

Bayesian inference for stochastic differential mixed effects models - initial steps

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Supervisors: RJB and AG

- ▶ Mixed Effects Stochastic Differential Equations (SDEs)
- ▶ Bayesian inference for SDEs
- ▶ Toy model (Roberts and Stramer 2001 paper)

- ▷ Consider an Itô process $\{\mathbf{X}_t, t \geq 0\}$ satisfying

$$d\mathbf{X}_t = \boldsymbol{\alpha}(\mathbf{X}_t, \boldsymbol{\theta})dt + \sqrt{\boldsymbol{\beta}(\mathbf{X}_t, \boldsymbol{\theta})}d\mathbf{W}_t$$

- \mathbf{X}_t is the value of the process at time t
- $\boldsymbol{\alpha}(\mathbf{X}_t, \boldsymbol{\theta})$ is the drift
- $\boldsymbol{\beta}(\mathbf{X}_t, \boldsymbol{\theta})$ is the diffusion coefficient
- \mathbf{W}_t is standard Brownian motion
- \mathbf{X}_0 is the vector of initial conditions

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- ▷ Seek a numerical solution via (for example) the Euler-Maruyama approximation

$$\Delta\mathbf{X}_t \equiv \mathbf{X}_{t+\Delta t} - \mathbf{X}_t = \boldsymbol{\alpha}(\mathbf{X}_t, \boldsymbol{\theta})\Delta t + \sqrt{\boldsymbol{\beta}(\mathbf{X}_t, \boldsymbol{\theta})}\Delta\mathbf{W}_t$$

where $\Delta\mathbf{W}_t \sim N(\mathbf{0}, \mathbf{I}\Delta t)$

CIR Model

$$dX_t = (\theta_1 - \theta_2 X_t)dt + \theta_3 \sqrt{X_t} dW_t$$

- ▶ Used to model short term interest rates
- ▶ The process is mean reverting

CIR Model

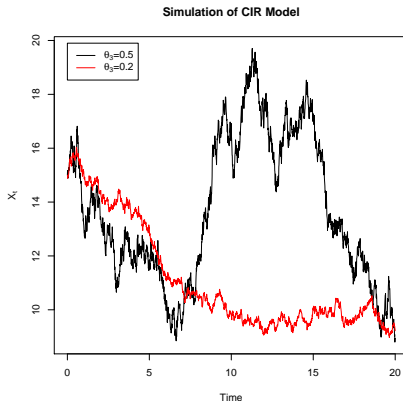


Figure: Numerical solution for CIR model, $\theta_1 = 1$, $\theta_2 = 0.1$, $X_0 = 15$

Aphid Growth Model



- ▷ Also known as plant lice, or greenfly
- ▷ They are small sap sucking insects
- ▷ Some species of ants “farm” aphids, for the honeydew they release. These “dairying ants”, “milk” the aphids by stroking them

Aphid Growth Model

$$\begin{pmatrix} dN_t \\ dC_t \end{pmatrix} = \begin{pmatrix} \lambda N_t - \mu N_t C_t \\ \lambda N_t \end{pmatrix} dt + \begin{pmatrix} \lambda N_t + \mu N_t C_t & \lambda N_t \\ \lambda N_t & \lambda N_t \end{pmatrix}^{\frac{1}{2}} d\mathbf{W}_t$$

- ▶ N_t is the aphid population size at time t
- ▶ C_t is the cumulative population at time t
- ▶ This model is an SDE approximation to an underlying stochastic kinetic model
- ▶ Birth rate of λN_t and a death rate of $\mu N_t C_t$

Aphid Growth Model

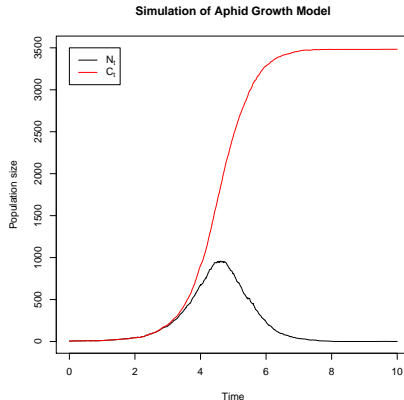


Figure: Numerical solution for Aphid model, $\lambda = 1.75$, $\mu = 0.001$

Mixed Effects SDE Models

- ▷ What if experimental units are not identical?
- ▷ Suppose the units have common parameters θ but different parameters \mathbf{b}^i
- ▷ We treat the \mathbf{b}^i as random effects with a population profile

Mixed Effects SDE Models

- ▶ What if experimental units are not identical?
- ▶ Suppose the units have common parameters θ but different parameters \mathbf{b}^i
- ▶ We treat the \mathbf{b}^i as random effects with a population profile
- ▶ This gives us a stochastic differential mixed-effects model for the experimental units:

$$d\mathbf{X}_t^i = \boldsymbol{\alpha}(\mathbf{X}_t^i, \boldsymbol{\theta}, \mathbf{b}^i)dt + \sqrt{\boldsymbol{\beta}(\mathbf{X}_t^i, \boldsymbol{\theta}, \mathbf{b}^i)}d\mathbf{W}_t^i, \quad i = 1, \dots, M$$

- ▶ Differences between units are down to different realisations of the Brownian motion paths \mathbf{W}_t^i and the random effects \mathbf{b}^i
- ▶ Allows us to split the total variation between within- and between-individual components

Bayesian inference for SDEs

- ▶ Problematic due to the intractability of the transition density characterising the process
- ▶ In other words, we typically can't solve an SDE analytically
- ▶ So we could just work with the Euler approximation
- ▶ Given data \mathbf{d} at equidistant times t_0, t_1, \dots, t_n , the Euler approximation may be unsatisfactory for $\Delta t = t_{i+1} - t_i$
- ▶ We therefore adopt a data augmentation scheme

- ▷ Introduce a partition of $[t_i, t_{i+1}]$ as

$$t_i = \tau_{im} < \tau_{im+1} < \dots < \tau_{(i+1)m} = t_{i+1}$$

where

$$\Delta\tau \equiv \tau_{im+1} - \tau_{im} = \frac{t_{i+1} - t_i}{m}$$

- ▷ Apply Euler approximation over each interval of width $\Delta\tau$
- ▷ Introduces $m - 1$ latent values between every pair of observations

- ▷ Formulate joint posterior for parameters and latent values

$$\rightarrow \quad \mathbf{d} \quad = (\mathbf{x}_{t_0}, \mathbf{x}_{t_1}, \dots, \mathbf{x}_{t_n})$$

$$\rightarrow \quad \mathbf{x} \quad = (\mathbf{x}_{\tau_1}, \mathbf{x}_{\tau_2}, \dots, \mathbf{x}_{\tau_{m-1}}, \mathbf{x}_{\tau_{m+1}}, \dots, \dots, \mathbf{x}_{\tau_{nm-1}})$$

= latent path

$$\rightarrow \quad (\mathbf{x}, \mathbf{d}) \quad = (\mathbf{x}_{\tau_0}, \mathbf{x}_{\tau_1}, \dots, \mathbf{x}_{\tau_m}, \mathbf{x}_{\tau_{m+1}}, \dots, \dots, \mathbf{x}_{\tau_{nm}})$$

= augmented path

Formulate joint posterior for parameters and latent data as

$$\begin{aligned}\pi(\boldsymbol{\theta}, \boldsymbol{x}|\boldsymbol{d}) &\propto \pi(\boldsymbol{\theta})\pi(\boldsymbol{x}, \boldsymbol{d}|\boldsymbol{\theta}) \\ &\propto \pi(\boldsymbol{\theta}) \prod_{i=0}^{nm-1} \pi(\boldsymbol{x}_{\tau_{i+1}}|\boldsymbol{x}_{\tau_i}, \boldsymbol{\theta})\end{aligned}$$

where

$$\pi(\boldsymbol{x}_{\tau_{i+1}}|\boldsymbol{x}_{\tau_i}, \boldsymbol{\theta}) = \phi(\boldsymbol{x}_{\tau_{i+1}}; \boldsymbol{x}_{\tau_i} + \boldsymbol{\alpha}(\boldsymbol{x}_{\tau_i}, \boldsymbol{\theta})\Delta t, \boldsymbol{\beta}(\boldsymbol{x}_{\tau_i}, \boldsymbol{\theta})\Delta t)$$

and $\phi(\cdot; \boldsymbol{\mu}, \boldsymbol{\Sigma})$ denotes the Gaussian density with mean $\boldsymbol{\mu}$ and variance $\boldsymbol{\Sigma}$

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- ▶ The posterior distribution is typically analytically intractable

A Gibbs sampling approach

- ▷ We therefore sample via an MCMC scheme
- ▷ E.g a Gibbs sampler, alternating between draws of
 - $\theta|x, d$
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- ▷ E.g a Gibbs sampler, alternating between draws of
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- ▷ The last step can be done (for example) in blocks of length $m - 1$ between observations
- ▷ Metropolis within Gibbs updates may be needed
- ▷ **Problem:** the mixing is poor for large m

Consider the SDE

$$d\mathbf{X}_t = \frac{1}{\sqrt{\theta}} d\mathbf{W}_t$$

- ▶ Suppose that we have observations $X_0 = x_0 = 0$ and $X_1 = x_1$
- ▶ Set $\tau_i = i/m$ for $i = 0, 1, \dots, m$ so that

$$(\mathbf{x}, \mathbf{d}) = x_0, x_{\frac{1}{m}}, x_{\frac{2}{m}}, \dots, x_{\frac{m-1}{m}}, x_1$$

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- ▶ Under the Euler approximation

$$X_{\frac{i}{m}} | x_{\frac{(i-1)}{m}}, \theta \sim N \left(x_{\frac{(i-1)}{m}}, \frac{1}{m\theta} \right)$$

▷ Hence

$$\begin{aligned}\pi(\mathbf{x}, \mathbf{d}|\theta) &\propto \prod_{i=1}^m \frac{\sqrt{m\theta}}{\sqrt{2\pi}} \exp \left\{ -\frac{m\theta \left(x_{\frac{i}{m}} - x_{\frac{(i-1)}{m}} \right)^2}{2} \right\} \\ &\propto \theta^{m/2} \exp \left\{ -\frac{1}{2} m\theta \sum_{i=1}^m \left(x_{\frac{i}{m}} - x_{\frac{(i-1)}{m}} \right)^2 \right\}\end{aligned}$$

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▷ Take prior $\theta \sim \text{Exp}(1)$

▷ The full conditional for θ is

$$\pi(\theta|\mathbf{x}, \mathbf{d}) \propto \pi(\theta)\pi(\mathbf{x}, \mathbf{d}|\theta)$$

$$\begin{aligned} &\propto \theta^{m/2} \exp \left\{ -\frac{1}{2} m \theta \sum_{i=1}^m \left(x_{\frac{i}{m}} - x_{\frac{(i-1)}{m}} \right)^2 - \theta \right\} \\ &\propto \theta^{m/2} \exp \left\{ -\theta \left(\frac{m \Sigma_X}{2} + 1 \right) \right\} \end{aligned}$$

where

$$\Sigma_X = \sum_{i=1}^m \left(x_{\frac{i}{m}} - x_{\frac{(i-1)}{m}} \right)^2$$

$$\begin{aligned} &\propto \theta^{m/2} \exp \left\{ -\frac{1}{2} m \theta \sum_{i=1}^m \left(x_{\frac{i}{m}} - x_{\frac{(i-1)}{m}} \right)^2 - \theta \right\} \\ &\propto \theta^{m/2} \exp \left\{ -\theta \left(\frac{m \Sigma_X}{2} + 1 \right) \right\} \end{aligned}$$

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► Therefore

$$\theta | \mathbf{x}, \mathbf{d} \sim \Gamma \left(\frac{m}{2} + 1, \frac{m \Sigma_X}{2} + 1 \right)$$

- ▶ Under the linear Gaussian structure of the simple SDE, the full conditional $x|\theta, d$ can be sampled using

$$X_{\frac{i}{m}}|x_1, \theta = \frac{ix_1}{m} + \frac{1}{\sqrt{\theta}}Z_{\frac{i}{m}}, \quad i = 1, 2, \dots, m$$

where $\{Z_t, 0 \leq t \leq 1\}$ is a standard Brownian bidge, that is a standard Brownian motion conditioned to hit 0 at time 0, at time 1

Simulated data

- ▷ Take $x_0 = 0$, $\theta = 1$
- ▷ Simulate x_1 using Euler scheme
- ▷ Get $x_1 = -0.6947 \Rightarrow \mathbf{d} = (0, -0.6947)$

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MCMC scheme

We perform a run of 1000 iterations of the scheme with no thin. Initialise with $\theta^{(0)} = 1$, the prior mean

Step 1 Update the discretised Brownian bridge which hits x_1 at $t = 1$

Step 2 Draw θ from its full conditional distribution,

$$\theta | \mathbf{x}, \mathbf{d} \sim \Gamma \left(\frac{m}{2} + 1, \frac{m \Sigma_X}{2} + 1 \right)$$

Toy model – Results

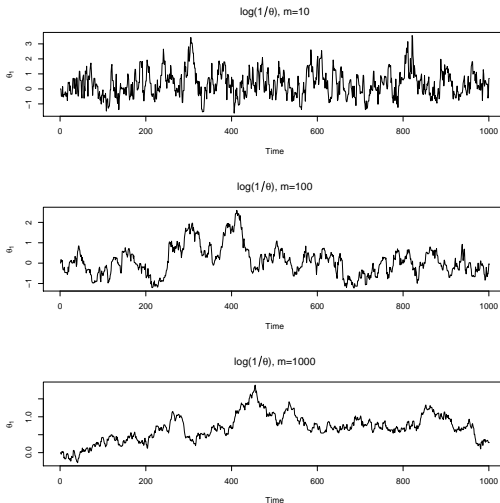


Figure: Trace plots for $\log(\frac{1}{\theta})$

Toy model – Results

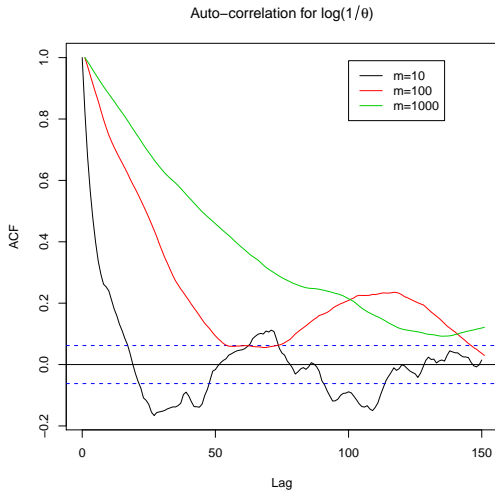


Figure: Auto-correlation plots for $\log(\frac{1}{\theta})$

Toy model – Results

- ▷ Mixing gets even worse for larger m
- ▷ Why is this happening?
- ▷ Try and quantify the mixing time by considering the parameter update
- ▷ The parameter update can be rewritten in a clever way ...

Toy model – What's going wrong?

▷ Recall that

$$X_{\frac{i}{m}} | x_1, \theta = \frac{ix_1}{m} + \frac{1}{\sqrt{\theta}} Z_{\frac{i}{m}}, \quad i = 1, 2, \dots, m$$

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▷ Now

$$\begin{aligned} \Sigma_X &= \sum_{i=1}^m \left(\frac{i}{m} x_1 + \frac{1}{\sqrt{\theta}} Z_{\frac{i}{m}} - \left[\frac{(i-1)}{m} x_1 + \frac{1}{\sqrt{\theta}} Z_{\frac{(i-1)}{m}} \right] \right)^2 \\ &= \sum_{i=1}^m \left(\frac{1}{\sqrt{\theta}} \left[Z_{\frac{i}{m}} - Z_{\frac{(i-1)}{m}} \right] + \frac{i}{m} x_1 - \frac{(i-1)}{m} x_1 \right)^2 \\ &= \sum_{i=1}^m \left(\frac{1}{\sqrt{\theta}} \left[Z_{\frac{i}{m}} - Z_{\frac{(i-1)}{m}} \right] + \frac{x_1}{m} \right)^2 \end{aligned}$$

Toy model – What's going wrong?

▷ Expanding out

$$\begin{aligned}\Sigma_X &= \sum_{i=1}^m \left(\frac{1}{\theta} \left[Z_{\frac{i}{m}} - Z_{\frac{(i-1)}{m}} \right]^2 + \frac{x_1^2}{m^2} + \frac{2}{\sqrt{\theta}} \frac{x_1}{m} \left[Z_{\frac{i}{m}} - Z_{\frac{(i-1)}{m}} \right] \right) \\ &= \frac{1}{\theta} \Sigma_Z + \frac{x_1^2}{m} + \frac{2}{\sqrt{\theta}} \frac{x_1}{m} \sum_{i=1}^m \left[Z_{\frac{i}{m}} - Z_{\frac{(i-1)}{m}} \right]\end{aligned}$$

▷ Now

$$\begin{aligned}\sum_{i=1}^m \left[Z_{\frac{i}{m}} - Z_{\frac{(i-1)}{m}} \right] &= \left(Z_{\frac{1}{m}} - Z_0 \right) + \left(Z_{\frac{2}{m}} - Z_{\frac{1}{m}} \right) + \dots \\ &\quad \dots + \left(Z_{\frac{(m-1)}{m}} - Z_{\frac{(m-2)}{m}} \right) + \left(Z_1 - Z_{\frac{(m-1)}{m}} \right)\end{aligned}$$

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Toy model – What's going wrong?

▷ So

$$\Sigma_X = \frac{1}{\theta} \Sigma_Z + \frac{x_1^2}{m} + \frac{2}{\sqrt{\theta}} \frac{x_1}{m} [Z_1 - Z_0]$$

But $Z_0 = Z_1 = 0$ since Z is a standard Brownian bridge

Toy model – What's going wrong?

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where

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▷ Using properties of Gaussian random variables we have

$$\Sigma_Z \sim \frac{\chi_{m-1}^2}{m}$$

Toy model – What's going wrong?

▷ Now

$$\theta|\mathbf{x}, \mathbf{d} \sim \Gamma\left(\frac{m}{2} + 1, \frac{m\Sigma_X}{2} + 1\right)$$

▷ If $H \sim \Gamma\left(\frac{m}{2} + 1, 1\right)$ then

$$\begin{aligned}\theta_{\text{new}} &= \frac{H}{\frac{m\Sigma_X}{2} + 1} \\ &= \frac{H}{\frac{m}{2} \left(\frac{x_1^2}{m} + \frac{\chi_{m-1}^2}{m\theta_{\text{old}}} \right) + 1}\end{aligned}$$

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Toy model – What's going wrong?

- ▶ For large m , approximate H and χ_{m-1}^2 with Normal random variables
- ▶ Roberts and Stramer then use a suitable Taylor expansion of the expression for θ_{new} to give

$$\theta_{\text{new}} \approx \theta_{\text{old}} \left\{ 1 + \left(\frac{2}{m} \right)^{\frac{1}{2}} (W_1 - W_2) - \frac{2 + x_1^2}{m} \theta_{\text{old}} + \frac{W_2^2}{m} \right\}$$

where W_1 and W_2 are independent $N(0, 1)$ random variables

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- ▶ $\theta_{\text{new}} = \theta_{\text{old}} \{1 + O(m^{-1})\}$

\Rightarrow Mixing time is $O(m)$

- ▷ Construct MCMC schemes for arbitrary nonlinear diffusion processes
 - Naive schemes with a block update for the path
 - “Better” schemes that use a reparameterisation
 - Joint update of path and parameters (pMCMC)
- ▷ Application to mixed effects SDEs
 - Aphid model, real data examples

- ▶ Roberts. G. O. and Stramer. O.
On inference for partially observed nonlinear diffusion models using the Metropolis-Hastings algorithm.
Biometrika, 88 (3) 603-621, 2001
- ▶ Gillespie, C. S. and Golightly, A.
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JRSS Series C, Applied Statistics, 59(2):341-357, 2010