## Inference for misspecified models

Richard Wilkinson<br>University of Sheffield



## Mechanistic models

Models describe hypothesised relationships between variables.

## Mechanistic model

- e.g. ODE/PDE models
- explains how/why the variables interact the way they do.
- parameters may have a physical meaning
- often imperfect representations of reality, but may be the only link between the quantity of interest and the data


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## e.g. Atrial fibrillation



## UQ in Patient Specific Cardiac Models

## With Sam Coveney, Richard Clayton, Steve Neiderer, Jeremy Oakley, ...

Atrial fibrillation (AF) - rapid and uncoordinated electrical activation (arrhythmia) leading to poor mechanical function.

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- Infer tissues properties, including regions of fibrotic material
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We use complex electrophysiology simulations, combine these with sparse and noisy clinical data, to

- Infer tissues properties, including regions of fibrotic material
- Predict AT pathways
- Aid clinical decision making (accounting for uncertainty) However, our simulator is imperfect. How should we proceed?


## Inference under discrepancy

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Note: Interest lies in inference of $\theta$

$$
\hat{\theta} \pm \sigma \quad \text { or } \quad \pi(\theta \mid y)
$$

not calibrated prediction:

$$
\pi\left(y^{\prime} \mid y\right)=\int F_{\theta}\left(y^{\prime}\right) \pi(\theta \mid y) \mathrm{d} \theta
$$



- $G$

[^0]
## Maximum likelihood

Maximum likelihood estimator

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\begin{aligned}
\hat{\theta}_{n} & \rightarrow \theta_{0} \text { almost surely as } n \rightarrow \infty \\
\sqrt{n}\left(\hat{\theta}_{n}-\theta_{0}\right) & \stackrel{d}{\Rightarrow} N\left(0, \mathcal{I}^{-1}\left(\theta_{0}\right)\right)
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Asymptotic consistency, efficiency, normality.

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$$
\begin{aligned}
\hat{\theta}_{n} \rightarrow \theta^{*} & =\arg \min _{\theta} D_{K L}\left(G, F_{\theta}\right) \text { a.s. } \\
& =\arg \min _{\theta} \int \log \frac{d G}{d F_{\theta}} d G \\
\sqrt{n}\left(\hat{\theta}_{n}-\theta^{*}\right) & \stackrel{d}{\Rightarrow} N\left(0, V^{-1}\right)
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## Bayes

Bayesian posterior

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Often with non-parametric models (eg GPs), we don't even get this convergence to the pseudo-true value due to lack of identifiability.

An appealing idea: model the discrepancy Kennedy an O'Hagan 2001

Can we model our way out of trouble by expanding $\mathcal{F}$ into a non-parametric world?

- Grey-box models



## An appealing idea: model the discrepancy

## Kennedy an O'Hagan 2001

Can we model our way out of trouble by expanding $\mathcal{F}$ into a non-parametric world?

- Grey-box models

One way to expand the class of models is by adding a Gaussian process (GP) to the simulator.

If $f_{\theta}(x)$ is our simulator, $y$ the observation, then perhaps we can correct $f$ using the model

$$
y=f_{\theta^{*}}(x)+\delta(x) \quad \text { where } \quad \delta(\cdot) \sim G P
$$


and jointly infer $\theta^{*}$ and $\delta(\cdot)$

## An appealing, but flawed, idea

Kennedy and O'Hagan 2001, Brynjarsdottir and O'Hagan 2014
Simulator

$$
f_{\theta}(x)=\theta x \quad g(x)=\frac{\theta x}{1+\frac{x}{a}} \quad \theta=0.65, a=20
$$

Solid=model with true theta, dashed=truth


## An appealing, but flawed, idea

Bolting on a GP can correct your predictions ${ }^{2}$, but won't necessarily fix your inference:

- No discrepancy:

$$
\begin{gathered}
y=f_{\theta}(x)+N\left(0, \sigma^{2}\right) \\
\theta \sim N(0,100), \sigma^{2} \sim \Gamma^{-1}(0.001,0.001)
\end{gathered}
$$

- GP discrepancy:

$$
\begin{aligned}
y=f_{\theta}(x) & +\delta(x)+N\left(0, \sigma^{2}\right), \\
\delta(\cdot) & \sim G P(\cdot, \cdot) \text { with objective priors }
\end{aligned}
$$



${ }^{2}$ as long as you are not extrapolating

## Dynamic discrepancy

Time structured problems give us many more opportunities to learn the model discrepancy.

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Consider the state space model:

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x_{t+1}=f_{\theta}\left(x_{t}\right)+e_{t}, \quad y_{t}=g\left(x_{t}\right)+\epsilon_{t}
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Can we correct errors in $f$ or $g$ ?

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Can we correct errors in $f$ or $g$ ? eg, $x_{t+1}=f_{\theta}\left(x_{t}\right)+\delta\left(x_{t}\right)+e_{t}$




Fitting a GP is challenging: PGAS works but is expensive, reduced rank methods better. Variational approaches (for parametric models) look promising...

## Dangers of non-parametric model extensions

There are (at least) two problems with this approach:

- We may still find $G \notin \mathcal{F}$
- Identifiability


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- A GP is an incredibly complex infinite dimensional model, which is not necessarily identified even asymptotically. The posterior can concentrate not on a point, but on some sub manifold of parameter space, and the projection of the prior on this space continues to impact the posterior even as more and more data are collected.
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ie We never forget the prior, but the prior is too complex to understand
- Brynjarsdottir and O'Hagan 2014 try to model their way out of trouble with prior information:

$$
\delta(0)=0 \quad \delta^{\prime}(x) \geq 0
$$

Great if you have this information.

## Inferential approaches

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Common approaches to inference:

- Maximum likelihood/minimum-distance
- Bayes(ish)
- History matching (HM)/ABC type methods (thresholding)


## Inferential approaches

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How do these approaches behave for well-specified and mis-specified models?

Try to understand why (at least anecdotally) HM and $A B C$ seem to work well in mis-specified cases.
Big question ${ }^{3}$ is what properties would we like our inferential approach to possess?

## ABC: approximate Bayesian computation

## Rejection Algorithm

- Draw $\theta$ from prior $\pi(\cdot)$
- Accept $\theta$ with probability $\pi(D \mid \theta)$

Accepted $\theta$ are independent draws from the posterior distribution, $\pi(\theta \mid D)$.

## ABC: approximate Bayesian computation

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Accepted $\theta$ are independent draws from the posterior distribution, $\pi(\theta \mid D)$.
If the likelihood, $\pi(D \mid \theta)$, is unknown:

## 'Mechanical' Rejection Algorithm

- Draw $\theta$ from $\pi(\cdot)$
- Simulate $X \sim f(\theta)$ from the computer model
- Accept $\theta$ if $D=X$, i.e., if computer output equals observation

The acceptance rate is $\int \mathbb{P}(D \mid \theta) \pi(\theta) \mathrm{d} \theta=\mathbb{P}(D)$.

## Rejection ABC

If $\mathbb{P}(D)$ is small (or $D$ continuous), we will rarely accept any $\theta$. Instead, there is an approximate version:

## Uniform Rejection Algorithm

- Draw $\theta$ from $\pi(\theta)$
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## Uniform Rejection Algorithm

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- Accept $\theta$ if $\rho(D, X) \leq \epsilon$
$\epsilon$ reflects the tension between computability and accuracy.
- As $\epsilon \rightarrow \infty$, we get observations from the prior, $\pi(\theta)$.
- If $\epsilon=0$, we generate observations from $\pi(\theta \mid D)$.


## $\epsilon=10$




$$
\begin{gathered}
\theta \sim U[-10,10], \quad X \sim N\left(2(\theta+2) \theta(\theta-2), 0.1+\theta^{2}\right) \\
\rho(D, X)=|D-X|, \quad D=2
\end{gathered}
$$

$$
\epsilon=7.5
$$

theta vs D


Density


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theta vs D


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$$
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$$
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## Rejection ABC

If the data are too high dimensional we never observe simulations that are 'close' to the field data - curse of dimensionality

Reduce the dimension using summary statistics, $S(D)$.
Approximate Rejection Algorithm With Summaries

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If $S$ is sufficient this is equivalent to the previous algorithm.

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Simple $\rightarrow$ Popular with non-statisticians

Handbooks of Modern Statistical Methods

> Handbook of
> Approximate Bayesian
> Computation

Edined by
Scott A. Sisson
Yanan Fan
Mark A. Beaumont

## History matching and $A B C$

History matching seeks to find a NROY set

$$
\mathcal{P}_{\theta}=\left\{\theta: S_{H M}\left(\hat{F}_{\theta, y}\right) \leq 3\right\}
$$

where

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S_{H M}\left(F_{\theta}\right)=\frac{\left|\mathbb{E}_{F_{\theta}}(Y)-y\right|}{\sqrt{\operatorname{Var}_{F_{\theta}}(Y)}}
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\pi_{\epsilon}(\theta) \propto \pi(\theta) \mathbb{E}\left(\mathbb{I}_{S\left(\hat{F}_{\theta}, y\right) \leq \epsilon}\right)
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for some choice of $S$ (typically $S\left(\hat{F}_{\theta}, y\right)=\rho\left(\eta(y), \eta\left(y^{\prime}\right)\right)$ where $\left.y^{\prime} \sim F_{\theta}\right)$ and $\epsilon$.

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They have thresholding of a score in common and are algorithmically comparable.

## History matching and $A B C$

These methods (anecdotally) seem to work better in mis-specified situations.

Why?


## History matching and $A B C$

These methods（anecdotally）seem to work better in mis－specified situations．

## Why？

They differ from likelihood based approaches in that
－They only use some aspect of the simulator output
－Typically we hand pick which simulator outputs to compare，and weight them on a case by case basis．
－Potentially use generalised scores／loss－functions
－The thresholding type nature potentially makes them somewhat conservative

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Do any of these approaches have favourable properties/characteristics for inference under discrepancy? Particularly when the discrepancy model is crude?

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- Coherence?
- Robustness to small mis-specifications?


## Generalized scores

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－A single outlier can make our inference arbitrarily bad
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$S$ is a proper score if

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i.e. predicting $G$ gives the best possibly score.

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Examples:

- Log-likelihood $S(F, y)=-\log f(y)$
- Tsallis-score $(\gamma-1) \int f(x)^{\alpha} d x-\gamma f(y)^{\alpha-1}$

Minimum scoring rule estimation (Dawid et al. 2014 etc) uses

$$
\hat{\theta}=\arg \min _{\theta} S\left(F_{\theta}, y\right)
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For proper scores

$$
\begin{aligned}
\mathbb{E}_{\theta_{0}}\left(\left.\frac{\partial}{\partial \theta} S\left(F_{\theta}, y\right)\right|_{\theta=\theta_{0}}\right) & =\left.\frac{\partial}{\partial \theta} \mathbb{E}_{\theta_{0}} S\left(F_{\theta}, y\right)\right|_{\theta=\theta_{0}} \\
& =0
\end{aligned}
$$

so we have an unbiased estimating equation, and hence get asymptotic consistency for well-specified models. We also get asymptotic normality.

Dawid et al. 2014 show that if

- $\nabla_{\theta} f_{\theta}(x)$ is bounded in $x$ for all $\theta$
- Bregman gauge of scoring rule is locally bounded then the minimum scoring rule estimator $\hat{\theta}$ is B -robust
- i.e. it has bounded influence function

$$
I F\left(x ; \hat{\theta}, F_{\theta}\right)=\lim _{\epsilon \rightarrow 0} \frac{\hat{\theta}\left(\epsilon \delta_{x}+(1-\epsilon) F_{\theta}\right)-\hat{\theta}\left(F_{\theta}\right)}{\epsilon}
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i.e. if $F_{\theta}$ is infected by outlier at $x$, this doesn't unduly affect the inference.

Note both ABC and HM are B-robust in this sense, but using the log-likelihood is not.

What type of robustness do we want here?

## Bayes like approaches

What about Bayesian like approaches with generalized scores?

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J. R. Statist. Soc. B (2016)

78, Part 5, pp. 1103-1130
A general framework for updating belief distributions
Bissiri et al. 2016 consider updating prior beliefs when parameter $\theta$ is connected to observations via a loss function $L(\theta, y)$.

## Bayes like approaches

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78, Part 5, pp. 1103-1130

## A general framework for updating belief distributions

Bissiri et al. 2016 consider updating prior beliefs when parameter $\theta$ is connected to observations via a loss function $L(\theta, y)$.
They argue the update must be of the form

$$
\pi(\theta \mid x) \propto \exp (-L(\theta, x)) \pi(\theta)
$$

via coherency arguments.
Note using log-likelihood as the loss function $\left(L(\theta, x)=-\log f_{\theta}(x)\right)$ recovers Bayes.

## Bayes like approaches

What about Bayesian like approaches with generalized scores?

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See also Jewson, Smith, Holmes 2018 who use general divergence criteria for Bayesian inference (rather than KL).

Advantages of this include

- Allows focus solely on the quantities of interest.
- Full Bayesian inference requires us to model the complete data distribution even when we're only interested in a low-dimensional summary statistic of the population.
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Relates to the Bayes linear approach of Goldstein and Wooff, which is also motivated by difficulties with specifying a complete model for the data.

## HM and ABC thresholding

History matching is an approach designed for inference for mis-specified models. Uses an implausibility measure:

$$
S_{H M}\left(F_{\theta}\right)=\frac{\left|\mathbb{E}_{F_{\theta}}(Y)-y\right|}{\sqrt{\operatorname{Var}_{F_{\theta}}(y)}}
$$

Often applied in a Bayes linear type setting, with $\operatorname{Var}_{F_{\theta}}(y)$ broken down into constituent parts

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\operatorname{Var}_{F_{\theta}}(y)=\mathbb{V a r}_{\text {sim }}+\mathbb{V a r}_{\text {discrep }}+\mathbb{V a r}_{\text {emulator }}
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Combined with the thresholding nature

$$
\mathcal{P}_{\theta}=\left\{\theta: S_{H M}\left(\hat{F}_{\theta, y}\right) \leq 3\right\}
$$

means we don't get asymptotic concentration.

- $A B C$ shares similar properties if $\epsilon$ fixed at something reasonable.

$$
\pi_{\epsilon}(\theta) \propto \pi(\theta) \mathbb{I}_{S\left(\hat{F}_{\theta}, y\right) \leq \epsilon}
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The indicator functions acts to add a ball of radius $\epsilon$ around the data, so that we only need to get within it.

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Both approaches also allow the user to focus on aspects/summaries of the simulator output that either are of interest, or for which we believe the simulator is better specified.

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Also

- Allow for crude/simple discrepancy characterization.
- Some form of robustness arises from the scores used.


## Brynjarsdottir et al. revisited

Simulator
Reality

$$
f_{\theta}(x)=\theta x \quad g(x)=\frac{\theta x}{1+\frac{x}{a}} \quad \theta=0.65, a=20
$$



Uniform MD on [-1,1]


GP prior on MD


Uniform MD on [-0.5,0.5]


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Thank you for listening!


[^0]:    ${ }^{1}$ Even if we can't agree about it!

