The background of the slide features a close-up photograph of two ladybugs on a green plant stem. One ladybug is positioned higher on the stem, and another is lower down. The background is a soft-focus green, suggesting foliage.

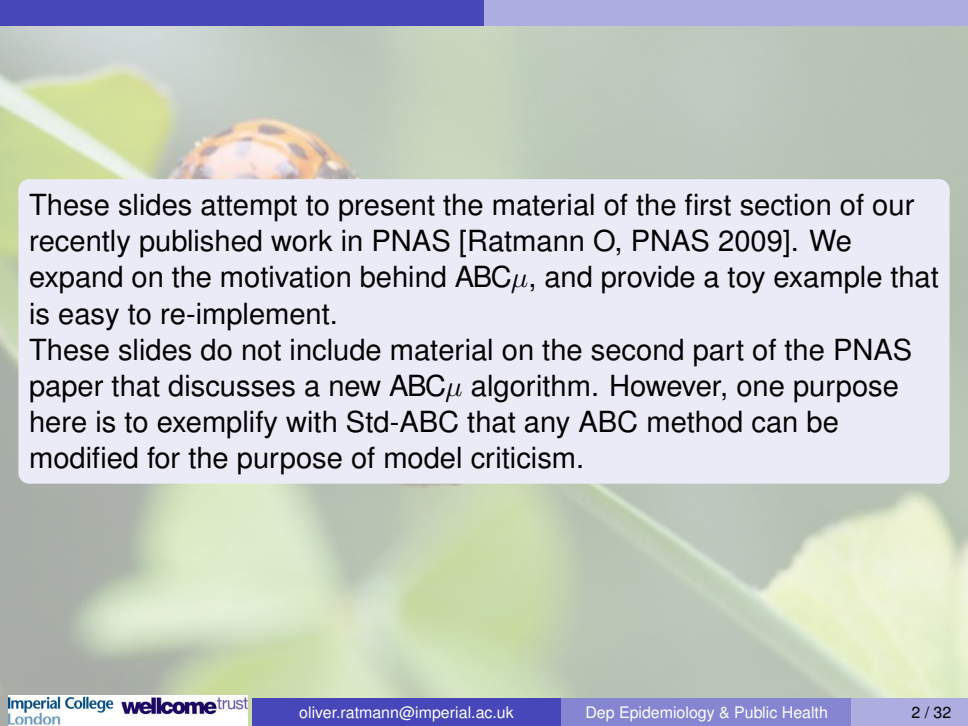
Approximate Bayesian Computation under model uncertainty (ABC μ)

Oliver Ratmann¹ C Andrieu² C Wiuf³ S Richardson¹

¹Centre for Biostatistics, Imperial College, London, UK

²Dep of Mathematics, University of Bristol, Bristol, UK

³Bioinformatics Research Centre, University of Aarhus, Aarhus, DK



These slides attempt to present the material of the first section of our recently published work in PNAS [Ratmann O, PNAS 2009]. We expand on the motivation behind ABC_{μ} , and provide a toy example that is easy to re-implement.

These slides do not include material on the second part of the PNAS paper that discusses a new ABC_{μ} algorithm. However, one purpose here is to exemplify with Std-ABC that any ABC method can be modified for the purpose of model criticism.

When use ABC / ABC μ ?

Inference on
data-generating stochastic
processes

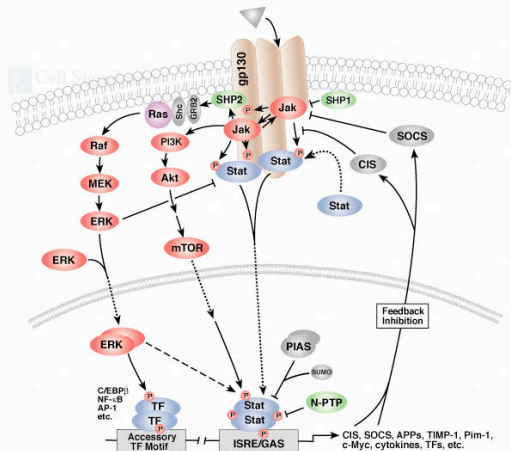
[Diggle PJ, RSB 1984]

... whose likelihood cannot
be readily evaluated

... within Bayesian
framework to
(a) infer model parameters,
(b) error terms to see if
model adequate

Signalling pathways, [Toni T RoySocInt 2008]

Jak/Stat Signaling: IL-6 Receptor Family



“All models are wrong but some are useful [Box GEP, JASA1976]”

Statistical reasoning in science

- 1 perform experiment
- 2 match model to data, assuming that model is correct
- 3 explore if model is adequate to explain the **data** in one or several aspects:
 - formally **test** null hypothesis that model is “adequate”
 - if not, use **diagnostics** to guide model developments
- 4 revise model, motivating new experiment



From ABC to ABC_{μ} : outline

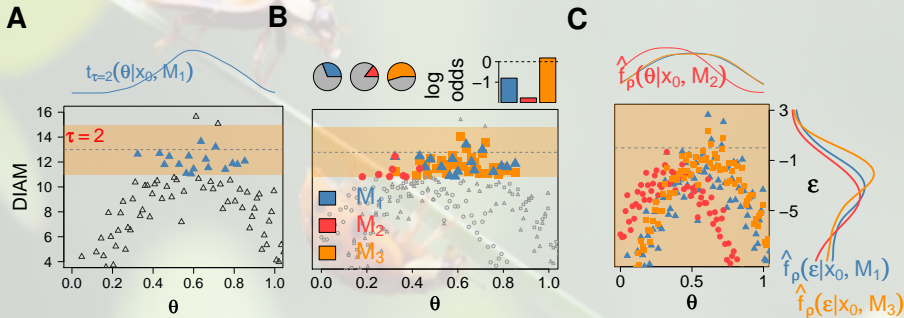
ABC

- exploit model predictions for sampling from approximate posterior densities of θ

ABC_{μ}

- Typically, model predictions form the basis for model criticism [Jeffreys, Th Probability 1961]
- use the data already generated in ABC for this purpose too
- dual use of model predictions reflected in an extension of the state space: we are interested in θ and $\varepsilon_{1:K}$
- to maintain Bayesian framework, crux is to derive a likelihood on the augmented space $(\theta, \varepsilon_{1:K})$.

From ABC to ABC μ : outline



From ABC to ABC μ : notation

- θ random variable, x_0 observed data
- M data-generating stochastic process
- $x \sim f(\cdot|\theta, M)$ simulated data under M given θ

- lower dimensional summaries $\mathbb{S} = \{\mathbf{S}_1, \dots, \mathbf{S}_K\}$
- introduce mismatch thresholds τ_k
- approximate likelihood here with

$$t_{\rho, \tau}(x_0|\theta, M) = \frac{1}{\prod_k \tau_k} \int \mathbf{1} \left\{ \bigcap_{k=1}^K |\mathbf{S}_k(x) - \mathbf{S}_k(x_0)| \leq \tau_k/2 \right\} f(x|\theta, M) dx$$

- apply Bayes' Theorem

$$t_{\rho, \tau}(\theta|x_0, M) = t_{\rho, \tau}(x_0|\theta, M) \pi_{\theta}(\theta|M) / f_{\rho, \tau}(x_0|M)$$

ABC under wrong model: example



How is ABC affected ...

... when the model is not adequate?



ABC under wrong model: example

Simulation study

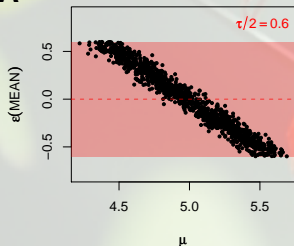
- observe dataset x_0 of 100 sample points, $\bar{x}_0 = 5$,
0.25 Quantile $Q_0 = 1.48$
- we believe $x_i \sim \mathcal{N}(\theta, 1)$ whereas in reality $x_i \sim \text{Exp}(0.2)$
- summarize data with 0.25 Quantile Q and \bar{x}
- consider ABC inference for separate as well as for joint summaries:

$$\rho(\bar{x}_0, Q(x_0); \bar{x}, Q(x)) = \frac{1}{\tau_1} \mathbf{1}\left\{|\bar{x}_0 - \bar{x}| \leq \tau_1/2\right\} \times \\ \frac{1}{\tau_2} \mathbf{1}\left\{|Q(x_0) - Q(x)| \leq \tau_2/2\right\},$$

- reconstruct $t_{\rho, \tau}(\theta | x_0, M)$ with Std-ABC

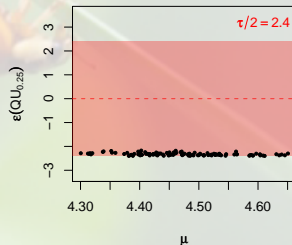
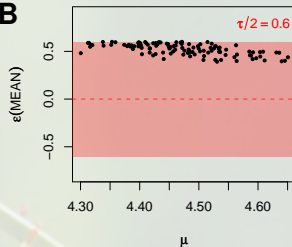
ABC under wrong model: example (cont'd)

A



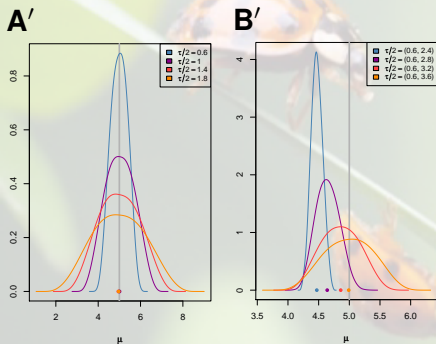
$$\mathcal{S} = \bar{X}$$

B



$$\mathcal{S} = \{\bar{X}, Q\}$$

ABC under wrong model: example (cont'd)



$$\mathbb{S} = \bar{X}$$

$$\mathbb{S} = \{\bar{X}, Q\}$$

- But what is the meaning of posterior estimates of θ ?

make use of the data already generated

Intuitively, want to make information in **B** (computed with ABC) available as job output & interpret; e.g. want to interpret posterior $\varepsilon_{\bar{X}}$ as retained points $\bar{X}_0 - \bar{X}$. Two steps:

- 1 Take ABC kernel as prior $\pi_{\varepsilon_{1:K}}$.
- 2 Propose new likelihood definition that provides desired interpretation.

ABC_μ : Derivation of augmented likelihood



Predictive error of summaries given θ

- **Given θ** , consider the “predictive error” ε with probability distribution

$$\mathbb{P}_{\theta, x_0}(\varepsilon \leq \mathbf{e}) = \int_{\mathcal{X}} \mathbf{1}\{\rho(\mathbb{S}(x), \mathbb{S}(x_0)) \leq \mathbf{e}\} f(x|\theta, M) dx$$

- \mathcal{X} often finite \rightarrow integral replaced by sum
- not difficult to generalize to multiple errors

ABC_μ : Derivation of augmented likelihood

Conditional predictive error density

- Assume $\mathbb{P}_{\theta, x_0}(\varepsilon \leq e)$ has density ξ_{θ, x_0} w.r.t. appropriate measure
- Write ξ_{θ, x_0} as elementary derivative (a. e.)

$$\begin{aligned}
 \xi_{\theta, x_0}(\varepsilon) &= \lim_{h \rightarrow 0} \frac{1}{2h} \int \mathbf{1} \left\{ \rho(\mathbb{S}(x), \mathbb{S}(x_0)) - h < \varepsilon \leq \right. \\
 &\quad \left. \rho(\mathbb{S}(x), \mathbb{S}(x_0)) + h \right\} f(x|\theta, M) dx \\
 &= \lim_{h \rightarrow 0} \int \delta_h \left(\rho(\mathbb{S}(x), \mathbb{S}(x_0)) - \varepsilon \right) f(x|\theta, M) dx \\
 &\stackrel{\text{def}}{=} \int \delta \{ \rho(\mathbb{S}(x), \mathbb{S}(x_0)) = \varepsilon \} f(x|\theta, M) dx
 \end{aligned}$$

ABC μ : Derivation of augmented likelihood

Augmented likelihood on (θ, ε)

- set

$$f_{\rho}(x_0|\theta, \varepsilon, M) \stackrel{\text{def}}{=} \xi_{\theta, x_0}(\varepsilon) = \int \delta\{\rho(\mathbb{S}(x), \mathbb{S}(x_0)) = \varepsilon\} f(x|\theta, M) dx$$

Interpretation

By construction:

$\rho \neq 0 \sim$ discrepancies between data and model

- “expect” mode of $\xi_{\theta, x_0}(\varepsilon) \approx 0$ for some θ
only if model matches data
- if mode “far away” from 0 for all θ ,
detect model mismatch under that discrepancy

Joint posterior density

Joint posterior density of θ and summary errors

Given prior $\pi(\theta, \varepsilon | M)$, apply Bayes' Theorem

$$f_{\rho}(\theta, \varepsilon | x_0, M) = \xi_{\theta, x_0}(\varepsilon) \pi(\theta, \varepsilon | M) / f_{\rho}(x_0 | M)$$

- take $\pi(\theta, \varepsilon | M_i) = \pi_{\varepsilon}(\varepsilon | M) \pi_{\theta}(\theta | M)$
- choose $\pi(\varepsilon | M)$ with mode at zero

“Standard” ABC μ algorithm

“Std”-ABC μ [Ratmann, PNAS 2009]

Std-ABC μ 1 Sample $\theta \sim \pi(\theta|M)$, simulate $x \sim f(\cdot|\theta, M)$ and compute $\varepsilon = \rho(\mathbb{S}(x), \mathbb{S}(x_0))$.

Std-ABC μ 2 Accept (θ, ε, x) with probability proportional to $\pi(\varepsilon|M)$, and go to Std-ABC μ 1.

Cmp'd to Standard-ABC almost unchanged (only record realized errors) & no additional cost

Marginally in (θ, ε) **we obtain samples from $f_\rho(\theta, \varepsilon|x_0, M)$ by construction**. Exploit expression in terms of predictive density to interpret $f_\rho(\theta, \varepsilon|x_0, M)$.

ABC μ : parameter inference

Approximation of true posterior $f(\theta|x_0, M)$

Under regularity assumptions on ξ_{θ, x_0}

$$f_{\rho}(\theta|x_0, M) \propto \pi_{\theta}(\theta|M) \int \pi_{\varepsilon}\left(\rho(\mathbb{S}(x), \mathbb{S}(x_0))\right) f(x|\theta, M) dx$$

- $\pi_{\varepsilon}(\varepsilon|M) = \mathbf{1}\{|\varepsilon| \leq \tau/2\}/\tau \Rightarrow$ “standard” ABC
- may interpret ABC kernel $\frac{1}{\tau}K(\rho(\mathbb{S}(x), \mathbb{S}(x_0))/\tau)$ as particular prior belief on M [Wilkinson R, 2008]

We suggest not to stop at prior interpretation of error ...

ABC_{μ} : model criticism

... Posterior ε is a compound variable that reflects stochastic fluctuations and systematic biases between the model and x_0 .

K real-valued summary errors

In contrast to ABC where error $\rho(\mathbb{S}(x), \mathbb{S}(x_0))$ is scalar & positive,

consider K real-valued summary errors ε_k
that correspond to $\rho_k(\mathbb{S}_k(x), \mathbb{S}_k(x_0))$.

→ gain of interpretability and diagnostic value

ABC μ : model criticism

Formally ...

- define multiple, real-valued prior error terms with density

$$\pi_{\varepsilon_{1:K}}(\varepsilon_{1:K}|M) \stackrel{\text{def}}{=} \prod_{k=1}^K 1/\tau_k K(\varepsilon_k/\tau_k)$$

think: $K(\varepsilon_k/\tau_k) = \mathbf{1}\{|\varepsilon_k| \leq \tau_k/2\}$

- set augmented likelihood to joint predictive density

$$f_{\rho,\tau}(x_0|\theta, \varepsilon_{1:K}, M) \stackrel{\text{def}}{=} \xi_{\theta,x_0}(\varepsilon_{1:K}).$$

ABC μ : model criticism

Interpretation of posterior $\varepsilon_{1:K}$

Under regularity assumptions on ξ_{θ, x_0}

$$\begin{aligned} f_{\rho, \tau}(\varepsilon_{1:K} | x_0, M) &= \int f_{\rho, \tau}(\theta, \varepsilon_{1:K} | x_0, M) d\theta \\ &= \pi_{\varepsilon_{1:K}}(\varepsilon_{1:K} | M) L_{\rho}(\varepsilon_{1:K} | M) / f_{\rho, \tau}(x_0 | M) \end{aligned}$$

where (prior) predictive density [Box, RSSA 1980] of summary errors is

$$L_{\rho}(\varepsilon_{1:K} | M) \stackrel{\text{def}}{=} \lim_{h \rightarrow 0} \int_{\mathcal{X}} \delta_h \left((\rho_k(\mathcal{S}_k(x), \mathcal{S}_k(x_0)) - \varepsilon_k)_{1:K} \right) \pi(x | M) dx$$

and (prior) predictive density of data is

$$\pi(x | M) \stackrel{\text{def}}{=} \int f(x | \theta, M) \pi(\theta | M) d\theta$$

ABC μ : model criticism

$$f_{\rho, \tau}(\varepsilon_{1:K} | x_0, M) = \int f_{\rho, \tau}(\theta, \varepsilon_{1:K} | x_0, M) d\theta \\ \propto L_{\rho}(\varepsilon_{1:K} | M) \pi_{\varepsilon_{1:K}}(\varepsilon_{1:K} | M)$$

- **towards more accurate model criticism**
inspect multiple errors jointly, no problem for Monte Carlo methods
- **“weighted” prior predictive error according to error magnitude**
 - focus on those θ actually inferred
 - criticize posterior model
 - alleviate sensitivity of $L_{\rho}(\varepsilon_{1:K} | M)$ on $\pi_{\theta}(\theta | M)$
- **computationally feasible**

ABC_{μ} : model criticism



Posterior mean shift

[unpublished material taken out]



ABC_{μ} : model criticism

[unpublished material taken out]

k th mean shift re-weighted according to mutual constraints

[unpublished material taken out]

ABC_{μ} : model criticism

Recall: summaries typically co-dependent and not clear if sufficient for θ under M

Exploit co-dependency of summary errors in ABC_{μ}

- If ε_k independent, then mutual constraints collapse [unpublished material taken out]
- Mutual constraints require co-dependent summaries
- Mutual constraints do not require summaries to be sufficient for θ under M

ABC_{μ} reveals model inconsistency with conflicting, co-dependent summaries

ABC_μ under wrong model: example 2

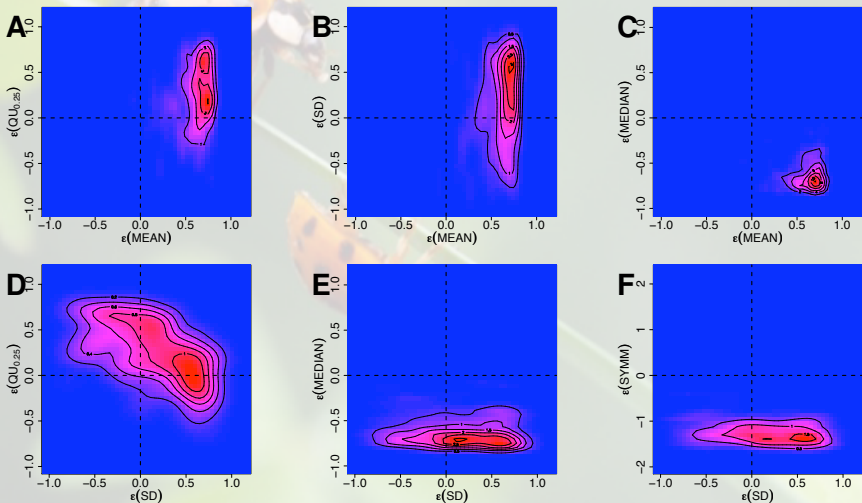


Simulation study

- observe dataset x_0 of 100 sample points, $\bar{x}_0 = 5$,
0.25 Quantile $Q_0 = 1.48$
- we believe $x_i \sim \mathcal{N}(\mu, \sigma^2)$ whereas in reality $x_i \sim \text{Exp}(0.2)$
[unpublished material taken out]

ABC_{μ} under wrong model: example 2

- Std- ABC_{μ} with 5 summary errors



Concluding remarks 1

Utility of posterior summary errors $\varepsilon_{1:K}$

- Exploit the data already generated in ABC also for model criticism
- Formally, dual use of simulated data is described by the joint posterior density

$$f_{\rho, \tau}(\theta, \varepsilon_{1:K} | x_0, M)$$

that rests on augmenting the likelihood in a particular way

- Marginally in θ , we perform approximate inference exactly as in ABC
- Marginally in $\varepsilon_{1:K}$, we criticize the **fitted** model with all summaries **simultaneously**

Concluding remarks 2

Choice of summary errors $\varepsilon_{1:K}$

- we always work with K errors that correspond to K summaries
- we always include as many summaries as possible to preserve co-dependencies;
compare to [Joyce P, Stat App Gen Mol Biol 2008]
- we do not combine summaries into scalar error
- we always choose τ so that co-dependencies are preserved

Concluding remarks 3



Diagnostics and hypothesis testing

- we use high probability density intervals of marginals $f_{\rho, \tau}(\varepsilon_k | x_0, M)$ to obtain precise information as to how improve a model
- these are not confidence intervals, and marginal posterior errors are typically not independent
- to test $H_0: \varepsilon_{1:K} = 0$ we compute a Bayes factor [unpublished material taken out]

Concluding remarks 4

Std-ABC $_{\mu}$ does not work well in other than very simple settings

- Prob1: $f_{\rho,\tau}(\theta, \varepsilon_{1:K} | x_0, M)$ typically very different from $\pi_{\theta}(\theta | M) \prod_{k=1}^K \pi_{\varepsilon_k}(\varepsilon_k | M)$
- Prob2: By simulating $x \sim f(\cdot | \theta, M)$, we extend the state space $(\theta, \varepsilon_{1:K})$ with x . The volatility of the stochastic process $f(\cdot | \theta, M)$ induces an extra price to pay.
- Prob3: Compared to scalar error, acceptance probability of ABC $_{\mu}$ based on several summary errors $\varepsilon_{1:K}$ is further reduced.

But straightforward to modify existing methods of ABC. We also implement a novel ABC $_{\mu}$ algorithm ...

Concluding remarks 5

- We use **repeated sampling** $x_b \sim f(\cdot|\theta, M)$, $b = 1, \dots, B$ to control the volatility of the data generating process. [Andrieu C, Ann Stat 2009]
- We also replaced the joint likelihood $\xi_{\theta, x_0}(\varepsilon_{1:K})$ with a conservative **combination of its marginals** $\min_k \xi_{k, \theta, x_0}(\varepsilon_k)$ [Ratmann O, PNAS 2009]
- For our applications, we found **MCMC samplers (with annealing & refined proposals)** to work well [Ratmann O, PLOS CompBiol 2007]

Two ladybugs are shown on a green stem. One is positioned higher and facing right, while the other is lower and facing left. The background is a soft-focus green.

Thank you

Sylvia Richardson, Christophe Andrieu, Carsten Wiuf

and

Wellcome Trust