

# IDM workshop: emulation and history matching Part 1: General Principles

Michael Goldstein, Ian Vernon\*

\*Thanks to MRc, for funding for example in presentation.



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The science in each of these applications is completely different. However, the underlying methodology for handling uncertainty is the same.



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Twenty behavioural and two epidemiologic inputs were varied for this study.



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The run time for a single simulation for the study varies between 10 minutes and 3 hours.

#### **Example: references**



Full details of example are in the paper:

Ioannis Andrianakis, Ian R. Vernon, Nicky McCreesh, Trevelyan J. McKinley, Jeremy E. Oakley, Rebecca N. Nsubuga, Michael Goldstein, Richard G. White (2015) Bayesian History Matching of Complex Infectious Disease Models Using Emulation: A Tutorial and a Case Study on HIV in Uganda, PLOS Computational Biology.

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(2015) Bayesian History Matching of Complex Infectious Disease Models Using Emulation: A Tutorial and a Case Study on HIV in Uganda,

PLOS Computational Biology.

More careful and detailed treatment in

Ioannis Andrianakis, Ian R. Vernon, Nicky McCreesh, Trevelyan J. McKinley, Jeremy E. Oakley, Rebecca N. Nsubuga, Michael Goldstein, Richard G. White

(2017) Efficient history matching of a high dimensional individual based HIV transmission model"

to appear in SIAM/ASA Journal on Uncertainty Quantification.

which applies a development of the same ideas to a much larger version of the model (96 inputs, 50 outputs).



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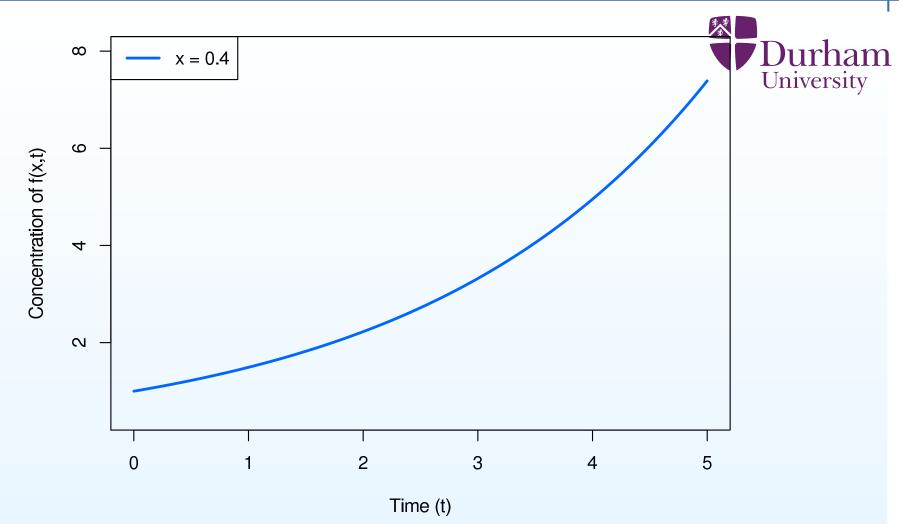
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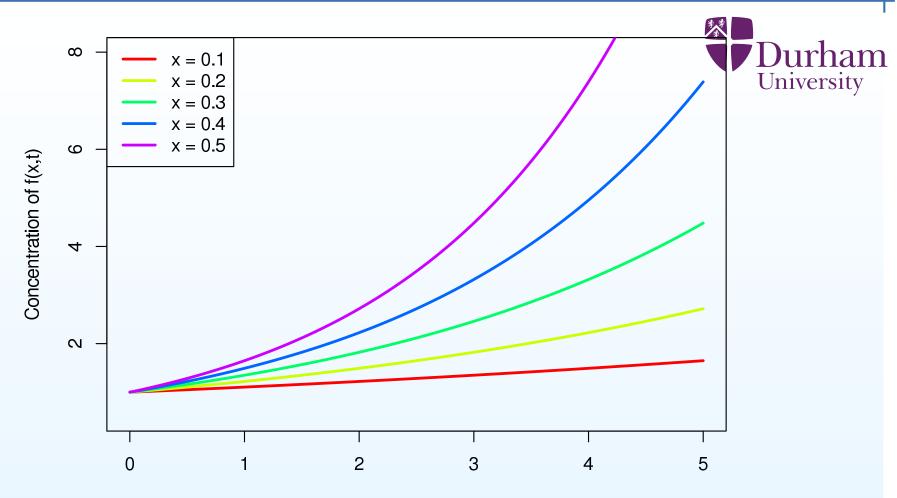
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Model features an input parameter x which we want to learn about.

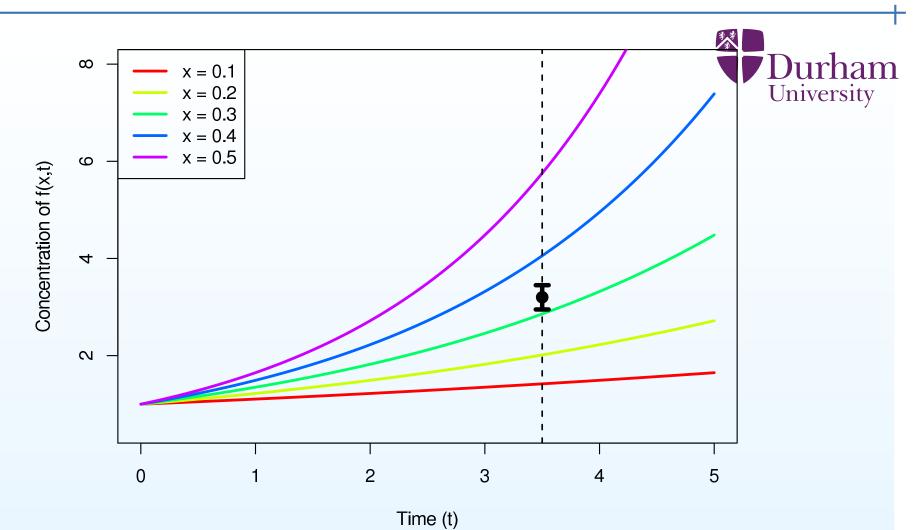


One "model run" with the input parameter x = 0.4



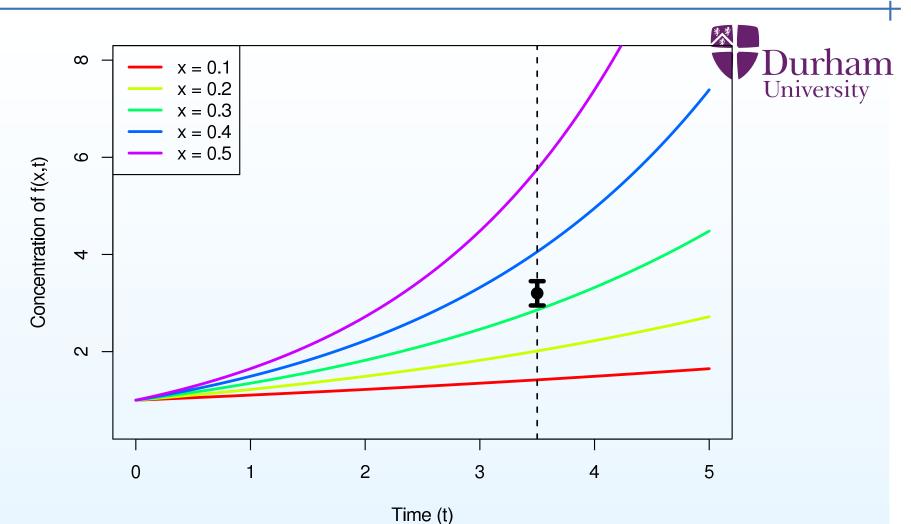
Time (t)

Five model runs with the input parameter varying from x = 0.1 to x = 0.5



We are going to measure f(x,t) at t = 3.5

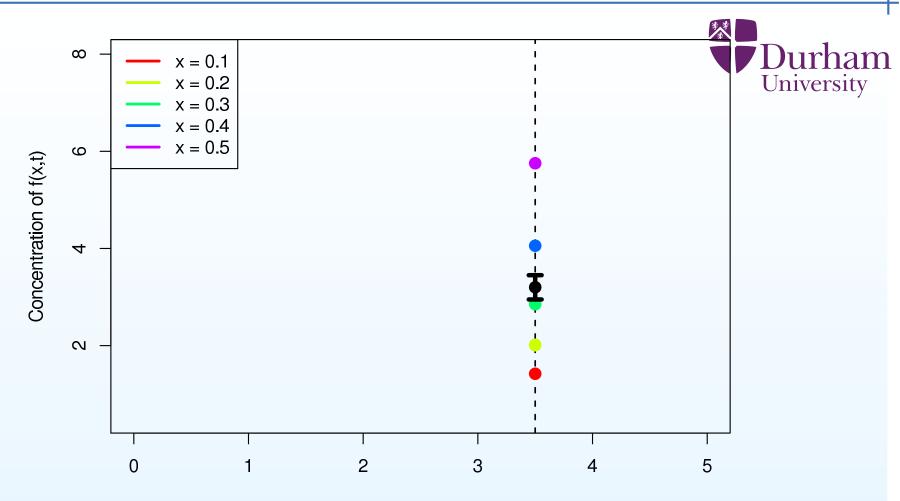
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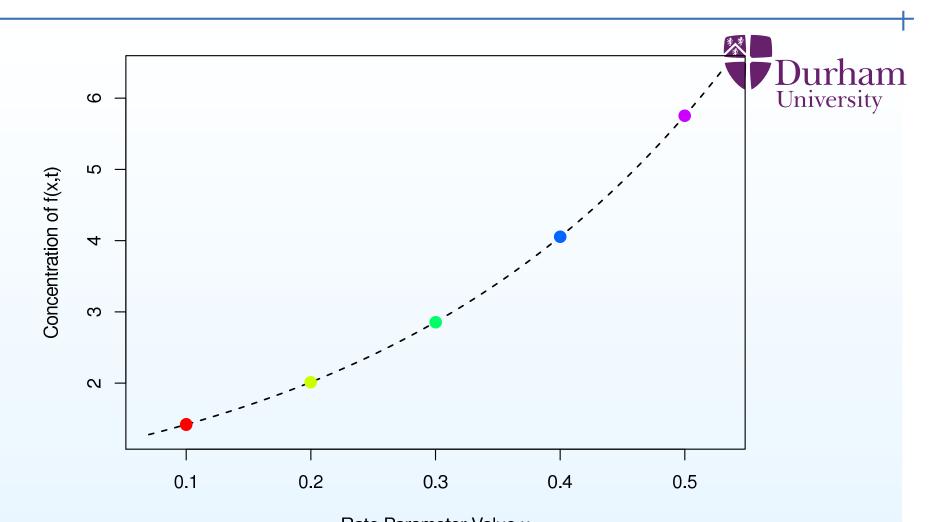
The measurement comes with measurement error.

For which values of x is the output f(x, t = 3.5) consistent with the observation?

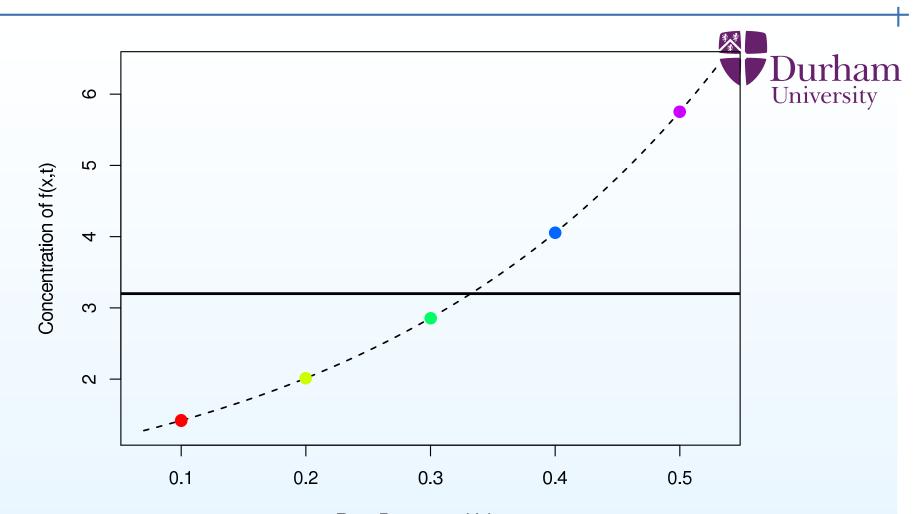


Time (t)

To answer this, we can now discard other values of f(x,t) and think of f(x,t=3.5) as a function of x only, that is take  $f(x) \equiv f(x,t=3.5)$ 

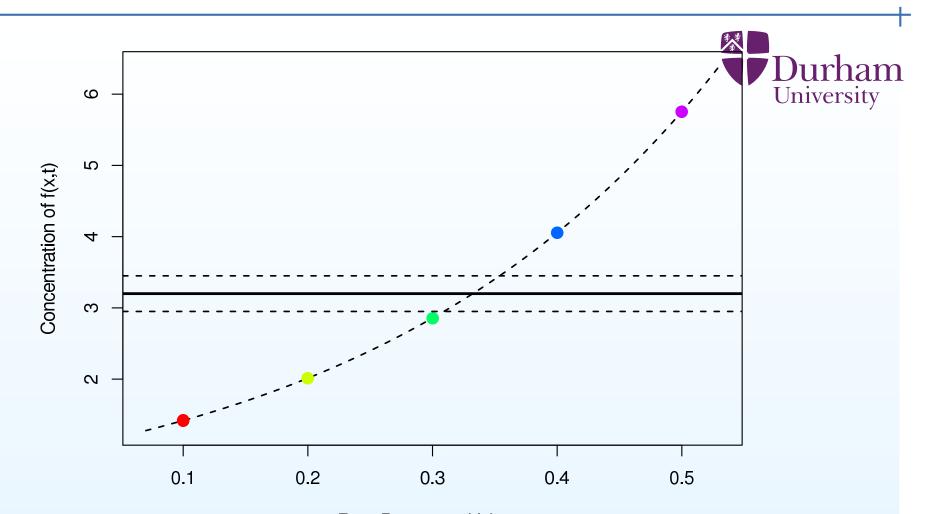


Rate Parameter Value x We plot the concentration f(x) as a function of the input parameter x.



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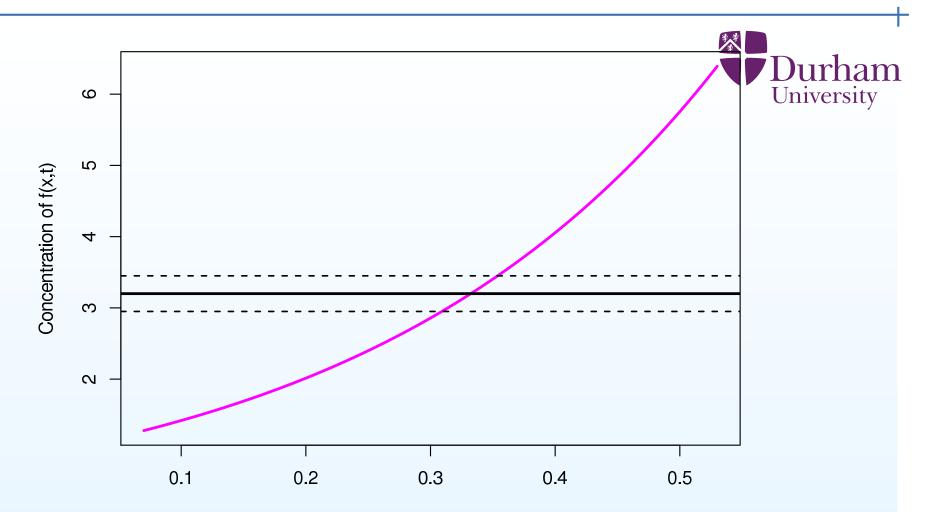
Black horizontal line: the observed measurement of f



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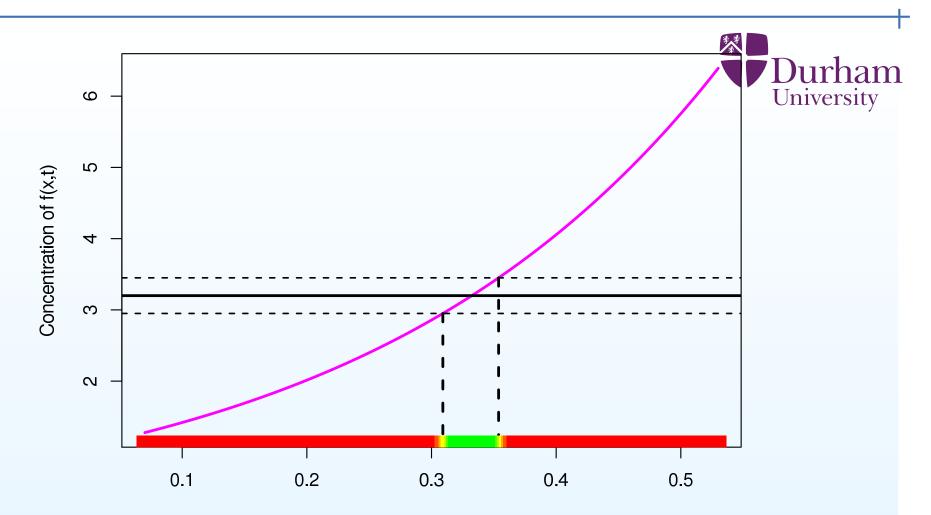
Black horizontal line: the observed measurement of f

Dashed horizontal lines: the measurement errors



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Hence we see a range (green/yellow) of possible values of x consistent with the measurements, with all the implausible values of x in red.

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Different physical models vary in many aspects, but the approaches for addressing these problems are very similar

(which is why there is a common underlying methodology).



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In almost all cases (i) evaluation of f(x) is expensive (ii) inferring  $x^*$  from z is hard (iii) relating  $f(x^*)$  to y is challenging.



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An excellent resource for work in this area is the Managing Uncertainty in Complex Models web-site, www.mucm.ac.uk



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Bayes linear adjustment may be viewed as an approximation to a full Bayes analysis or the appropriate analysis given a partial specification.



The Bayes linear adjusted expectation and variance for vector  $\boldsymbol{y}$  given vector  $\boldsymbol{z}$  are

$$\mathsf{E}_{z}[y] = \mathsf{E}(y) + \operatorname{Cov}(y, z)\operatorname{Var}(z)^{-1}(z - \mathsf{E}(z)),$$
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#### For a detailed treatment, see

Bayes linear Statistics: Theory and Methods, 2007, (Wiley) Michael Goldstein and David Wooff

#### For a quick overview, see

Bayes linear analysis, 2015, Michael Goldstein, in Wiley StatsRef: Statistics Reference Online (7 pages)

And all of our papers in this area contain examples of Bayes linear computations.



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Unlike the original simulator, the emulator is fast to evaluate for any choice of inputs. This allows us to explore model behaviour for all physically meaningful input specifications.

#### Form of the emulator



We may represent beliefs about component  $f_i$  of f, using an emulator:

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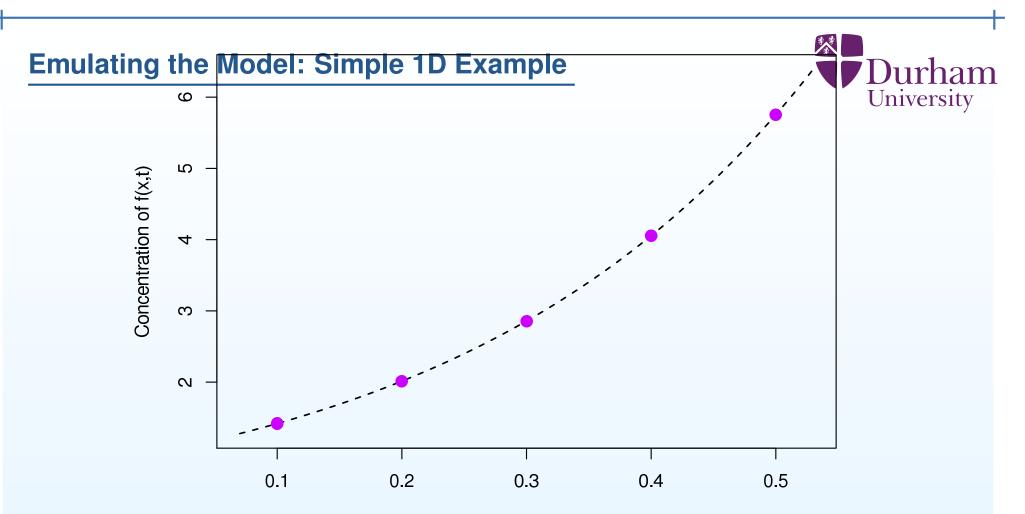
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#### **Local Variation**

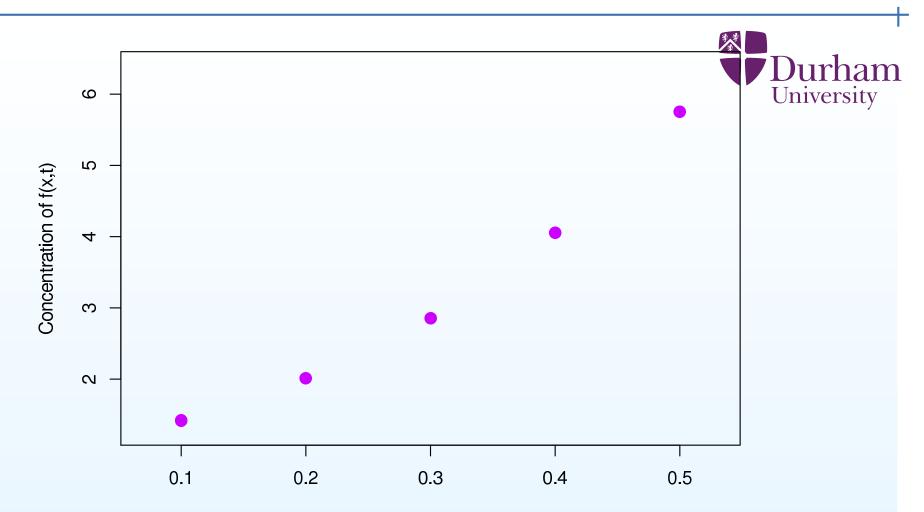
 $u_i(x)$  is a second order stationary stochastic process, with (for example) correlation function

$$\operatorname{Corr}(u_i(x), u_i(x')) = \exp(-(\frac{\|x - x'\|}{\theta_i})^2)$$



Rate Parameter Value x

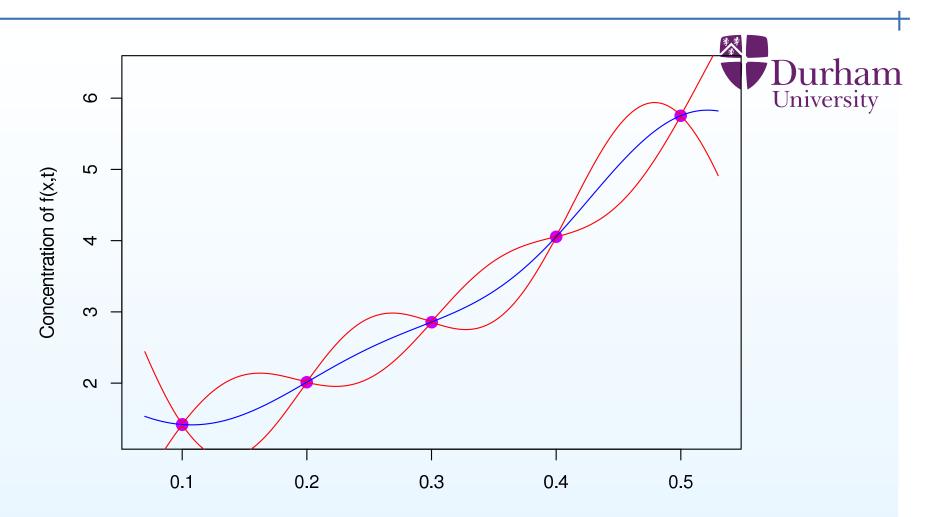
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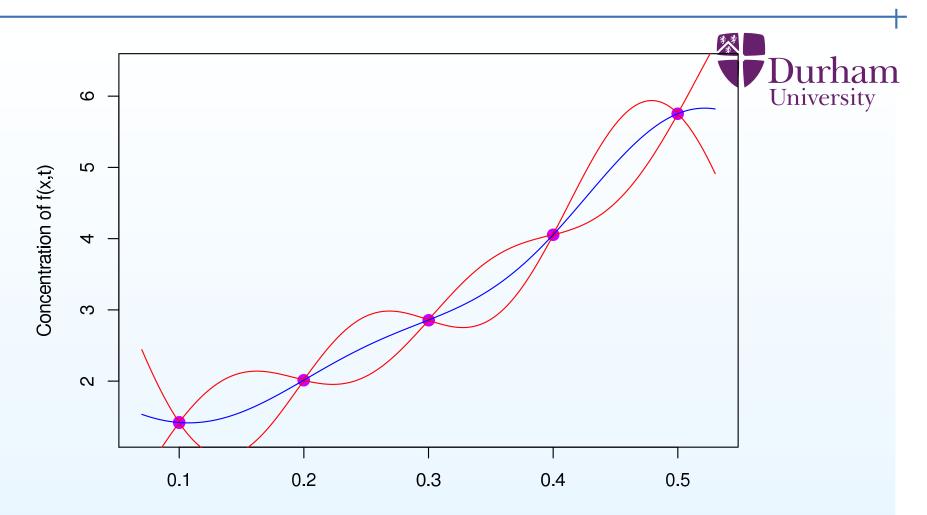
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Instead we only have a finite number of runs of the model, in this case five.

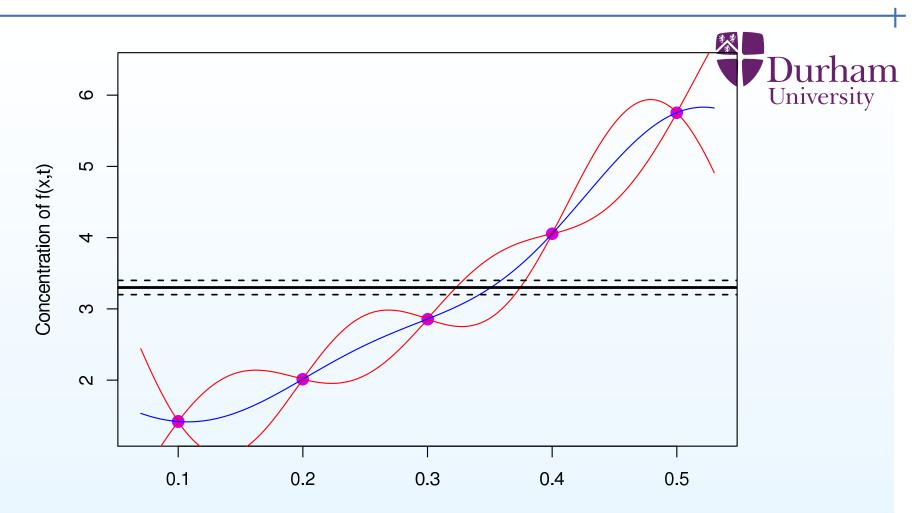


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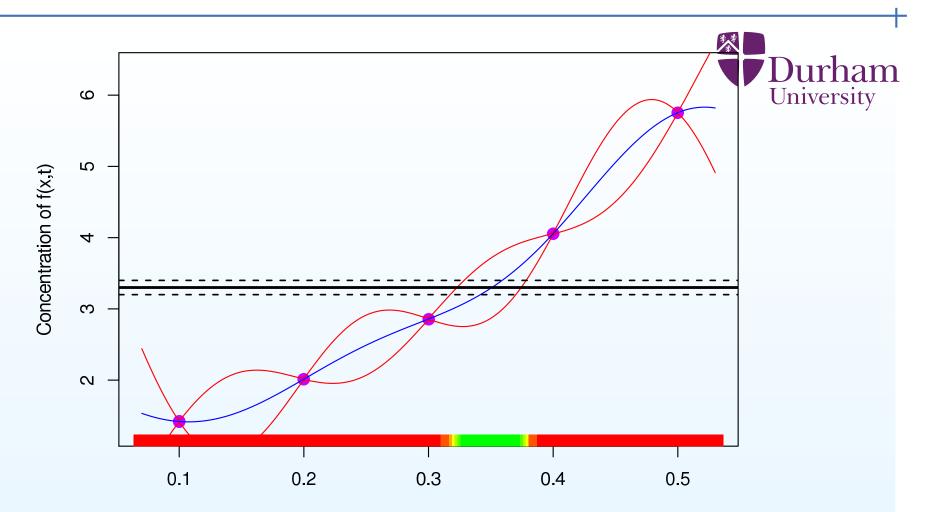
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It gives both the expected value of f(x) (the blue line) along with a credible interval for f(x) (the red lines) representing the uncertainty about the model's behaviour.



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The uncertainty on x now includes uncertainty coming from the emulator.



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We use efficient space filling (multi-level) designs to generate the set of simulator evaluations to carry out in order to fit the emulators. (For example, maximin Latin Hypercubes.)



We fit the emulators, given a collection of carefully chosen model evaluations, using our favourite statistical tools - generalised least squares, maximum likelihood, Bayes - with a generous helping of expert judgement.

If the model is slow to evaluate, we typically create an informed prior assessment based on a fast approximation, then combine with a carefully designed set of runs of the full simulator to construct the emulator.

We use efficient space filling (multi-level) designs to generate the set of simulator evaluations to carry out in order to fit the emulators. (For example, maximin Latin Hypercubes.)

We use careful diagnostics to test the validity of our emulators (for example, assessing the reliability of the emulator for predicting the simulator at new evaluations).



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- (iii) full probabilistic calibration analysis may be very difficult/non-robust for complex simulators.
- (because the likelihood surface is complicated and multi-modal, and the Bayes answer often depends on features of the prior distribution which are hard to specify meaningfully)

# **History matching**



A conceptually simple procedure is "history matching".

This means finding the collection, C(z), of all input choices x for which the match of the simulator outputs  $f_h(x)$  to observed data, z, is good enough, taking into account all of the uncertainties in the problem.

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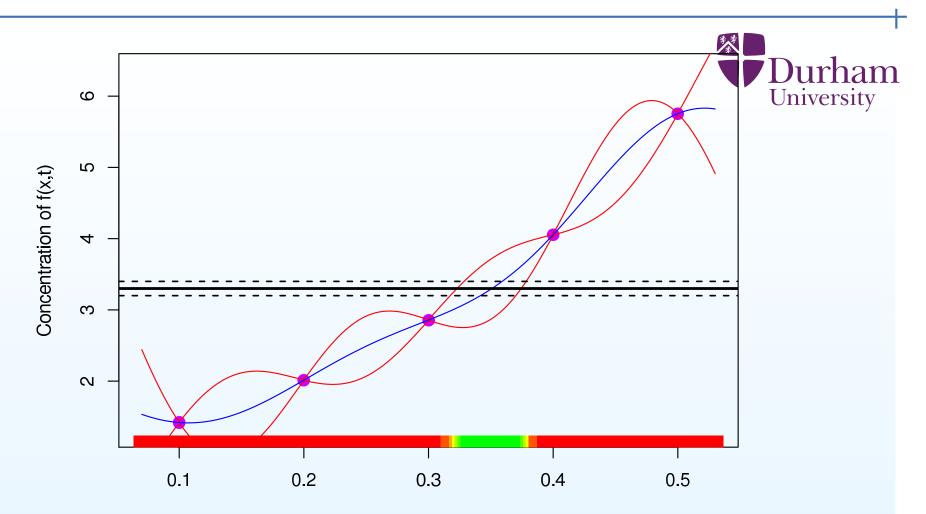
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If the data is informative for the parameter space, then C(z) will typically form a tiny percentage of the original parameter space.

Therefore, even if we do wish to calibrate the simulator, history matching is a useful preliminary step.



Comparing the emulator to the observed measurement we have identified the set of x values (the green values) which match the observed history, when we take into account all of the uncertainties (here, measurement and emulator error).

# History matching by implausibility



We use an 'implausibility measure' I(x) based on a probabilistic metric such as

$$I(x) = \frac{(z - \mathcal{E}(f_h(x)))^2}{\operatorname{Var}(z - \mathcal{E}(f_h(x)))}$$

(where the variance in the denominator is the sum of all of the individual variance terms e.g. measurement error, emulator error, discrepancy error and so forth.)

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The implausibility calculation can be performed univariately, or by multivariate calculation over sub-vectors.

The implausibilities are then combined to identify x with large I(x) as implausible, i.e. unlikely to be appropriate choices for system inputs.



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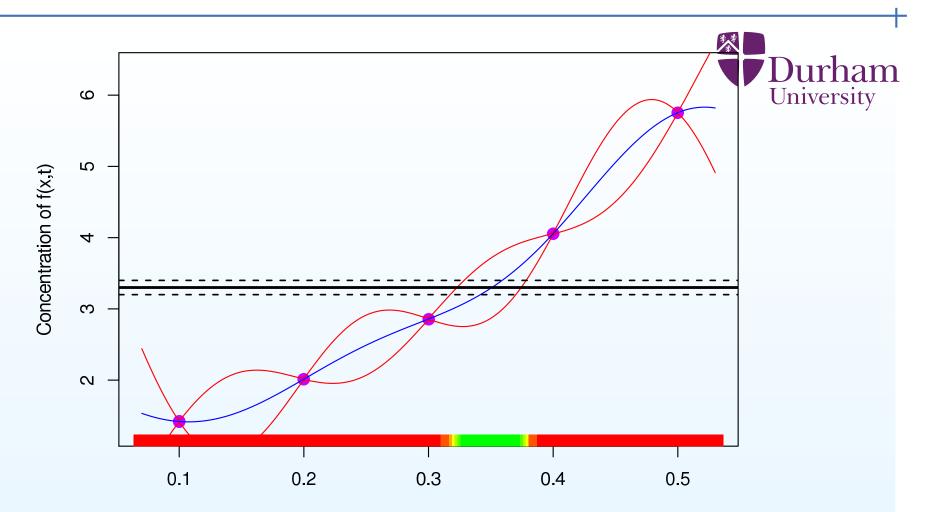
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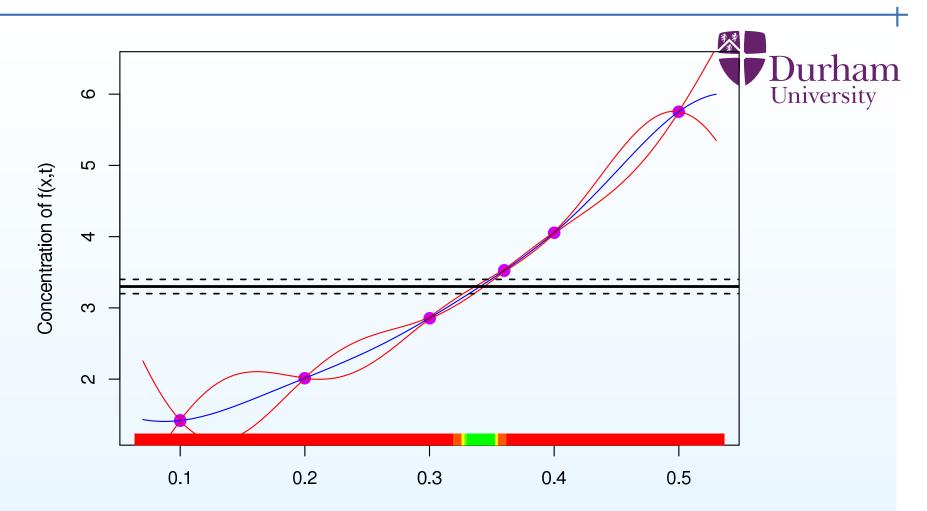
and repeating the implausibility analysis.

We continue until (hopefully) we identify the region of acceptable matches.

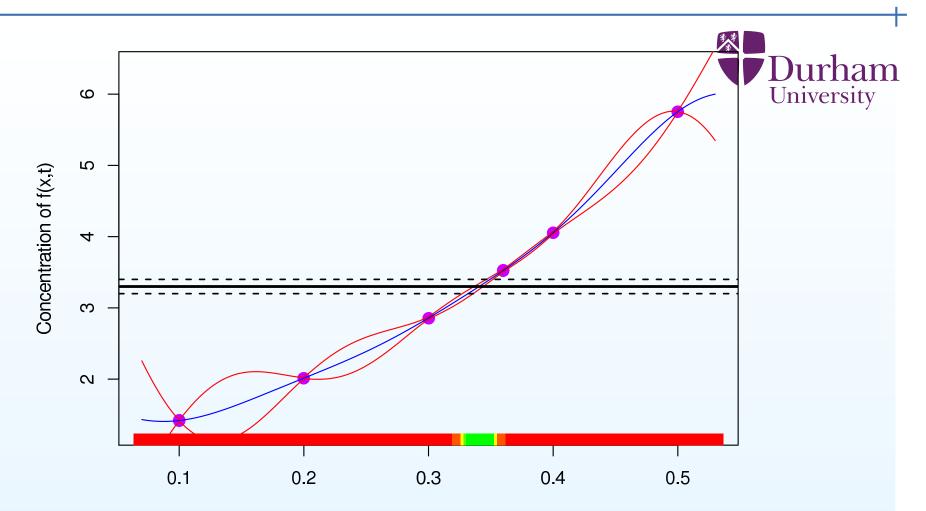
(This is a form of iterative global search.)



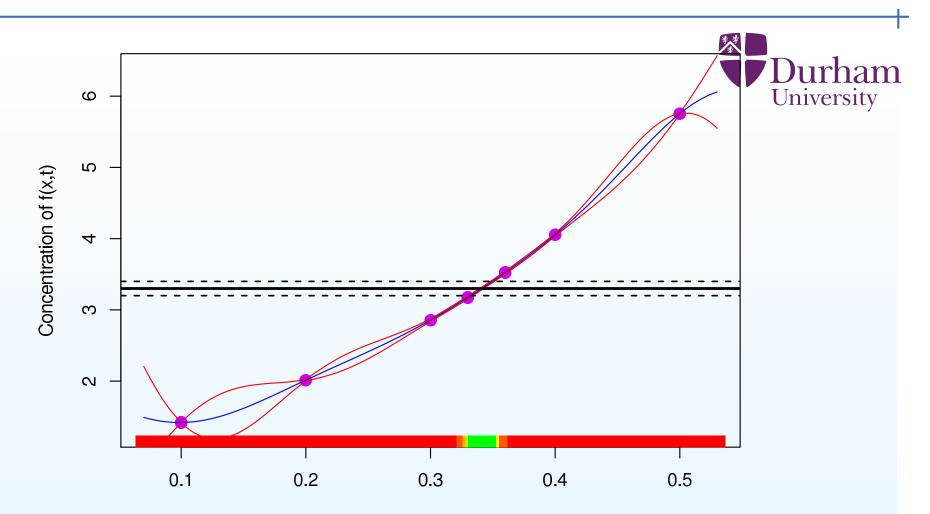
We now remove all of the implausible x values (the red values) and resample and re-emulate within the green region.



We perform a 2nd iteration or wave of runs to improve emulator accuracy.



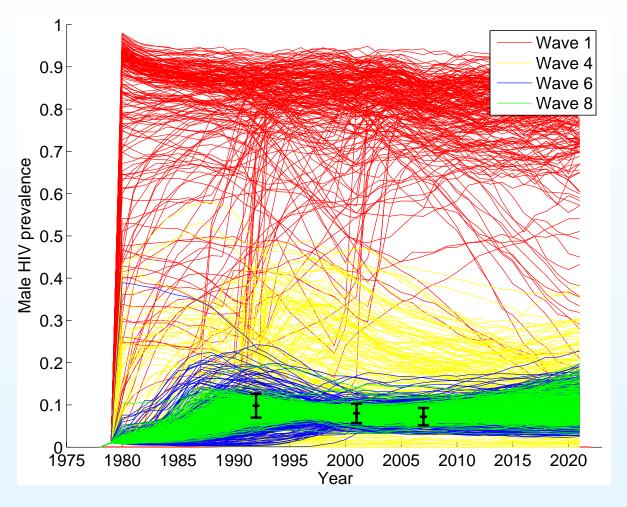
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We perform a 2nd iteration or wave of runs to improve emulator accuracy. The runs are located only at non-implausible (green/yellow) points. Now the emulator is more accurate than the observation, and we can identify the set of all x values of interest.

## History matching for the case study





In the case study, after 10 waves, we have reduced the space to about  $10^{-11}$  of original space. Around 65% of the simulator evaluations in the final space give runs with acceptable matches to the historical data.



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Neither of these approximations invalidates the modelling process.

Problems only arise when we forget these simplifications and confuse the analysis of the model with the corresponding analysis for the physical system itself.

## Relating the model and the system





Actual system

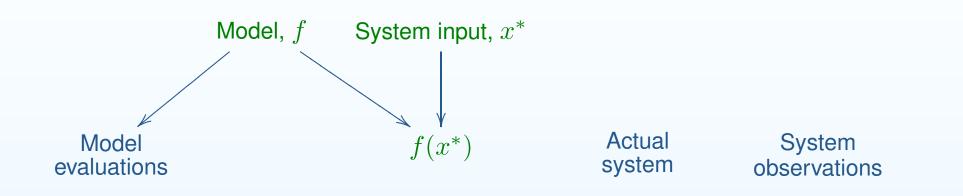
System observations

1. We start with a collection of model evaluations, and some observations on actual system

- 2. We link the model evaluations to the evaluation of the model at the (unknown) system values  $x^*$  for the inputs
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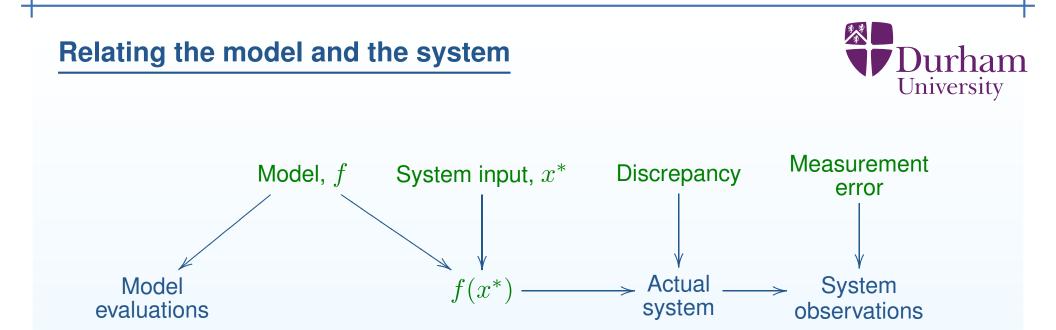




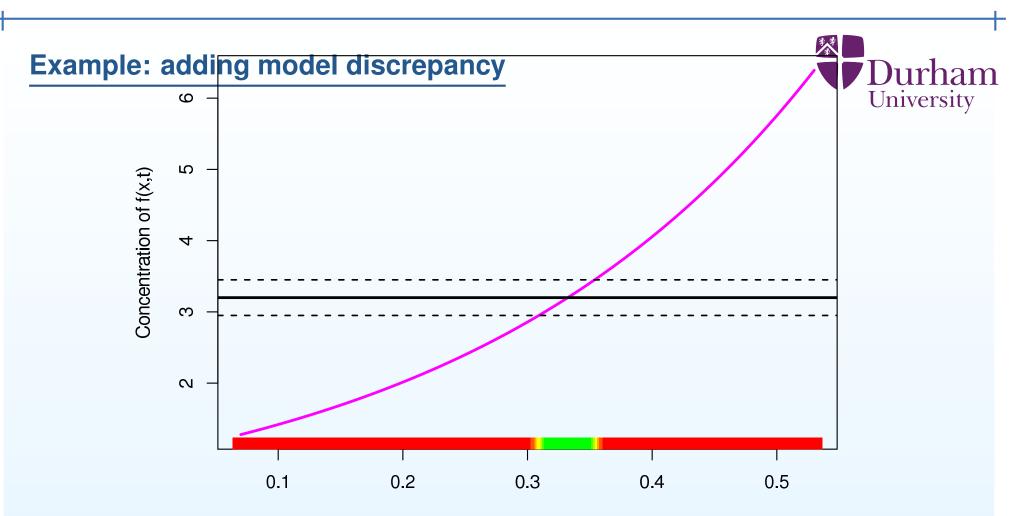
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# Model f System input, $x^*$ Discrepancy System Model $f(x^*)$ Actual system System

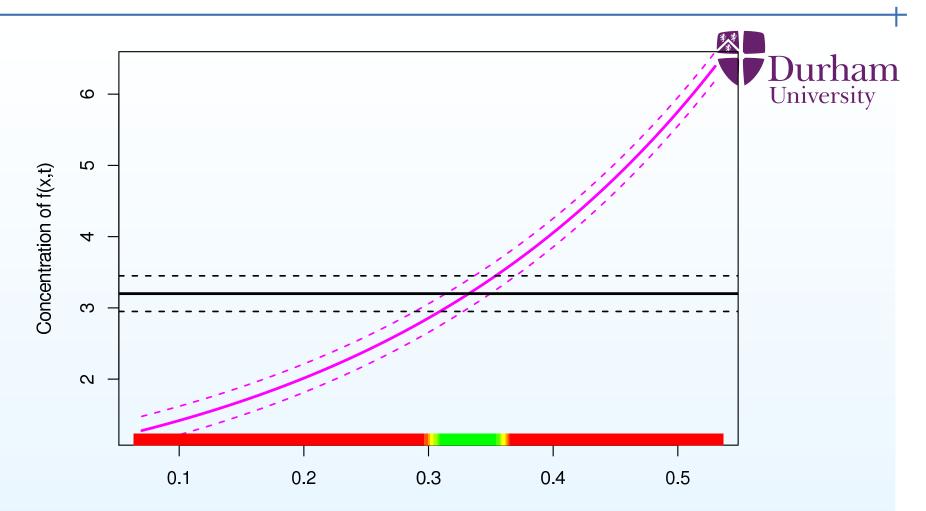
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Rate Parameter Value x The notion of model discrepancy is related to how accurate we believe the model to be.



Rate Parameter Value x

Model discrepancy is represented as uncertainty around the model output f(x) itself: here the purple dashed lines.

This results in more uncertainty in x, and hence a larger range of x values.

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### (i) Internal discrepancy

Any aspect of discrepancy we can assess by direct experiments on the computer simulator.

#### For example,

we may vary parameters held fixed in the standard analysis,

we may add random noise to the state vector which the model propagates,

we may allow parameters to vary over time.

we may add noise to the forcing functions used to evaluate the simulator

Durham University

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Note, in particular, that this method gives an order of magnitude assessment for the correlation between discrepancy values across time.

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The simplest way to incorporate external discrepancy is to add an extra component of uncertainty to the simulator outputs. For example we may introduce, say, 10% additional error to account for structural discrepancy.

(This is simple, but much better than ignoring external discrepancy.)



Better is to consider what we know about the limitations of the model, and build a probabilistic representation of additional features of the relationship between system properties and behaviour.

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For example, we can treat our actual simulator as a prior for the reified form. This is similar to the way in which we use fast simulators to act as priors for slow simulators.



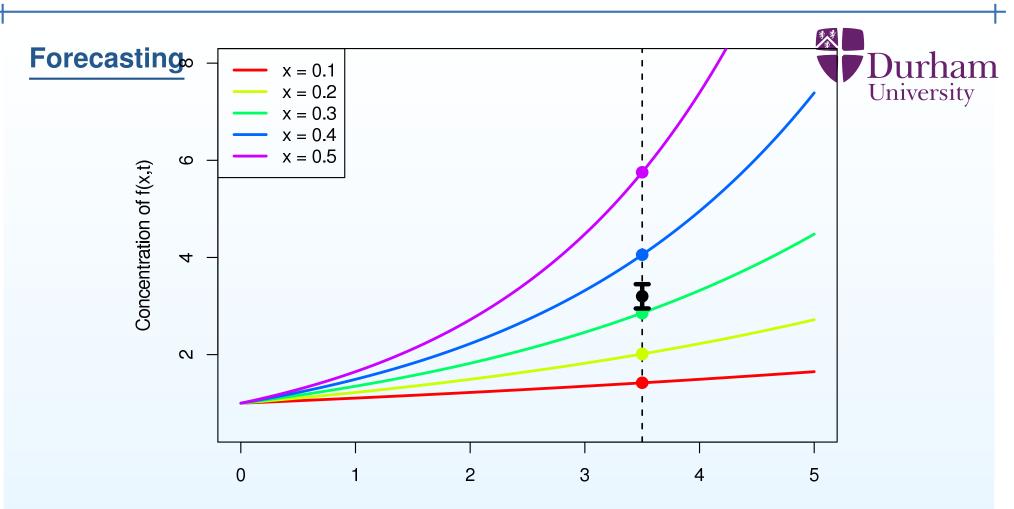
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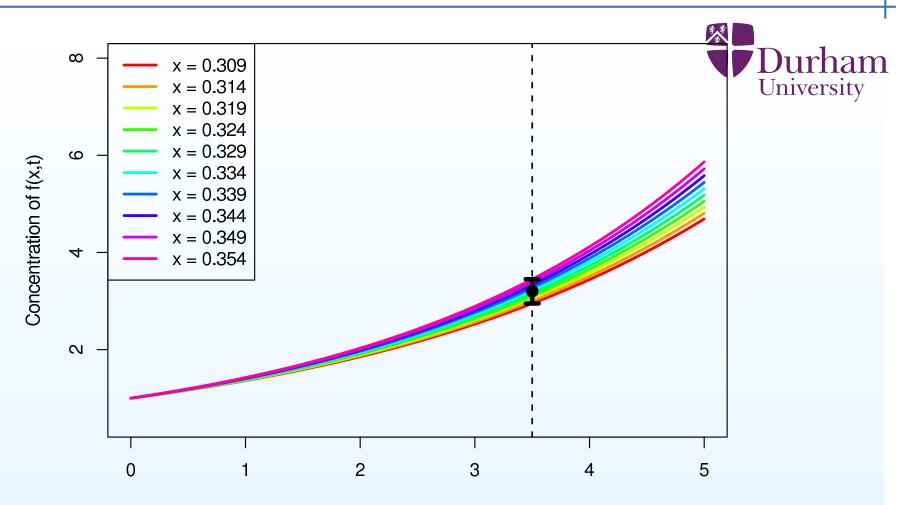
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So, the methods for history matching based on emulation will work in the same way using the reified emulator.



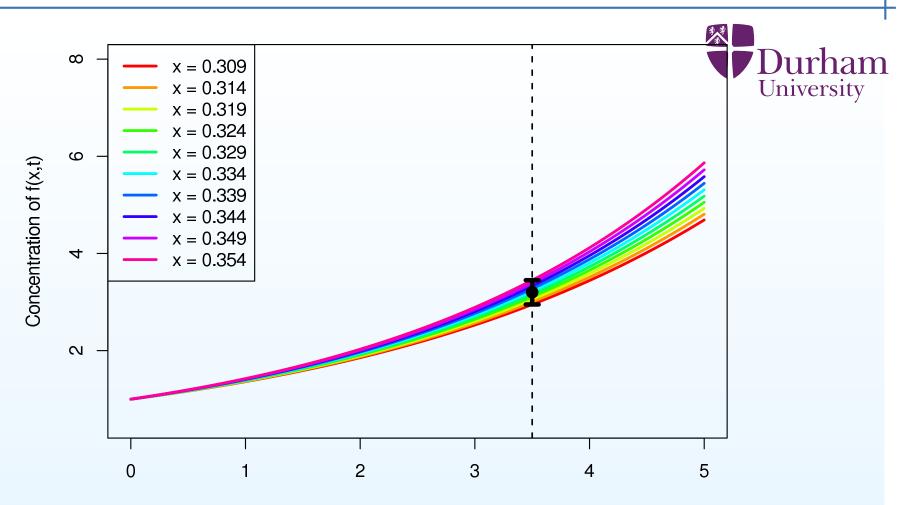
Time (t)

Constraints on x from observations impose constraints on f(x, t) in the future.



Time (t)

We choose values of x consistent with the measurement of f(x,t) at t = 3.5, and perform corresponding runs of the simulator, possibly at a variety of control choices. If the simulator is expensive, we may emulate these future outcomes.



Time (t)

These are future projections within the simulator. To transfer these to future projections for the world we need to add the effects of structural discrepancy.



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(iii) help us to make reliable control choices for future outcomes.

(by recognising the real world risks of our various control choices).



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(iii) **structural discrepancy modelling**, to make reliable uncertainty statements about the real world

#### References



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