

MCMC Sampling for Intractable MJP Models of Chemical Kinetics via the Linear Noise Approximation

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Outline

- 1 Motivating Example and Background
- 2 Diffusion Approximation
- 3 Linear Noise Approximation
- 4 Challenges for Statistical Inference
- 5 Differential Geometric Monte Carlo
- 6 MCMC with Riemann Manifold Methods
- 7 Illustrative Experiment for Gene Autoregulation
- 8 Conclusions





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$$\frac{d\boldsymbol{p}(\boldsymbol{x},t|\boldsymbol{x}_{0},t_{0})}{dt} = \sum_{j=1}^{M} \left[f_{j}(\boldsymbol{x}-\boldsymbol{s}_{j},\boldsymbol{\theta},t)\boldsymbol{p}(\boldsymbol{x}-\boldsymbol{s}_{j},t|\boldsymbol{x}_{0},t_{0}) - f_{j}(\boldsymbol{x},\boldsymbol{\theta},t)\boldsymbol{p}(\boldsymbol{x},t|\boldsymbol{x}_{0},t_{0}) \right].$$

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- Parameter inference scheme Boys, Wilkinson, Kirkwood (2006)



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- Langevin Equation

$$d\boldsymbol{x}_{t} = \boldsymbol{S}\boldsymbol{f}(\boldsymbol{x}_{t},\theta)dt + \frac{1}{\sqrt{\Omega}}\boldsymbol{S}\sqrt{\operatorname{diag}(\boldsymbol{f}(\boldsymbol{x}_{t},\theta))}dB_{t}$$
(3)



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- Employed extensively in chemical physics and genome research
- · May provide intermediate scheme for MCMC based inference



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· which is a linear SDE with analytic solution

$$\boldsymbol{\xi}_{t} = \boldsymbol{\Phi}(t_{0}, t) \left(\boldsymbol{\xi}_{0} + \int_{t_{0}}^{t} \boldsymbol{\Phi}(s, t)^{-1} \boldsymbol{S} \sqrt{\text{diag}(\boldsymbol{f}(\boldsymbol{\phi}_{s}, \boldsymbol{\theta}))} d\boldsymbol{B}_{s} \right)$$

where Φ(t0, s) the solution to

$$d\Phi(t0,s) = \boldsymbol{SJ}_{\boldsymbol{f}}(\phi_s,\theta)\Phi(t0,s)ds, \ \Phi(t0,t0) = \boldsymbol{I}$$



Likelihood for the Linear Noise Approximation

$p(\pmb{x}^{(TS)}|\pmb{ heta}) \propto \mathcal{N}(\pmb{\mu}(\pmb{ heta}), \pmb{\Sigma}(\pmb{ heta}))$

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$$\boldsymbol{x}_t \sim \mathcal{N}(\boldsymbol{\phi}_t, \boldsymbol{V}_t), \ \frac{d \boldsymbol{V}_t}{dt} = \boldsymbol{S} \boldsymbol{J}_f(\boldsymbol{\phi}_t, \boldsymbol{\theta}) \boldsymbol{V}_t + \boldsymbol{V}_t \boldsymbol{J}_f(\boldsymbol{\phi}_t, \boldsymbol{\theta})^T \boldsymbol{S}^T + \boldsymbol{S} \text{diag}(\boldsymbol{f}(\boldsymbol{x}_t, \boldsymbol{\theta})) \boldsymbol{S}^T$$



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Likelihood for the Linear Noise Approximation

 $p(\mathbf{x}^{(TS)}|\mathbf{ heta}) \propto \mathcal{N}(\mu(\mathbf{ heta}), \mathbf{\Sigma}(\mathbf{ heta}))$

• $\mu(\theta) = (\phi_{t1}, \dots, \phi_{tn})^T$ a *nN* vector with solutions of the MRE.

- $\Sigma(\theta)$ a $nN \times nN$ block matrix with blocks $\Sigma(\theta)^{i,j} N \times N$
 - $\boldsymbol{x}_{t} \sim \mathcal{N}(\boldsymbol{\phi}_{t}, \boldsymbol{V}_{t}), \ \frac{d\boldsymbol{V}_{t}}{dt} = \boldsymbol{S}\boldsymbol{J}_{\boldsymbol{f}}(\boldsymbol{\phi}_{t}, \boldsymbol{\theta})\boldsymbol{V}_{t} + \boldsymbol{V}_{t}\boldsymbol{J}_{\boldsymbol{f}}(\boldsymbol{\phi}_{t}, \boldsymbol{\theta})^{\mathsf{T}}\boldsymbol{S}^{\mathsf{T}} + \boldsymbol{S}\text{diag}(\boldsymbol{f}(\boldsymbol{x}_{t}, \boldsymbol{\theta}))\boldsymbol{S}^{\mathsf{T}}$

$$\boldsymbol{\Sigma}(\boldsymbol{\theta})^{i,j} = \begin{cases} \boldsymbol{V}_{t_i} & \text{if } i = j \\ cov(\boldsymbol{x}_{t_i}, \boldsymbol{x}_{t_j}) = \boldsymbol{\Sigma}(\boldsymbol{\theta})^{i,j-1} \boldsymbol{\Phi}(t_{j-1}, t_j)^T \end{cases}$$



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Fisher Information

$$FI(\theta)_{m,n} = \frac{\partial \mu(\theta)^{T}}{\partial \theta_{m}} \boldsymbol{\Sigma}^{-1}(\theta) \frac{\partial \mu(\theta)}{\partial \theta_{n}} + \frac{1}{2} \operatorname{tr} \left(\boldsymbol{\Sigma}^{-1}(\theta) \frac{\partial \boldsymbol{\Sigma}(\theta)}{\partial \theta_{m}} \boldsymbol{\Sigma}^{-1}(\theta) \frac{\partial \boldsymbol{\Sigma}(\theta)}{\partial \theta_{n}} \right)$$

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Likelihood for the Linear Noise Approximation

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 Augment the MRE for φ with the lower triangular elements of V and solve the augmented system with forward sensitivity analysis.







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Mechanistic modelling



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Figure: A diagram of the Goodwin Circadian Oscillator model network.

$$\begin{aligned} \frac{dx_1}{dt} &= \frac{k_1}{1+x_n^0} - m_1 x_1 \\ \frac{dx_2}{dt} &= k_2 x_1 - m_2 x_2 \\ \vdots \\ \frac{dx_n}{dt} &= k_n x_{n-1} - m_n x_n \end{aligned}$$







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Figure: A diagram of the Goodwin Circadian Oscillator model network.

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Figure: A diagram of the Goodwin Circadian Oscillator model network.

Dependence
on kinetic
parameters

$$\rightarrow \theta$$
 $\frac{dx_1}{dt} = \frac{k_0}{1 + x_n^{h}} - (m)x_1$
 $\frac{dx_2}{dt} = (k_0x_1 - (m_0x_2))$
 \vdots
 $\frac{dx_n}{dt} = (k_0x_{n-1} - (m_0x_n))$



Necessary condition for stable oscillations



Figure: A diagram of the Goodwin Circadian Oscillator model network.



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Example: log likelihood landscape



Figure: A diagram of the Goodwin Circadian Oscillator model network.

$$\begin{pmatrix} \frac{dx_1}{dt} &= \frac{k_1}{1+x_n^p} - m_1 x_1\\ \frac{dx_2}{dt} &= k_2 x_1 - m_2 x_2\\ \vdots\\ \frac{dx_n}{dt} &= k_n x_{n-1} - m_n x_n \end{pmatrix}$$



Example: log likelihood landscape



Figure: A diagram of the Goodwin Circadian Oscillator model network.





Posterior landscape $P(\theta|\mathcal{M}, \mathcal{D})$



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MCMC trajectories

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Journal of the Royal Statistical Society



J. R. Statist. Soc. B (2011) 73, Part 2, pp. 123–214

Riemann manifold Langevin and Hamiltonian Monte Carlo methods

Mark Girolami and Ben Calderhead

University College London, UK

[Read before The Royal Statistical Society at a meeting organized by the Research Section on Wednesday, October 13th, 2010, Professor D. M. Titterington in the Chair]

Summary. The paper proposes Metropolis adjusted Langevin and Hamiltonian Monte Carlo sampling methods defined on the Riemann manifold to resolve the shortcomings of existing Monte Carlo algorithms when sampling from target densities that may be high dimensional and exhibit strong correlations. The methods provide fully automated adaptation mechanisms that circumvent the costly pilot runs that are required to tune proposal densities for Metropolis-Hastings or indeed Hamiltonian Monte Carlo and Metropolis adjusted Langevin algorithms. This allows for highly efficient sampling even in very high dimensions where different scalings may be required for the transient and stationary phases of the Markov chain. The methodology proposed exploits the Riemann geometry of the parameter space of statistical models and thus automat-







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Geometric Concepts in MCMC



• Tangent space - local metric defined by $\delta \theta^{\mathsf{T}} \mathbf{G}(\theta) \delta \theta = g_{kl} \delta \theta_k \delta \theta_l$



Geometric Concepts in MCMC



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- Christoffel symbols characterise Levi-Civita connection on manifold





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$$\Gamma_{kl}^{i} = \frac{1}{2} \sum_{m} g^{im} \left(\frac{\partial g_{mk}}{\partial \theta^{l}} + \frac{\partial g_{ml}}{\partial \theta^{k}} - \frac{\partial g_{kl}}{\partial \theta^{m}} \right)$$





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· Geodesics - shortest path between two points on manifold





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$$\frac{d^2\theta^i}{dt^2} + \sum_{k,l} \Gamma^i_{kl} \frac{d\theta^k}{dt} \frac{d\theta^l}{dt} = 0$$

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Illustration of Geometric Concepts

• Consider Normal density $p(x|\mu, \sigma) = \mathcal{N}_x(\mu, \sigma)$



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Illustration of Geometric Concepts

- Consider Normal density p(x|μ, σ) = N_x(μ, σ)
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- Metric on tangent space

$$\delta \boldsymbol{\theta}^{\mathsf{T}} \mathbf{G}(\boldsymbol{\theta}) \delta \boldsymbol{\theta} = \frac{(\delta \mu^2 + 2\delta \sigma^2)}{\sigma^2}$$

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Illustration of Geometric Concepts

- Consider Normal density p(x|μ, σ) = N_x(μ, σ)
- Local inner product on tangent space defined by metric tensor, i.e. $\delta \theta^{\mathsf{T}} \mathbf{G}(\theta) \delta \theta$, where $\theta = (\mu, \sigma)^{\mathsf{T}}$
- Metric is Expected Fisher Information

$$\mathbf{G}(\mu,\sigma)=\left[egin{array}{cc} \sigma^{-2} & \mathbf{0}\ \mathbf{0} & 2\sigma^{-2} \end{array}
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- Components of connection $\partial_{\mu}\mathbf{G} = \mathbf{0}$ and $\partial_{\sigma}\mathbf{G} = -\operatorname{diag}(2\sigma^{-3}, 4\sigma^{-3})$
- Metric on tangent space

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Metric tensor for univariate Normal defines a Hyperbolic Space


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- Metric tensor for univariate Normal defines a Hyperbolic Space
- Consider densities $\mathcal{N}(0,1)$ & $\mathcal{N}(1,1)$ and $\mathcal{N}(0,2)$ & $\mathcal{N}(1,2)$



Normal Density - Euclidean Parameter space



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Normal Density - Riemannian Functional space





M.C. Escher, Heaven and Hell, 1960





Langevin Diffusion on Riemannian manifold

• Discretised Langevin diffusion on manifold defines proposal mechanism

$$\boldsymbol{\theta}_{d}^{\prime} = \boldsymbol{\theta}_{d} + \frac{\epsilon^{2}}{2} \left(\boldsymbol{G}^{-1}(\boldsymbol{\theta}) \nabla_{\boldsymbol{\theta}} \mathcal{L}(\boldsymbol{\theta}) \right)_{d} - \epsilon^{2} \sum_{i,j}^{D} \boldsymbol{G}(\boldsymbol{\theta})_{ij}^{-1} \Gamma_{ij}^{d} + \epsilon \left(\sqrt{\mathbf{G}^{-1}(\boldsymbol{\theta})} \mathbf{z} \right)_{d}$$



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Langevin Diffusion on Riemannian manifold

· Discretised Langevin diffusion on manifold defines proposal mechanism

$$\boldsymbol{\theta}_{d}^{\prime} = \boldsymbol{\theta}_{d} + \frac{\epsilon^{2}}{2} \left(\boldsymbol{G}^{-1}(\boldsymbol{\theta}) \nabla_{\boldsymbol{\theta}} \mathcal{L}(\boldsymbol{\theta}) \right)_{d} - \epsilon^{2} \sum_{i,i}^{D} \boldsymbol{G}(\boldsymbol{\theta})_{ij}^{-1} \Gamma_{ij}^{d} + \epsilon \left(\sqrt{\boldsymbol{G}^{-1}(\boldsymbol{\theta})} \boldsymbol{z} \right)_{d}$$

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MALA proposal with preconditioning

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Proposal mechanism diffuses approximately along the manifold









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Simplified Manifold MALA

- For $\boldsymbol{\theta} \in \mathbb{R}^D$ with density $\pi(\boldsymbol{\theta}), \, \mathcal{L}(\boldsymbol{\theta}) \equiv \log \pi(\boldsymbol{\theta})$
- Define proposal distribution

$$q(\boldsymbol{\theta}^*|\boldsymbol{\theta}^{t-1}) = \mathcal{N}\left(\boldsymbol{\theta}^*|\boldsymbol{\mu}(\boldsymbol{\theta}^{t-1},\epsilon),\epsilon^2 \mathbf{G}^{-1}(\boldsymbol{\theta}^{t-1})\right)$$



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Simplified Manifold MALA

- For $\boldsymbol{\theta} \in \mathbb{R}^{D}$ with density $\pi(\boldsymbol{\theta}), \mathcal{L}(\boldsymbol{\theta}) \equiv \log \pi(\boldsymbol{\theta})$
- Define proposal distribution

$$q(\boldsymbol{\theta}^*|\boldsymbol{\theta}^{t-1}) = \mathcal{N}\left(\boldsymbol{\theta}^*|\boldsymbol{\mu}(\boldsymbol{\theta}^{t-1},\epsilon),\epsilon^2 \mathbf{G}^{-1}(\boldsymbol{\theta}^{t-1})\right)$$

Where

$$\mu(\theta,\epsilon) = \theta + \frac{\epsilon^2}{2} \left(\mathbf{G}^{-1}(\theta) \nabla_{\theta} \mathcal{L}(\theta) \right), \quad \mathbf{G}^{-1}(\theta) = Fl(\theta)$$



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• Accept θ^* with probability

$$\min\left\{1, \frac{\pi(\boldsymbol{\theta}^*)}{\pi(\boldsymbol{\theta}^{t-1})} \frac{q(\boldsymbol{\theta}^{t-1}|\boldsymbol{\theta}^*)}{q(\boldsymbol{\theta}^*|\boldsymbol{\theta}^{t-1})}\right\}$$

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Single gene expression model



• Single gene expression model with negative feedback.



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- Single gene expression model with negative feedback.
- System state $(R(t), P(t))^T$ models the population of RNA and protein.



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Simulated Data



- Simulated data generated with SSA.
- 10 independent sample paths for each time point.
- · Parameters set to

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Trace Plots



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Effective Sample Size

	10,000 posterior samples							
	γ_R	γ_P	k _P	b_0	b_1	b_2	b_3	
RMHMC	6532	6593	6614	5112	5384	6595	6642	
SMMAL	A 2990	3270	3454	3124	3164	3316	3195	
CWMH	201	71	73	465	339	420	239	

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Effects of System Size

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• The propensity functions

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Corresponding state change matrix is

$$\mathbf{S} = \left(\begin{array}{rrrr} -1 & -2 & 2 & 0 \\ 0 & 1 & -1 & -1 \\ 0 & 0 & 0 & 1 \end{array} \right)$$

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• Assume initial conditions known $S_1(t_0) = 5\Omega$, $S_2(t_0) = S_3(t_0) = 0$, $t_0 = 0$. Reaction rate parameters to $c_1 = 1$, $\hat{c_2} = 2\Omega^{-1}$, $c_3 = 0.5$ and $c_4 = 0.04$

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Ω	M.H.	SMMALA	RMHMC
1	121 (3.6)	150 (3.9)	245 (0.06)
2	226 (6.7)	2163 (57.2)	4775 (1.3)
5	132 (3.9)	3539 (93.6)	4618 (1.2)
10	180 (5.3)	3397 (89.8)	5954 (1.6)
100	214 (6.4)	3725 (98.5)	6066 (1.7)

min. ESS vs. Ω

Posterior mean and SD. vs. Ω

Ω	C 1	Ĉ ₂	C 3	<i>C</i> ₄
1	0.88 (0.031)	1.72 (0.253)	0.39 (0.039)	0.003 (0.002)
2	1.3 (0.041)	0.69 (0.066)	0.35 (0.016)	0.014 (0.002)
5	0.93 (0.019)	0.39 (0.028)	0.48 (0.025)	0.034 (0.002)
10	1.0 (0.015)	0.18 (0.008)	0.47 (0.015)	0.037 (0.001)
100	0.99 (0.004)	0.01 (0.0002)	0.52 (0.004)	0.039 (0.0003)



Failure Modes



Figure: Simulated time point data using SSA for the Schlögl reaction set and LNA predictions. Dots correspond to simulated data. The bold and dashed red lines correspond to the LNA prediction for the means and standard deviations using the true parameters. Doted blue lines correspond the LNA predictions using the posterior means for the rate parameters. (Online version in colour.)



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