# New Insights into History Matching via Sequential Monte Carlo

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Denote  $\theta$  as the parameter of a deterministic model  $M(\theta)$  or stochastic model  $M(\theta, u)$  where *u* are the random numbers needed in the simulation.

Have some observed data y believed to be generated by M.

Wish to obtain a collection of plausible  $\theta$  based on *y*.

However, the model *M* is too expensive to use say Markov chain Monte Carlo or sequential Monte Carlo.

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Aim: Determine 'plausible' region of the parameter space relatively quickly.

History matching (eg Craig et al 1997) determines a non-implausible region of the parameter space relatively quickly by using an *emulator* of the model outputs or 'distance' between model outputs and *y*.

The emulator not only provides predictions at untrained points but also quantifies the uncertainty in the predictions.

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# History Matching Notation/Details

Denote  $\pi(\theta)$  as a 'prior' distribution. For simplicity we will assume there is only n = 1 model output, eg:

- The model really does have only 1 output
- The output is a distance between outputs and data.
- The output is a (approximate) likelihood function.

Parameter  $\theta$  is deemed as non-implausible if  $\mathcal{I}(\theta) < c$  for cut-off c where  $\mathcal{I}(\theta)$  includes emulation uncertainty, eg Andrianakis et al 2015

$$\mathcal{I}(\theta) = \frac{|\mathbf{y}_{\theta} - \mathbf{y}_{\text{obs}}|}{\sqrt{s_{m,\theta}^2 + s_{e,\theta}^2 + s_d^2}}.$$
 (1)

Uncertainty:  $s_{m,\theta}^2$  simulator,  $s_{e,\theta}$  emulator,  $s_d^2$  model.

## History Matching Procedure

Steps involved in the history matching algorithm:

- **1** Generate  $N_w$  training samples  $\{\theta_j\}_{j=1}^{N_w} \sim \pi(\theta)$  using a space filling design and simulate the model at each  $\theta_j$  to generate the collection of outputs  $\{y_{\theta_j}\}_{j=1}^{N_w}$ .
- 2 Fit an emulator  $E_w$  to the training data  $\{\theta_j, y_{\theta_j}\}_{j=1}^{N_w}$ .
- 3 Use the emulator  $E_w$  to define an implausibility function  $\mathcal{I}_w(\theta)$ . If  $\mathcal{I}_w(\theta) > c_w$  for some chosen  $c_w$  then  $\theta$  is deemed as implausible by emulator  $E_w$ .
- 4 Use all emulators  $\{E_r\}_{r=1}^w$  to define the non-implausible region  $\Theta_w = \{\theta \in \Theta | \cap_{r=1}^w \mathcal{I}_w(\theta) < c_w\}.$
- 5 Increase wave counter w = w + 1. Generate  $N_w$  training samples  $\{\theta_j\}_{j=1}^{N_w}$  from  $\Theta_w$  and simulate the model at each  $\theta_j$  to generate  $\{y_{\theta_j}\}_{j=1}^{N_w}$ .
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  If the stopping rule is satisfied then finish otherwise return stopping rule is satisfied then finish otherwise rule is satisfied then finish otherwi

History matching does have at least two issues:

- 1 The cut-off values  $c_w$  may not be easy to select in practice and there is no existing automated method for doing so.
- 2 Sampling uniformly from  $\Theta_w$  as *w* increases becomes increasingly difficult.

Solution: We use sequential Monte Carlo (SMC, eg Chopin 2002) to help address the above 2 issues.

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SMC samples from a sequence of distributions (connecting easy and target distributions) by iteratively applying re-weighting, re-sampling and move steps.

Advantages of SMC approach over MCMC:

- Naturally adaptive
- Easily parallelisable
- More capable of dealing with multimodal or complex posterior distributions

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## SMC History Matching

For history matching, we define the sequence of distributions as

$$p_{w}( heta) \propto \pi( heta) \prod_{k=1}^{w} \mathbb{I}(\mathcal{I}_{k}( heta) \leq c_{k}).$$

Assume that we have a collection of 'particles',  $\{W_w^i, \theta_w^i\}_{i=1}^M$  from  $p_w(\theta)$ . To push the particles to target w + 1 we apply re-weighting step:

$$W_{w+1}^i \propto W_w^i \mathbb{I}(\mathcal{I}_{w+1}(\theta_w^i) \leq c_{w+1}).$$

Here  $W_w$  are all equal, so the weights  $W_{w+1}$  will either be constant or equal to zero. Thus we can select  $c_{w+1}$  so that a certain proportion,  $\alpha$ , have non-zero weight. Same as ensuring that the effective sample size (ESS) =  $1/\sum_{i=1}^{M} (W_{w+1}^i)^2$ , is  $M_{acense}$ .

After re-weighting the ESS drops to roughly  $\alpha M$ .

Resampling M times from the surviving particles allows the ESS to return to M. However, the drawback is that some particles will be duplicated.

Diversify particles with an MCMC kernel *R* times to each of the resampled particles. Can adaptively choose *R*. Can use population of particles to inform efficient MCMC proposal. This move step only uses the emulator (no expensive model simulations).

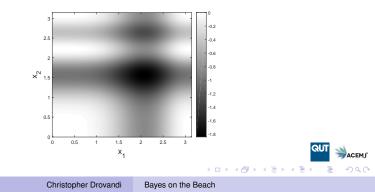
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### Simple Example

Consider the function

$$y = -\sin(x_1)\sin(x_1^2/\pi)^2 - \sin(x_2)\sin(2x_2^2/\pi)^2$$

where  $\theta = (x_1, x_2) \in (0, \pi) \times (0, \pi)$ . We wish to find the regions where *y* is small.



We use N = 50 training samples at each wave.

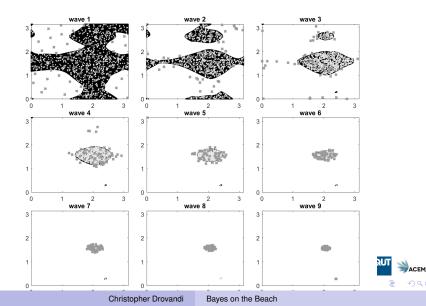
Implausability measure  $\mathcal{I}(\theta) = y_p(\theta) - r \times s_p(\theta)$  where  $y_p(\theta)$  and  $s_p(\theta)$  is the prediction and the standard deviation from the currently fitted GP, respectively. Set r = 3.

We solve the history matching problem 'perfectly' by taking  $2^{20}$  draws from parameter space and use rejection sampling. The cut-offs  $c_w$  are chosen so that half the remaining particles survive at each wave.

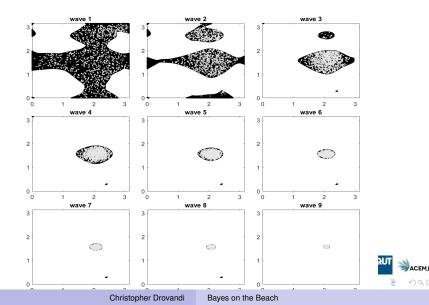
We investigate how well SMC can uniformly sample from non-implausible space implied by 'perfect' history matching.

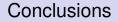
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#### **Results from SMC History Matching**



### **Results from Typical History Matching Approach**



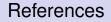


Our algorithm reveals the significant complexity of the non-implausible probability distributions arising from history matching.

The SMC algorithm provides a semi-automated and principled method for performing history matching.

More work needs to be done on designing efficient MCMC kernels when derivative information about the target distribution is not available.

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