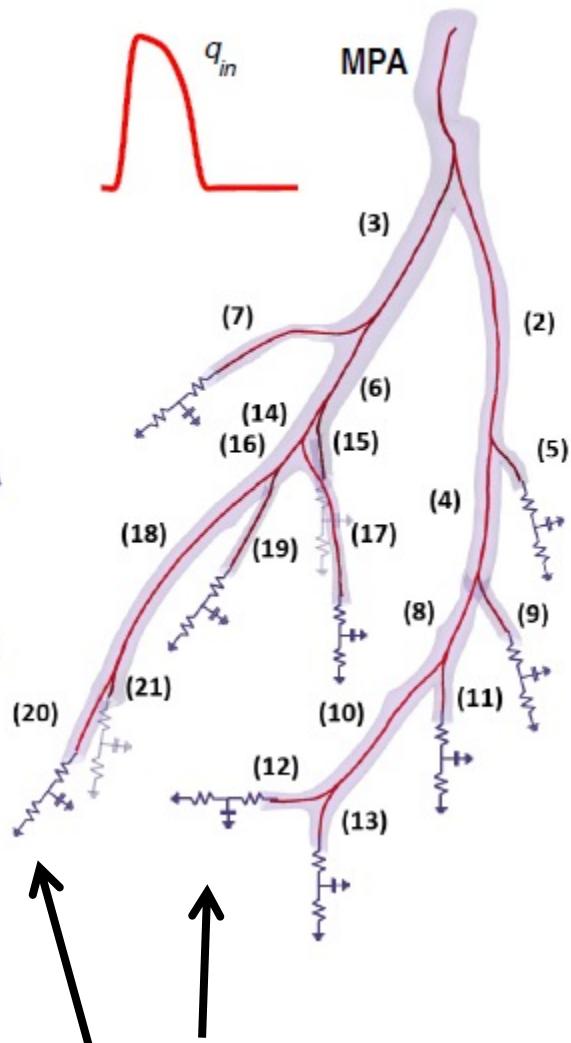
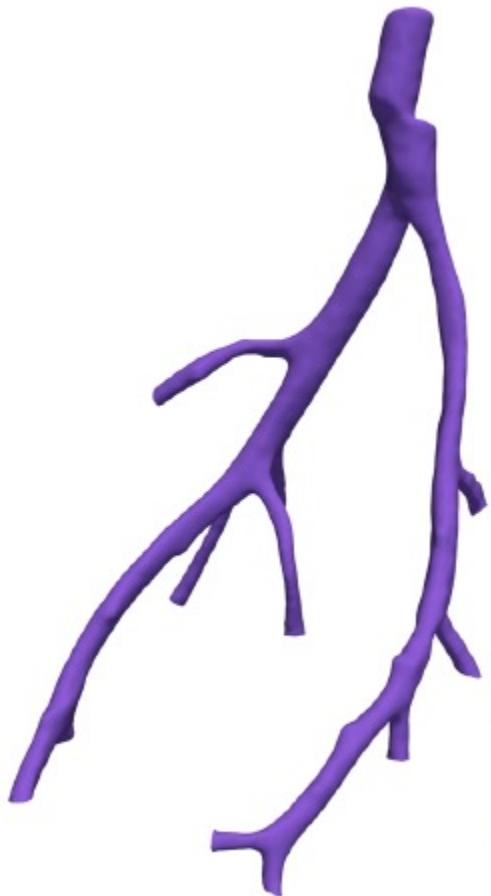
Boundary conditions at vessel junctions:

$$p_p = p_{d_i}$$

where p – parent, d_1 and d_2 - daughters

$$q_p = \sum_i q_i$$

Mathematical model



Windkessel models

3-element Windkessel model
(outlet boundary conditions)

$$\frac{dp(L,t)}{dt} - R_1 \frac{dq(L,t)}{dt}$$

$$= q(L,t) \left(\frac{R_1 + R_2}{R_2 C} \right) - \frac{p(L,t)}{R_2 C}$$

Flow

Resistances

Capacitance

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Parameter estimation

$$p = \frac{4Eh}{3r_0} \left(\sqrt{\frac{A}{A_0}} - 1 \right)$$

Unknown parameters

$$\frac{dp(L,t)}{dt} - R_1 \frac{dq(L,t)}{dt} = q(L,t) \left(\frac{R_1 + R_2}{R_2 C} \right) - \frac{p(L,t)}{R_2 C}$$

$$R_1^j = r_1 R_{01}^j, \quad R_2^j = r_2 R_{02}^j, \quad C^j = c C_0^j. \quad \text{for } j^{\text{th}} \text{ terminal vessel.}$$

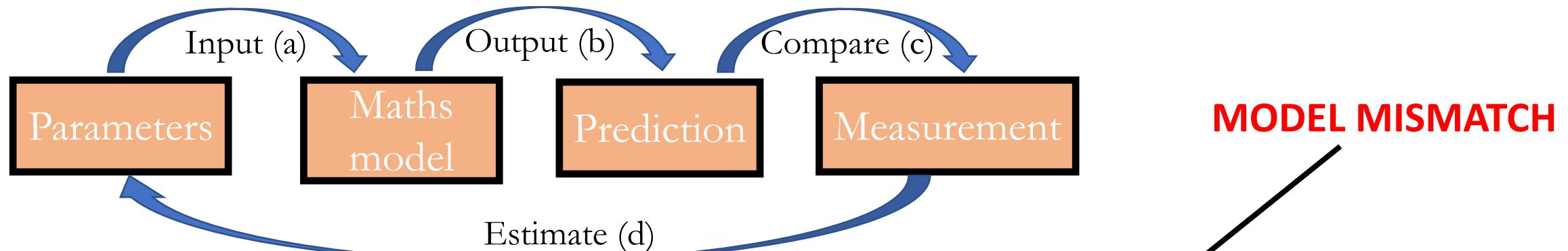
global scaling factors (common to all terminal vessels)

Estimate parameters from measured data ('**inverse problem**')

4D parameter vector:

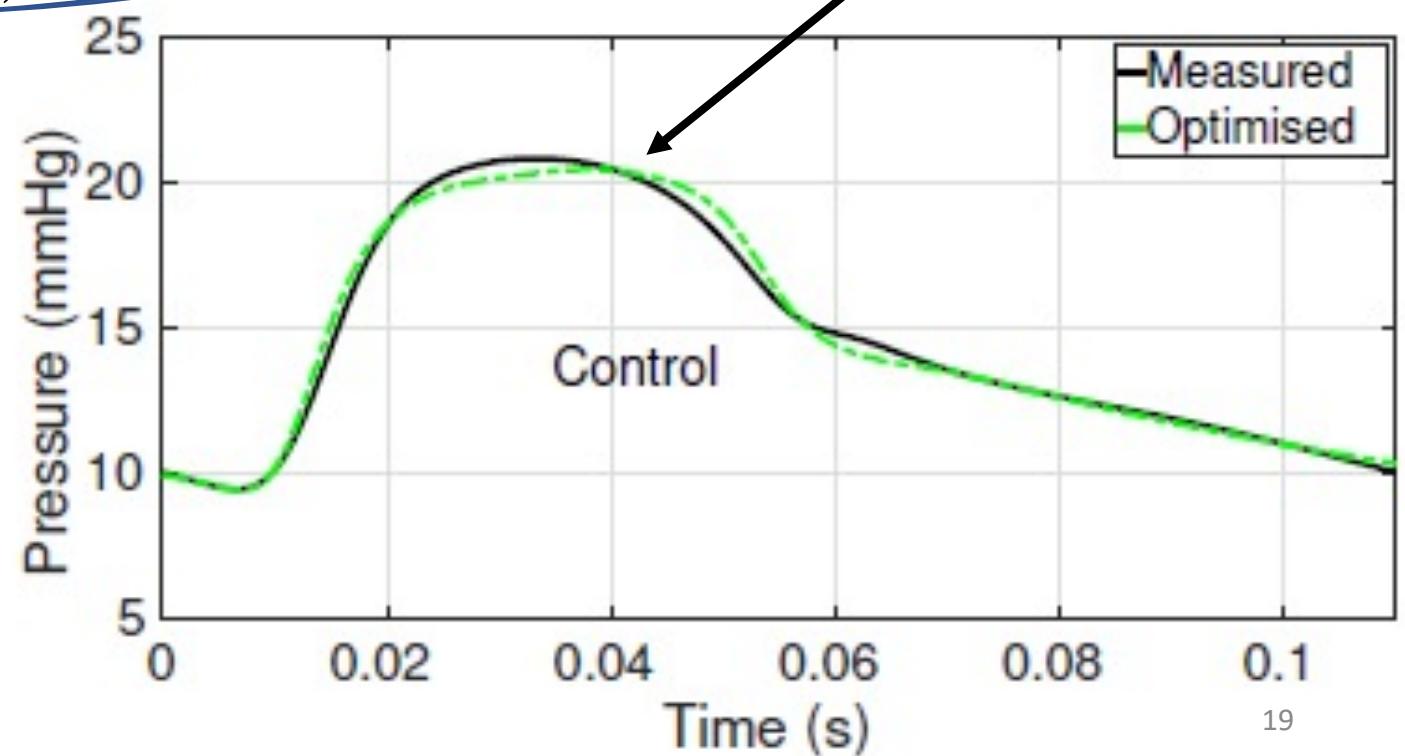
$$\frac{Eh}{r_0}, r_1, r_2, c$$

Conventional parameter estimation method



Conventional method: maximise agreement between measurement and prediction ('least squares fit')

WRONG for mis-specified maths models!



Conventional parameter estimation method

Iid errors: $\mathbf{y} \sim \mathcal{MVN}(\mathbf{m}(\boldsymbol{\theta}), \sigma^2 \mathbf{I})$, i.e.

Measurement PDE solution (prediction) Variance of measurement errors

↑ ↑ ↑

Parameters

Data likelihood

$$p(\mathbf{y}|\boldsymbol{\theta}, \sigma^2) = \left(\frac{1}{\sqrt{2\pi\sigma^2}} \right)^n \exp \left(-\frac{\sum_{i=1}^n (y_i - m_i(\boldsymbol{\theta}))^2}{2\sigma^2} \right),$$

where

Sum of squared errors (SSE)

$$\sum_{i=1}^n (y_i - m_i(\boldsymbol{\theta}))^2 = (\mathbf{y} - \mathbf{m}(\boldsymbol{\theta}))^\top (\mathbf{y} - \mathbf{m}(\boldsymbol{\theta}))$$

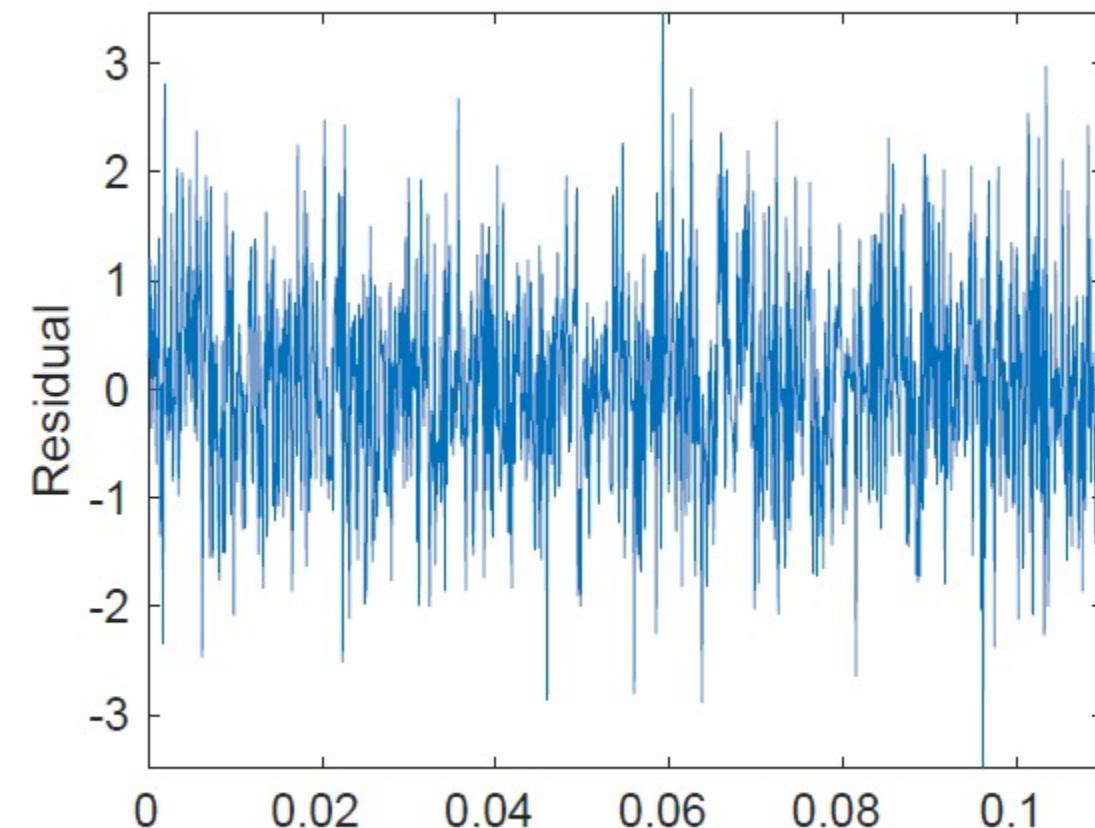
The conventional method minimises SSE
(maximises data likelihood for iid errors)

iid: independent and identically distributed
MVN: multivariate normal distribution

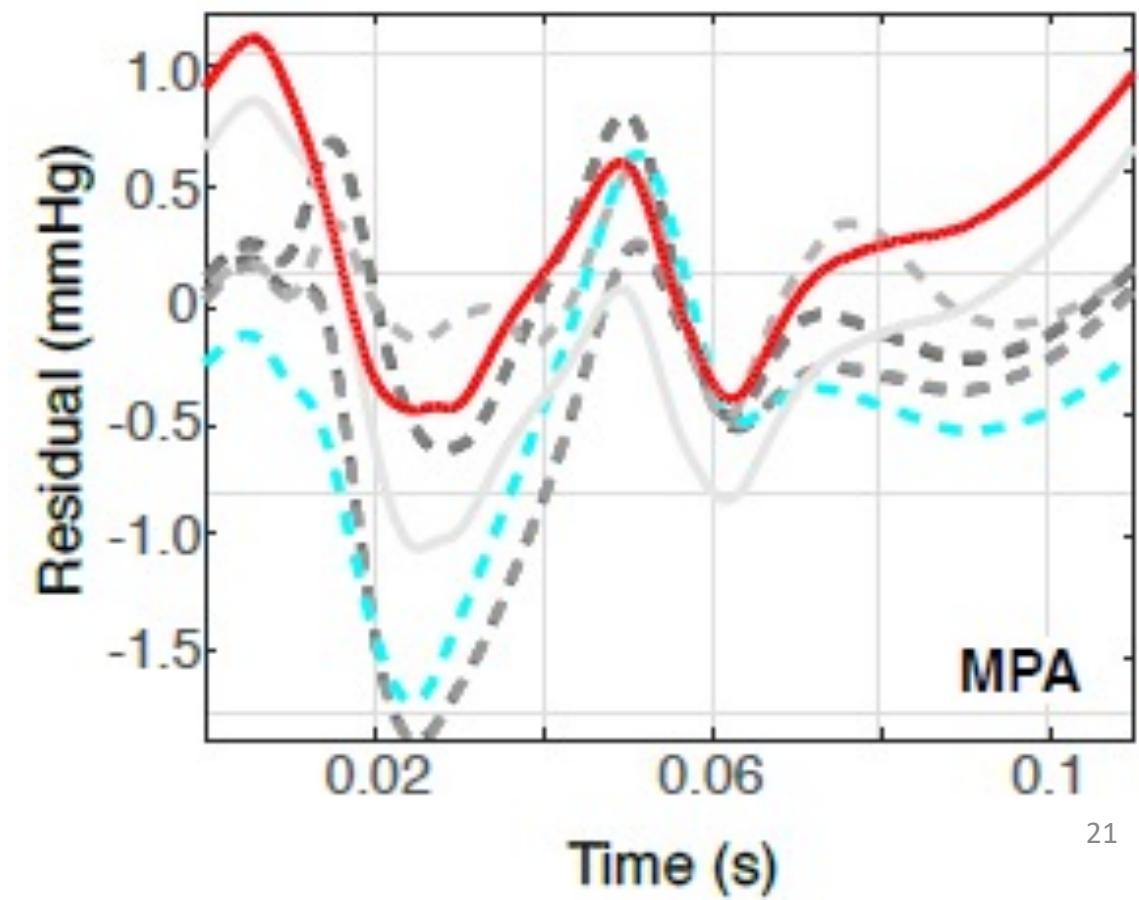
Visualising errors (residuals)

Errors: $\epsilon = y - m(\theta)$

NO MODEL MISMATCH
IID ERRORS



MODEL MISMATCH
CORRELATED ERRORS



Parameter estimation method when model mismatch is present

Correlated errors: $\mathbf{y} \sim \mathcal{MVN}(\mathbf{m}(\theta), \mathbf{C})$, i.e.

Data likelihood

$$p(\mathbf{y}|\theta, \mathbf{C}) = \det(2\pi\mathbf{C})^{-\frac{1}{2}} \exp\left(-\frac{1}{2}(\mathbf{y} - \mathbf{m}(\theta))^T \mathbf{C}^{-1} (\mathbf{y} - \mathbf{m}(\theta))\right)$$

Maximise data likelihood

Parameters that maximise data likelihood do **not** minimise SSE !

Now estimate θ and \mathbf{C}
(\mathbf{C} found using Gaussian Processes)

Modelling model mismatch

- ❖ Ignoring model mismatch (iid errors):

$$\mathbf{y}(\mathbf{t}) = \mathbf{m}(\boldsymbol{\theta}, \mathbf{t}) + \mathbf{u}(\mathbf{t}), \quad \mathbf{u}(\mathbf{t}) \sim \mathcal{MVN}(\mathbf{0}, \sigma^2 \mathbf{I})$$

- ❖ Modelling model mismatch (correlated errors):

model mismatch function

$$\begin{aligned} \mathbf{y}(\mathbf{t}) &= \mathbf{m}(\boldsymbol{\theta}, \mathbf{t}) + \underline{\mathbf{\Gamma}(\mathbf{t})} = \mathbf{m}(\boldsymbol{\theta}, \mathbf{t}) + \mathbf{f}(\mathbf{t}) + \mathbf{u}(\mathbf{t}), \\ \mathbf{f}(\mathbf{t}) &\sim \mathcal{GP}(\mathbf{0}, \mathbf{K}|\boldsymbol{\eta}), \quad \mathbf{u}(\mathbf{t}) \sim \mathcal{MVN}(\mathbf{0}, \sigma_n^2 \mathbf{I}), \end{aligned}$$

- ❖ Covariance matrix of the residuals:

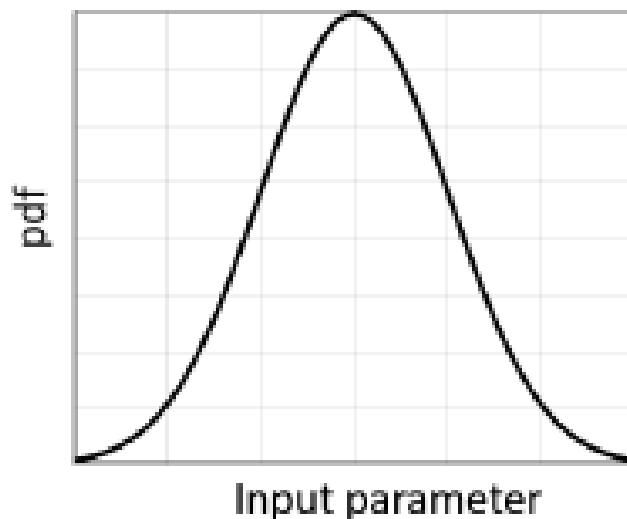
$$\mathbf{C} = \mathbf{K} + \sigma_n^2 \mathbf{I}$$

Outline

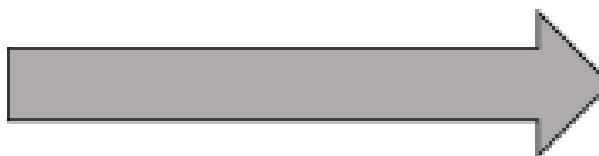
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Uncertainty quantification (UQ)

Input uncertainty

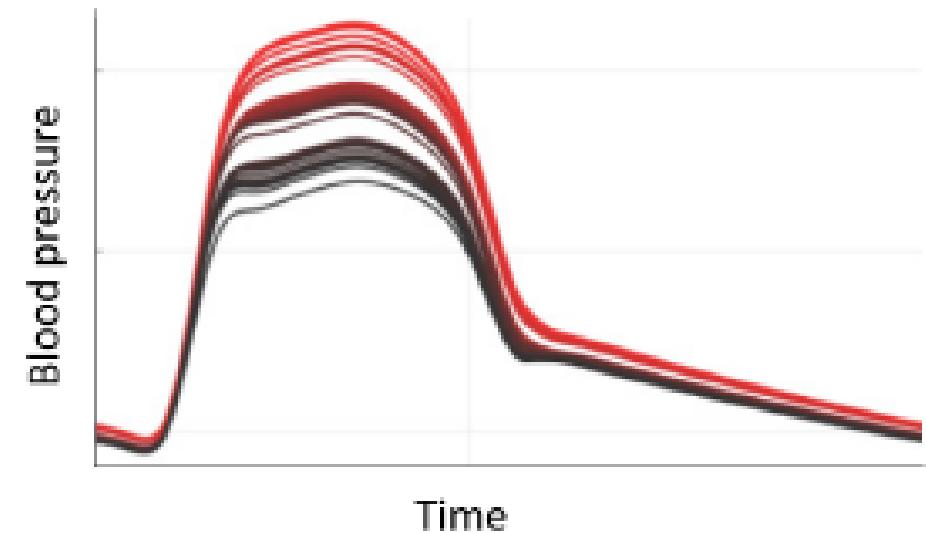


Mathematical model



Uncertainty in parameters due to: natural variability, limited and noisy data

Multi-output uncertainty



Uncertainty in parameters propagates to uncertainty in predictions

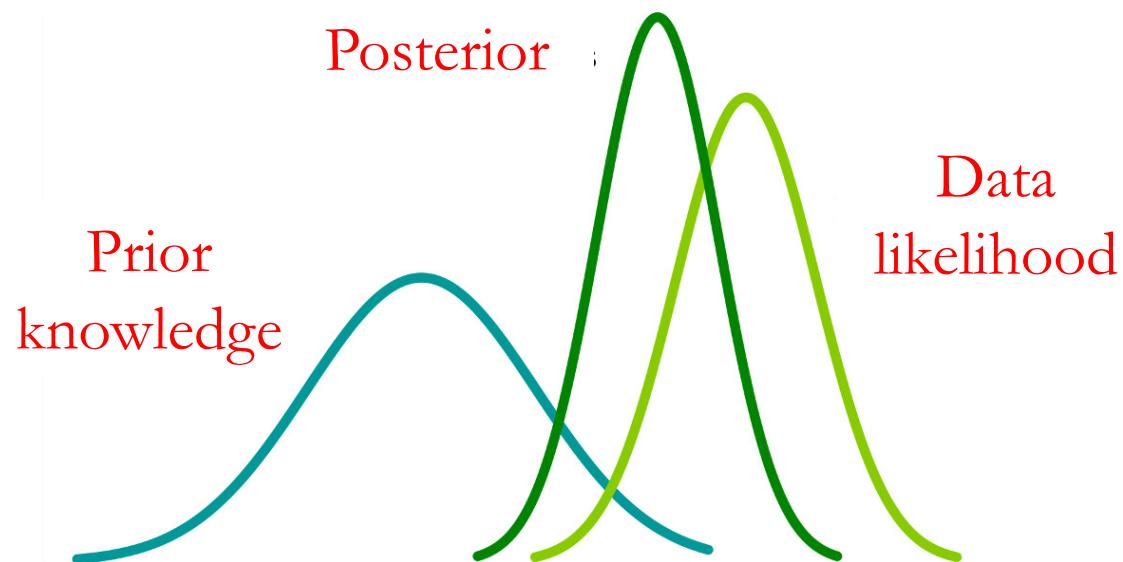
Quantify the uncertainty in parameters and predictions

Bayesian analysis

Posterior distribution Data likelihood Prior distribution

$$p(\theta, \eta | \mathbf{y}) \propto p(\mathbf{y} | \theta, \eta) p(\theta, \eta),$$

where θ are the biophysical parameters and η are the error parameters.



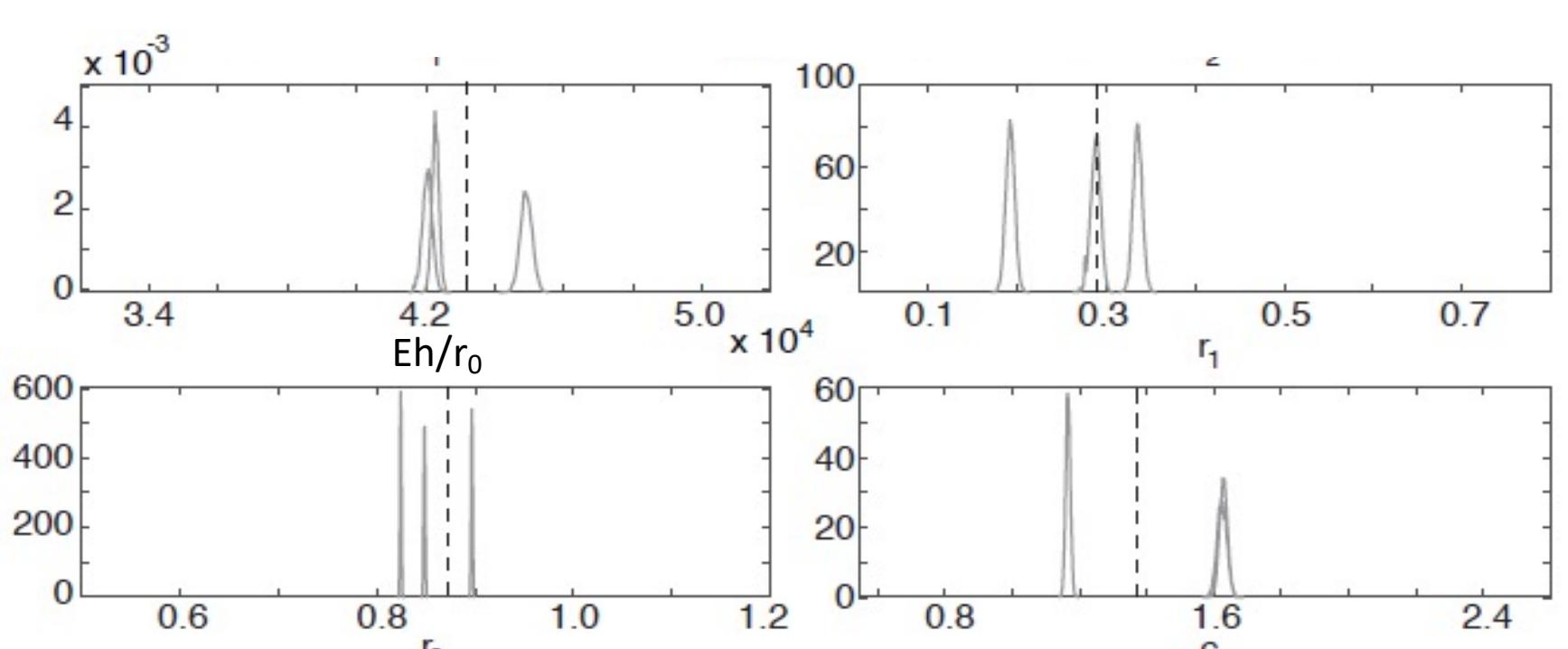
Define C , covariance matrix of residuals
(modelled with Gaussian Processes)

Idea: Sample parameter values
(approximately) from the posterior
distribution (using Markov Chain Monte
Carlo, MCMC).

Outline

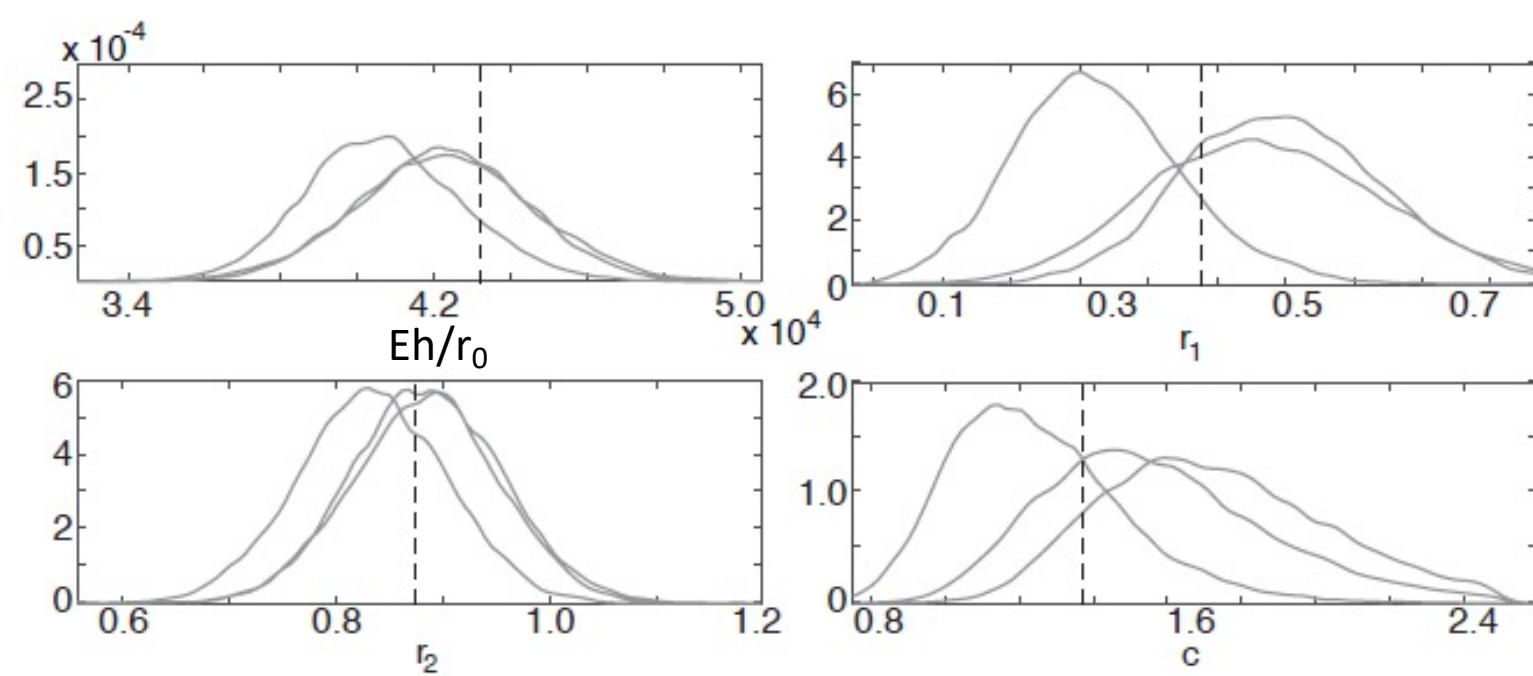
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Synthetic data
(3 data sets generated with the same ‘true’ parameter
values)



No model mismatch

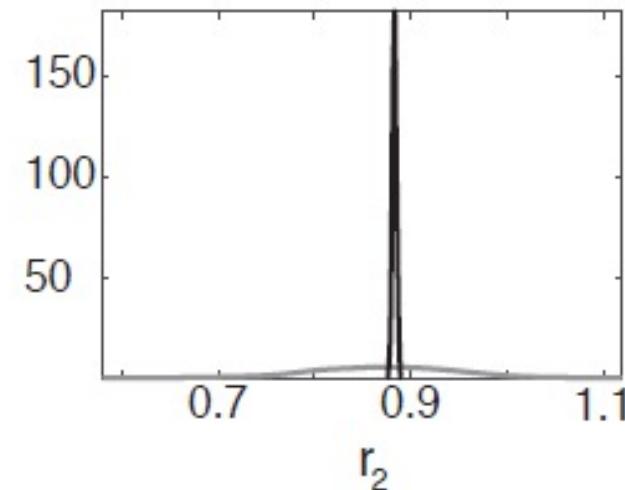
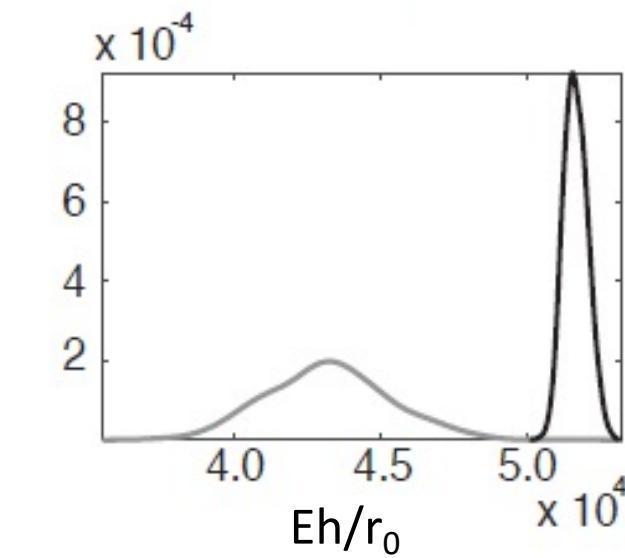
Model mismatch



Vertical dashed line: true parameter values

Measured (real) data

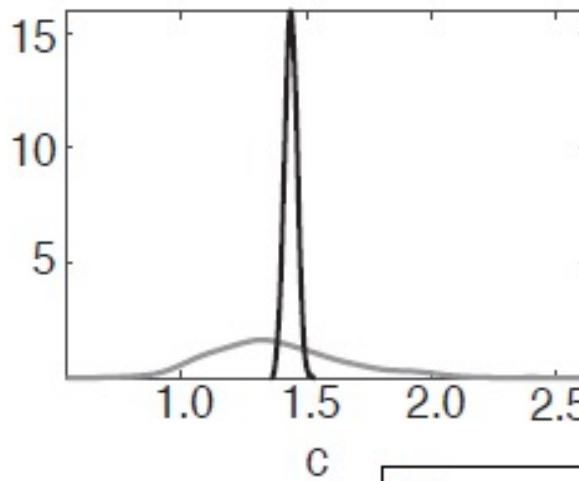
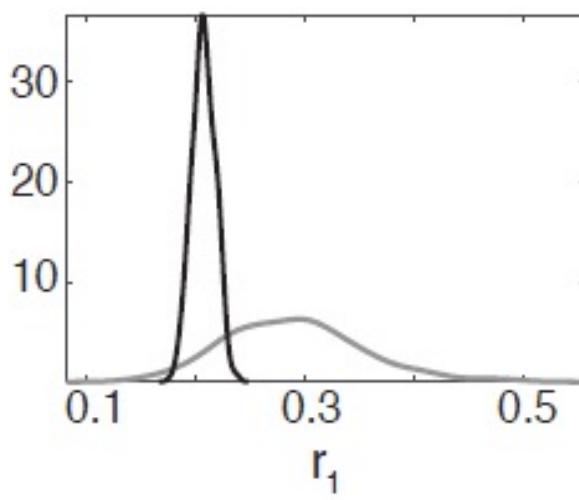
OUTPUT SPACE



C.I. = credible interval

P.I. = prediction interval

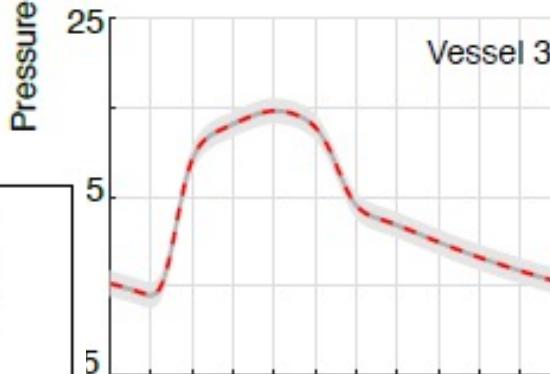
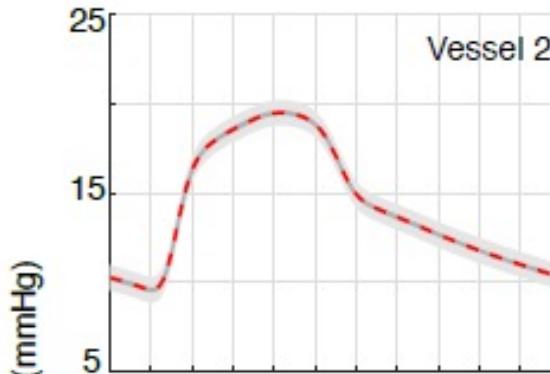
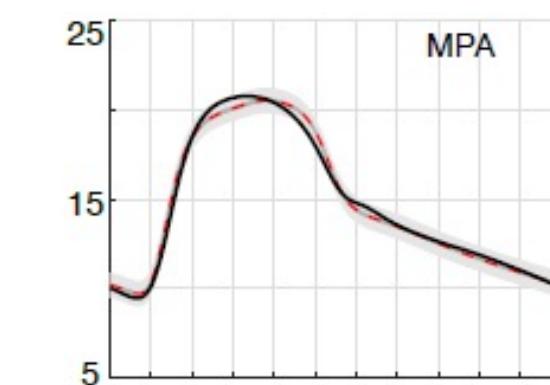
model mismatch	—
no model mismatch	—



Data	—
Median	—
95% C. I.	—
95% P. I.	—

No model mismatch

Model mismatch



Euclidean distance: 66 VS 571

Statistical model selection

- Watanabe–Akaike information criterion (WAIC) computed based on MCMC posterior samples.
- WAIC:
 - Ignoring model mismatch: 408
 - Accounting for model mismatch: -4515

Model with **lower** WAIC score is preferred.

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Conclusions

For mis-specific mathematical models, optimising the sum of squared errors ignores the model mismatch, and leads to:

- wrong parameter values and model predictions,
 - uncertainty underestimation in input and output space.
-
- These issues can be overlooked if a point-estimate analysis is used.

Thank you! ☺

Any questions?

Gaussian Processes

$$\mathbf{y}|\mathbf{f} \sim \mathcal{MVN}(\mathbf{f}, \sigma^2 \mathbf{I}),$$

$$\mathbf{f}(\mathbf{X})|\gamma \sim \mathcal{GP}(\mathbf{m}(\mathbf{X}), \mathbf{K}|\gamma),$$

$$\gamma, \sigma^2 \sim p(\gamma)p(\sigma^2),$$

$$\mathbf{C} = \begin{pmatrix} \text{Cov}(r_1, r_1) & \text{Cov}(r_1, r_2) & \text{Cov}(r_1, r_3) & \dots & \text{Cov}(r_1, r_n) \\ \text{Cov}(r_2, r_1) & \text{Cov}(r_2, r_2) & \text{Cov}(r_2, r_3) & \dots & \text{Cov}(r_2, r_n) \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ \text{Cov}(r_n, r_1) & \text{Cov}(r_n, r_2) & \text{Cov}(r_n, r_3) & \dots & \text{Cov}(r_n, r_n) \end{pmatrix}$$

$$k(\mathbf{x}, \mathbf{x}' | \gamma) = \sigma_m^2 \exp\left(-\frac{\|\mathbf{x} - \mathbf{x}'\|^2}{2l^2}\right)$$