

Bayesian Experimental Design for Stochastic Biological Models with Computational Demanding Likelihoods

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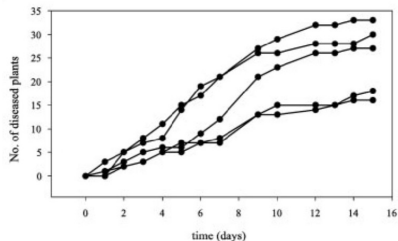
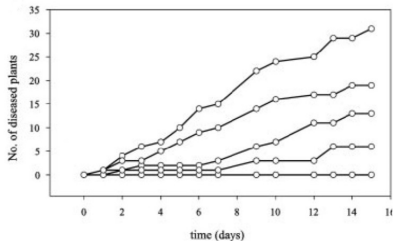
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Motivating Example

Kleczkowski et al (1996); Gibson et al (2004)



- Two competing models for each dataset (left treatment applied)
- When should samples be taken to estimate parameters or discriminate between models?

Models proposed for data

Markov Model (in deterministic DE form)

$$\frac{dl}{dt} = (r_p + r_s l) e^{-at} (N - l)$$

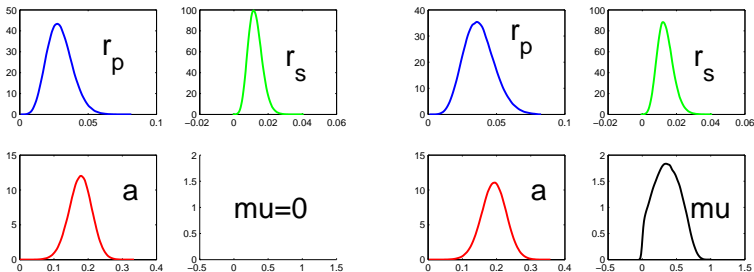
- Likelihood is expensive (if evaluated many times).
 $\theta = (r_p, r_s, a)$.

Non-Markov Model (in deterministic DE form)

$$\frac{dE}{dt} = (r_p + r_s l) e^{-at} (N - l - E)$$

- An exposed (E) plant becomes infectious (I) after μ units of time. $\theta = (r_p, r_s, a, \mu)$
- Likelihood intractable
- Easy to simulate from both models.
- Models are nested

The problem



Priors for future experiment with no treatment, Markov (left), non-Markov (right)

Models fitted via (auxiliary variable) MCMC.

- Posterior becomes the prior for future experiments (figure above)
- Objective: Determine optimal sampling times for estimating parameters and discriminating between models

Static Experimental Design

- Set-up: Have prior distribution $p(\boldsymbol{\theta})$ for parameter of statistical model, with likelihood $p(\mathbf{y}|\boldsymbol{\theta}, \mathbf{d})$.
- Wish to design for next n observations. Design variable: $\mathbf{d} = (d_1, \dots, d_n)$.
- Define (general) utility function $u(\mathbf{d}, \mathbf{y}, \boldsymbol{\theta})$. \mathbf{y} is future data.

$$u(\mathbf{d}) = E_{\boldsymbol{\theta}, \mathbf{y}}[u(\mathbf{d}, \mathbf{y}, \boldsymbol{\theta})] = \int_{\mathbf{y}} \int_{\boldsymbol{\theta}} u(\mathbf{d}, \mathbf{y}, \boldsymbol{\theta}) p(\mathbf{y}|\mathbf{d}, \boldsymbol{\theta}) p(\boldsymbol{\theta}) d\boldsymbol{\theta} d\mathbf{y},$$

- Model uncertainty can be included ($m = 1, \dots, K$ models)

$$u(\mathbf{d}) = \sum_{m=1}^K \int_{\mathbf{y}} \int_{\boldsymbol{\theta}} u(\mathbf{d}, \mathbf{y}, \boldsymbol{\theta}, m) p(\mathbf{y}|\mathbf{d}, \boldsymbol{\theta}_m, m) p(\boldsymbol{\theta}_m|m) p(m) d\boldsymbol{\theta} d\mathbf{y},$$

- Objective

$$\mathbf{d}^* = \arg \max_{\mathbf{d} \in \mathcal{D}} u(\mathbf{d}).$$

Too difficult to do directly

MCMC approach - Muller 1999

- Turn optimisation problem into simulation problem

$$h(\mathbf{d}, \boldsymbol{\theta}, \mathbf{y}) \propto u(\mathbf{d}, \mathbf{y}, \boldsymbol{\theta})p(\mathbf{y}|\mathbf{d}, \boldsymbol{\theta})p(\boldsymbol{\theta}).$$

Admits $\propto u(\mathbf{d})$ as marginal distribution

- Optimal design \mathbf{d}^* is mode of marginal
- If marginal is flat, may be beneficial to sample from

$$h(\mathbf{d}, \boldsymbol{\theta}_1, \dots, \boldsymbol{\theta}_J, \mathbf{y}_1, \dots, \mathbf{y}_J) \propto \prod_{j=1}^J u(\mathbf{d}, \mathbf{y}_j, \boldsymbol{\theta}_j)p(\mathbf{y}_j|\mathbf{d}, \boldsymbol{\theta}_j)p(\boldsymbol{\theta}_j),$$

for large J . A marginal is $\propto u(\mathbf{d})^J$

- Muller (1999) proposes MCMC for sampling.
 $q(\mathbf{d}^*, \mathbf{y}^*, \boldsymbol{\theta}^*|\mathbf{d}, \mathbf{y}, \boldsymbol{\theta}) = p(\mathbf{y}^*|\boldsymbol{\theta}^*, \mathbf{d}^*)p(\boldsymbol{\theta}^*)q(\mathbf{d}^*|\mathbf{d}).$
- Easy to extend to model uncertainty

Bayesian utility functions

- Based on posterior, so independent of θ , $u(\mathbf{d}, \mathbf{y})$. Still need θ for proposal in above MCMC

Parameter Estimation Design

- **Kullback-Leibler divergence** $u(\mathbf{d}, \mathbf{y}) = KL(p(\theta|\mathbf{y}, \mathbf{d})||p(\theta))$.
Mutual info between θ and \mathbf{y} .
- Concentration of Posterior distribution
 - Set $u(\mathbf{d}, \mathbf{y})$ as entropy of $p(\theta|\mathbf{y}, \mathbf{d})$
 - **Posterior precision** $u(\mathbf{d}, \mathbf{y}) = 1/\det(\text{var}(\theta|\mathbf{y}, \mathbf{d}))$

Bayesian utility functions (cont...)

Model Discrimination Design

- For nested models focus on **estimating 'extra' parameters**.
- General Bayesian model discrimination utility is **mutual information** between M and \mathbf{y} . Drovandi et al 2013 JCGS show that

$$u(\mathbf{d}, \mathbf{y}, m) = \log p(m|\mathbf{y}, \mathbf{d}).$$

Even if likelihood available then $u(\mathbf{d}, \mathbf{y})$ and $p(m|\mathbf{y}, \mathbf{d})$ must still be estimated.

Muller Algorithm

- A closer look at the Muller algo... \mathbf{d}^{i-1} is current, with utility $u^{i-1} = u(\mathbf{d}^{i-1}, \boldsymbol{\theta}^{i-1}, \mathbf{y}^{i-1})$
- Propose $\mathbf{d}^* \sim q(\mathbf{d}|\mathbf{d}^{i-1})$, $\boldsymbol{\theta}^* \sim p(\boldsymbol{\theta})$, $\mathbf{y}^* \sim p(\mathbf{y}|\boldsymbol{\theta}^*, \mathbf{d}^*)$.
Compute $u^* = u(\mathbf{d}^*, \boldsymbol{\theta}^*, \mathbf{y}^*)$
- Compute $\alpha = \min\left(1, \frac{u^* q(\mathbf{d}^{i-1}|\mathbf{d}^*)}{u^{i-1} q(\mathbf{d}^*|\mathbf{d}^{i-1})}\right)$
- Likelihood functions cancel as per ABC MCMC (Marjoram et al (2003))
- Now simply need utility function that does not require likelihood evaluation

Bayesian utilities in presence of intractable likelihoods

- Bayesian utilities based on posterior.
- Approximate true posterior via ABC (soon)

Parameter Estimation

- KLD between prior and ABC posterior (normal approximation)
- Concentration of ABC posterior (e.g. entropy or precision).
Precision straightforward to calculate:

$$u(\mathbf{d}, \mathbf{y}) = 1/\det(\text{Var}_{\theta|\mathbf{y},\mathbf{d}}(\theta)).$$

Model Discrimination

- KLD or concentration of ABC posterior of extra parameters.
- For mutual info use ABC posterior model probability



Approximate Bayesian Computation

- Simulation based method that does not involve likelihood evaluations
- Involves a joint 'approximate' posterior distribution

$$p(\boldsymbol{\theta}, \mathbf{x} | \mathbf{y}, \epsilon) \propto g(\mathbf{y} | \mathbf{x}, \epsilon) p(\mathbf{x} | \boldsymbol{\theta}) p(\boldsymbol{\theta})$$

- where $g(\mathbf{y} | \mathbf{x}, \epsilon)$ is a weighting function. Popular choice
 $g(\mathbf{y} | \mathbf{x}, \epsilon) = 1(\rho(\mathbf{y}, \mathbf{x}) \leq \epsilon)$
- How to choose $\rho(\mathbf{y}, \mathbf{x})$?
 - Usually based on a set of low-dimensional set of summary statistics
 - Here use full discrete data (low-dimensional designs)
- Choice of ϵ trade-off between accuracy and efficiency (and Monte Carlo error)

The effect of the approximation

- Idea is that marginal $p(\boldsymbol{\theta} | \mathbf{y}, \epsilon) \approx p(\boldsymbol{\theta} | \mathbf{y})$
- Errors from insufficient summaries and $\epsilon_i > 0$

ABC Algorithms

- Many algorithms available: ABC rejection (Beaumont et al 2002), MCMC ABC (Marjoram et al (2003)), SMC ABC (Sisson et al (2007), CD+TP (2011))
- MCMC ABC and SMC ABC suitable for single dataset analysis
- In context of static design, need ABC posterior at each iteration for a new \mathbf{y}
- ABC rejection amendable to multiple datasets from same model
- Easy to extend ABC rejection to include model uncertainty (Grelaud et al 2009)

ABC Rejection

- 1 Generate $\theta^i \sim p(\theta)$ for $i = 1, \dots, N$
- 2 Simulate $\mathbf{x}^i \sim p(\mathbf{y}|\theta^i, \mathbf{d})$ for $i = 1, \dots, N$
- 3 Compute discrepancies $\rho^i = \rho(\mathbf{y}, \mathbf{x}^i)$ for $i = 1, \dots, N$, creating particles $\{\theta^i, \rho^i\}_{i=1}^N$
- 4 Sort the particle set via the discrepancy ρ
- 5 Calculate $\epsilon = \rho^{\lfloor \alpha N \rfloor}$ (where $\lfloor \cdot \rfloor$ denotes the floor function).
The ABC posterior samples consist of the set $\{\theta^i | \rho^i \leq \epsilon\}_{i=1}^N$

Steps 1 and 2 are independent of data and $\{\theta^i, \mathbf{x}^i\}_{i=1}^N$ can be stored.

Discrepancy function:

$$\rho(\mathbf{y}, \mathbf{x}) = \sum_{i=1}^D \frac{|y_i - x_i|}{\text{std}_{p(\theta)}(x_i)}, \quad (1)$$



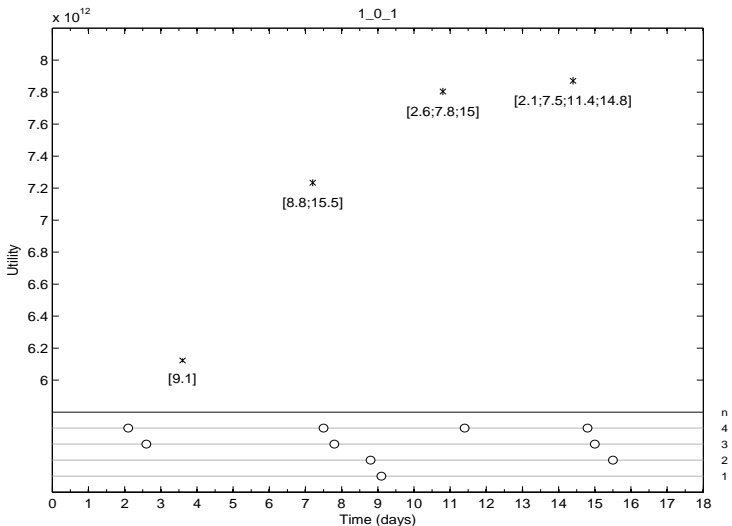
Discretising the Design Space

- Discretise design (time) space: $t_{min}, t_{max}, t_{inc}$
- Prior simulations get recorded at each design point in design space
- Prior simulations are done before Muller algorithm
- Metropolis-Hastings sample over discrete design space

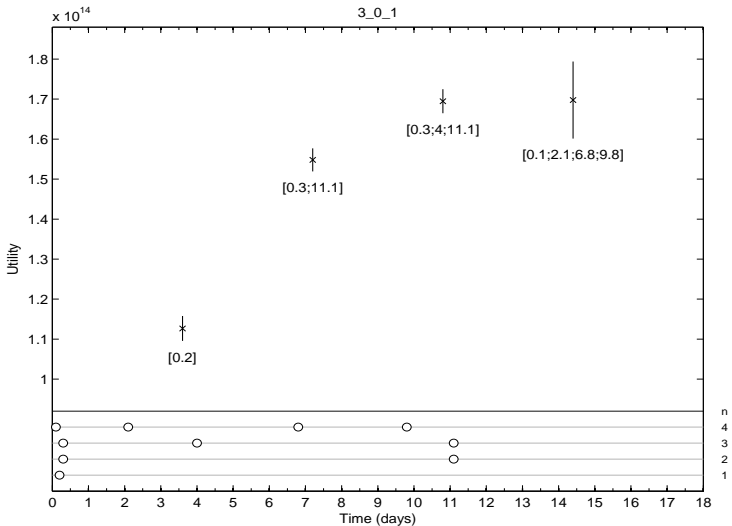
Algorithm Settings

- Run Muller algorithm for 100,000 iterations
- 100,000 prior simulations recorded at time interval $t_{inc} = 0.1$.
- Determine ϵ based on 200th smallest discrepancy.
- $J = 5$ for better identification of model
- Non-parametric estimate of multivariate density function of \mathbf{d} , then maximise. (see Cook et al (2008))

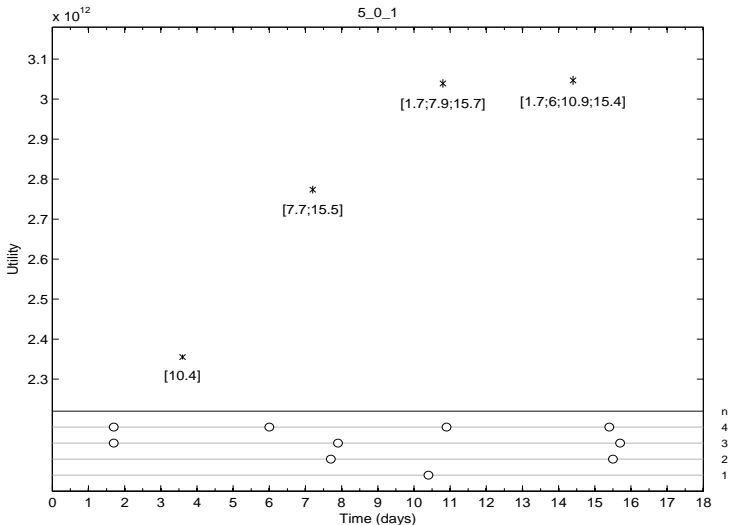
Parameter Estimation (Markov model)



Parameter Estimation (non-Markov model)



Parameter Estimation (non-Markov model excluding μ)



Parameter Estimation (non-Markov model just μ)

RESULTS STILL TO COME

Model Discrimination

RESULTS STILL TO COME



Limitations and Future Work

Limitations

- Only suitable for low dimensional designs and low number of sampling times.
- Initial condition needs to be fixed, or only a few options
- ABC rejection needs informative priors → but so does Bayesian design in general

Future Work

- Extend to larger number of sampling times (EP-ABC Algorithm)

Key References

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