

More of the SAME?

Sequential and Pseudomarginal Monte Carlo for
Point Estimation in Latent Variable Models

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- Background:
 - Marginal MLEs
 - SAME: An MCMC Scheme
- Sequential Monte Carlo
 - The SMC Method
 - A Population-Based SAME Method
 - Examples
- Pseudomarginal Methods
 - The Pseudomarginal Method
 - More of the SAME: multiple extensions of the space
 - Example
 - Even more of the SAME: complex extensions of the space
 - Examples

Background

- Marginal MLEs
- SAME: An MCMC Scheme

- Consider a model with:
 - parameters, θ ,
 - latent variables, x , and
 - observed data, y .
- Aim to maximise marginal likelihood

$$p(y|\theta) = \int p(x, y|\theta) dx$$

or posterior

$$p(\theta|y) \propto \int p(x, y|\theta)p(\theta) dx.$$

- Traditional approach is Expectation-Maximisation (EM)
 - Requires objective function in closed form.
 - Susceptible to trapping in local optima.

- Optimization and probability \rightsquigarrow simulated annealing.
- A distribution of the form

$$\pi(\theta|y) \propto p(\theta)p(y|\theta)^\gamma$$

will become concentrated, as $\gamma \rightarrow \infty$ on the maximisers of $p(y|\theta)$ under weak conditions.

- Why not target $\pi(\theta|y)$ using MCMC?

Adapted from (Hwang, 1980; Theorem 2.1).

Assume:

- $p(\theta)$ and $p(y|\theta)$ are α -Lipschitz continuous in θ
- $\log(p(\theta)) \in \mathcal{C}^3(\mathbb{R}^n)$ and $\log p(y|\theta) \in \mathcal{C}^3(\mathbb{R}^n)$.
- Θ_{ML} is a non-empty, countable set which is nowhere dense;
- $p(\theta) \leq M < \infty$; $p(\theta) > 0 \forall \theta \in \Theta_{ML}$
- $p(y|\theta) \leq M' < \infty$
- For some $k < \sup p(y|\theta)$, $\{\theta : p(y|\theta) \geq k\}$ is compact.

Then:

$$\lim_{\gamma \rightarrow \infty} \pi_\gamma(dt) \propto \sum_{\theta_{ml} \in \Theta_{ML}} \alpha(\theta_{ml}) \delta_{\theta_{ml}}(dt), \quad (1)$$

$$\alpha(\theta_{ml}) = \det \left[- \frac{\partial^2 \log p(y|\theta)}{\partial \theta_m \partial \theta_n} \Big|_{\theta = \theta_{ml}} \right]^{-1/2} \quad (2)$$

Data Augmentation: Synthetic distributions of the form:

$$\bar{\pi}_\gamma(\theta, x_{1:\gamma}|y) \propto p(\theta) \prod_{i=1}^{\gamma} p(x_i, y|\theta)$$

admit the marginals

$$\bar{\pi}_\gamma(\theta|y) \propto p(\theta)p(y|\theta)^\gamma.$$

SAME Algorithm (Doucet, Godsill and Robert, 2002):

- $t = 0$: Initialise $(\theta_0, X_{0,1})$ arbitrarily.
- For $t = 1, \dots, T$:
 - If $\gamma(t) > \gamma(t-1)$: Set $(X_{t-1, \gamma(t-1)+1}, \dots, X_{t-1, \gamma(t)})$ arbitrarily.
 - Sample $(\theta_t, X_{t,1}, \dots, X_{t, \gamma(t)}) \sim K_{\gamma(t)}(\theta_{t-1,1}, X_{t-1,1}, \dots, X_{t-1, \gamma(t)}, \cdot)$.

Where K_γ is $\bar{\pi}_\gamma$ -invariant.

NB An inhomogeneous Markov chain.

Sequential Monte Carlo

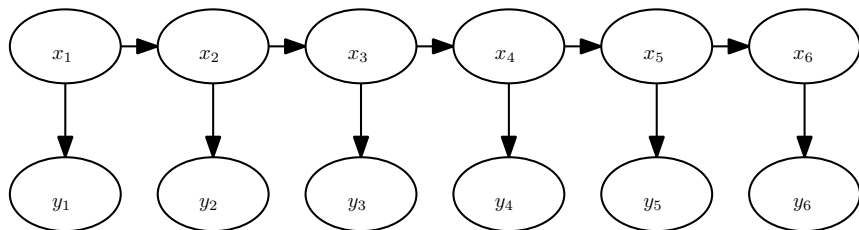
- The SMC Method
- A Population-Based SAME Method
- Examples

SMC: A Motivating Example — Filtering

- Let X_1, \dots denote the position of an object which follows Markovian dynamics.
- Let Y_1, \dots denote a collection of observations:

$$Y_i | \{X_i = x_i\} \sim g(\cdot | x_i).$$

- We wish to estimate, as observations arrive, $p(x_{1:t} | y_{1:t})$.
- A recursion obtained from Bayes rule exists but is intractable in most cases.



- Really tracking a sequence of distributions, $p_t \dots$
- on increasing state spaces.
- Other problems with the same structure exist.
- Any problem of sequentially approximating a sequence of such distributions, p_t , can be addressed in the same way.

Sequential Importance Resampling

At time t , $t \geq 2$. (Given $\{X_{1:t-1}^{(i)}\}_{i=1}^N$ approximating $p_{t-1}(x_{1:t-1})$).

Sampling Step

For $i = 1 : N$:

$$\text{sample } X_t^{(i)} \sim q_t(\cdot | X_{1:t-1}^{(i)}).$$

Resampling Step

For $i = 1 : N$:

$$\text{compute } w_t(X_{1:t}^{(i)}) = \frac{p_t(X_{1:t}^{(i)})}{p_{t-1}(X_{1:t-1}^{(i)})q_t(X_t^{(i)} | X_{1:t-1}^{(i)})}$$

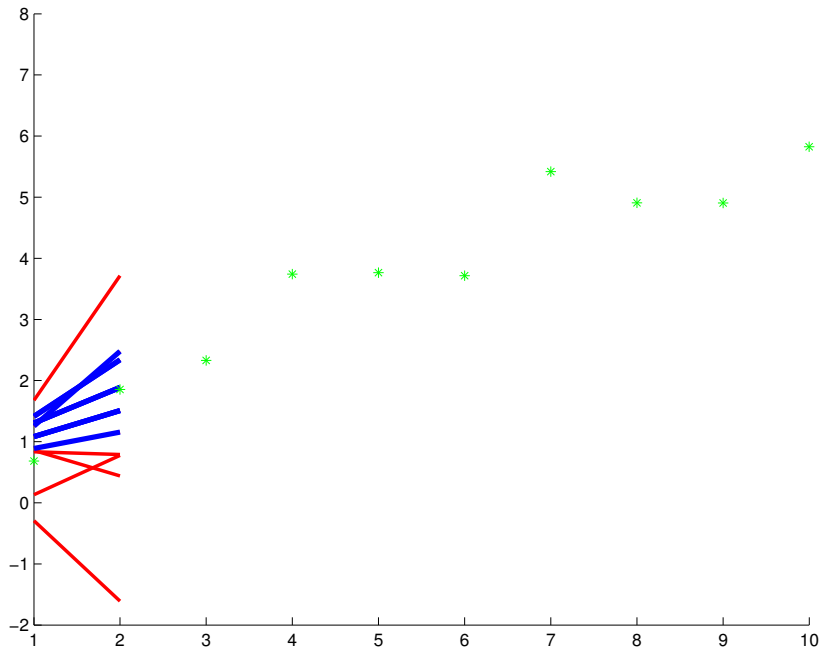
$$\text{and } W_t^{(i)} = \frac{w_t(X_{1:t}^{(i)})}{\sum_{j=1}^N w_t(X_{1:t}^{(j)})}$$

For $i = 1 : N$:

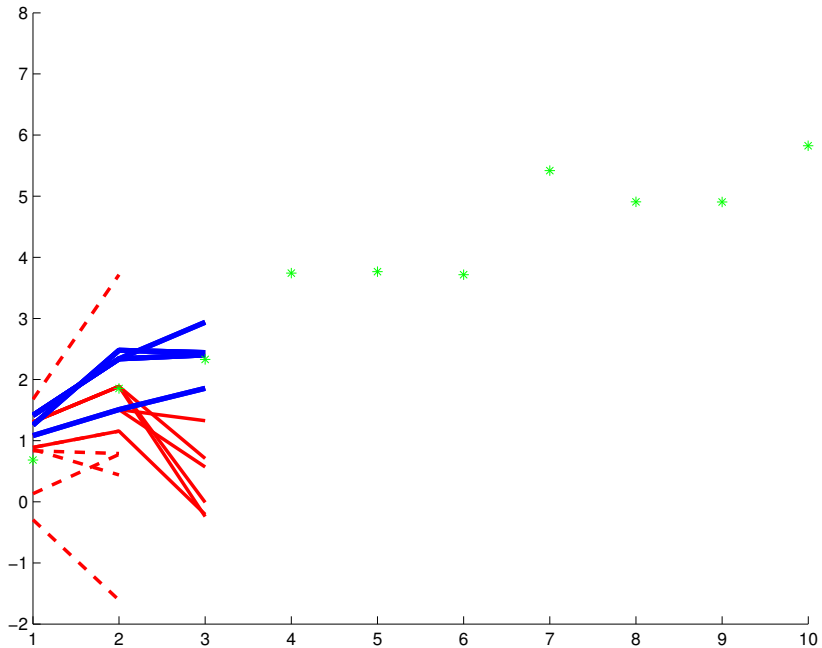
$$\text{sample } A_t^{(i)} \sim \sum_{j=1}^N W_t^{(j)} \delta_j$$

retain $\{X_{1:t}^{(A_t^i)}\}_{i=1}^N$

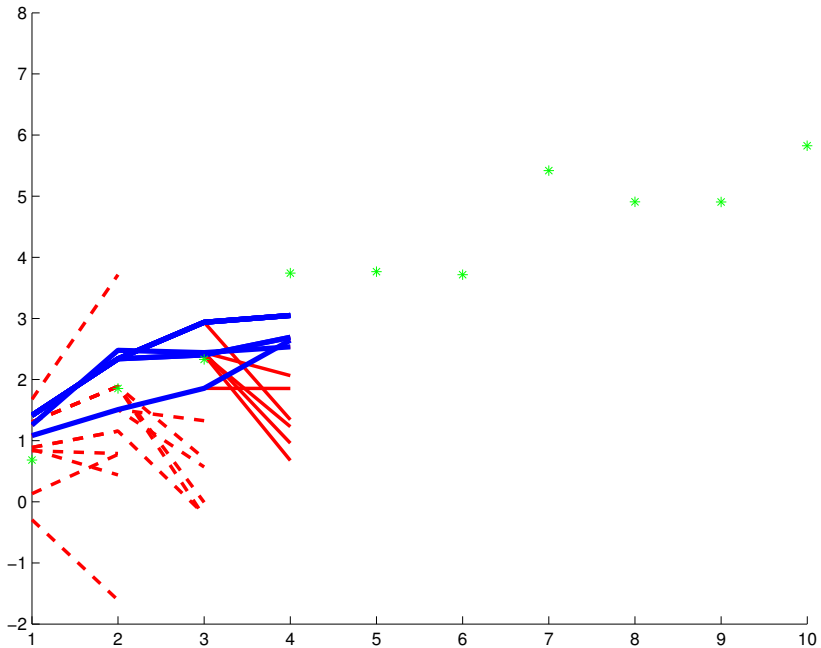
Iteration 2



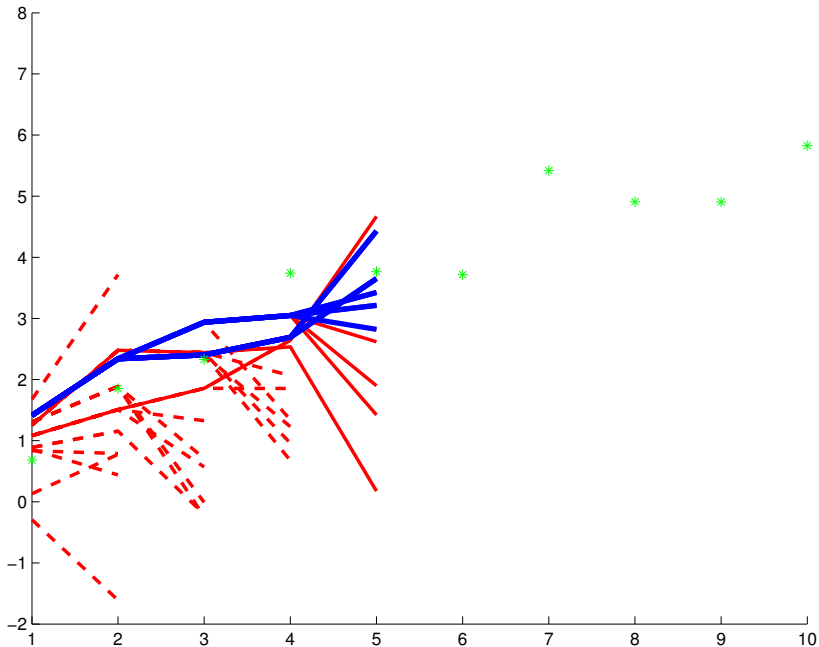
Iteration 3



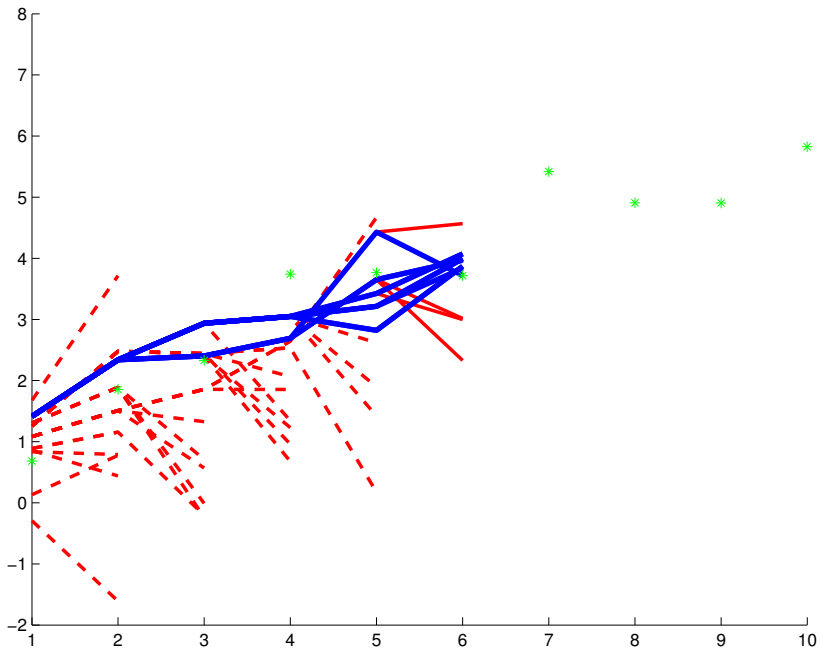
Iteration 4



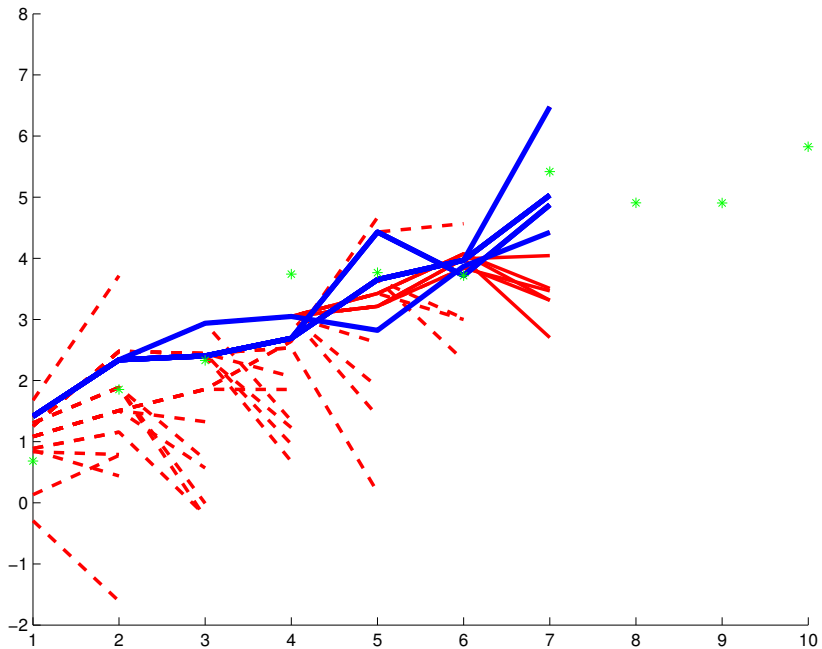
Iteration 5



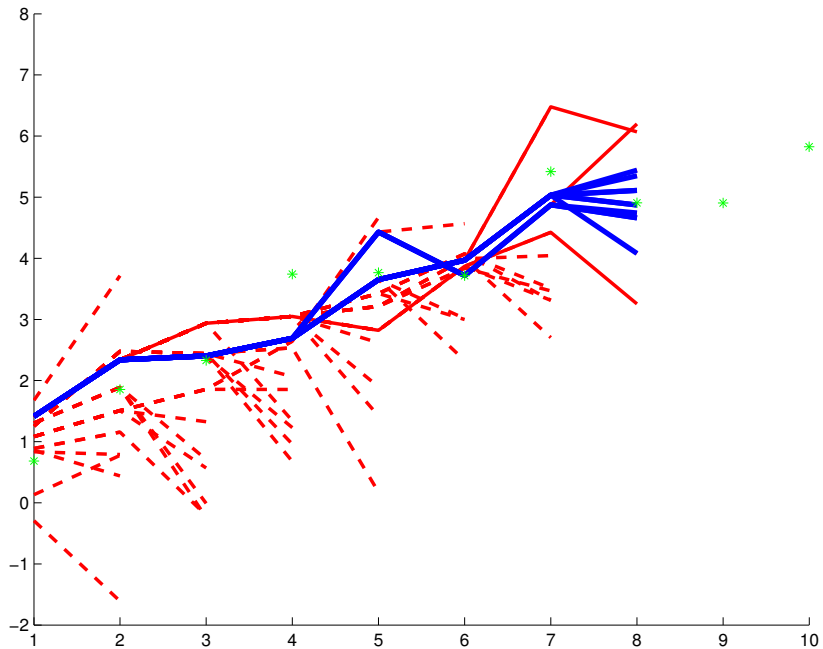
Iteration 6



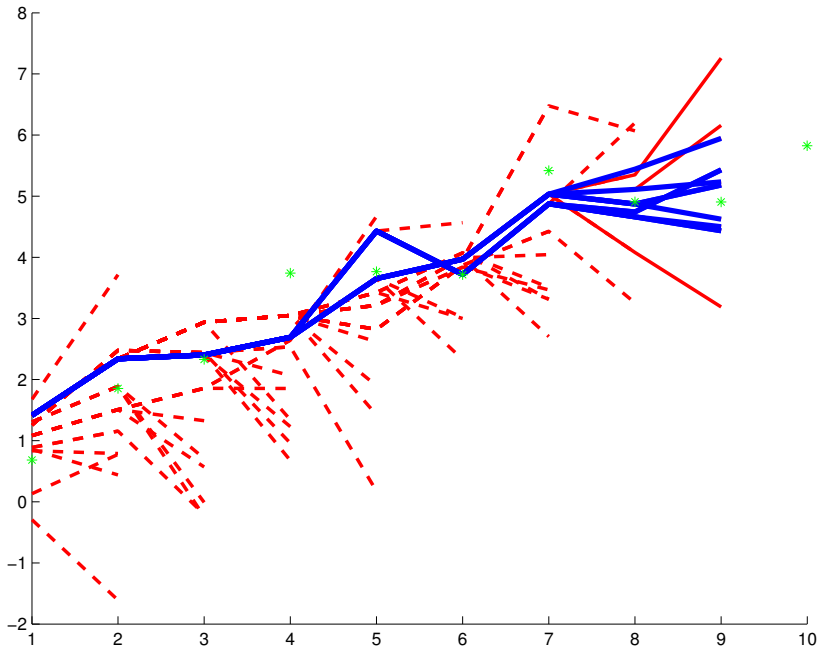
Iteration 7



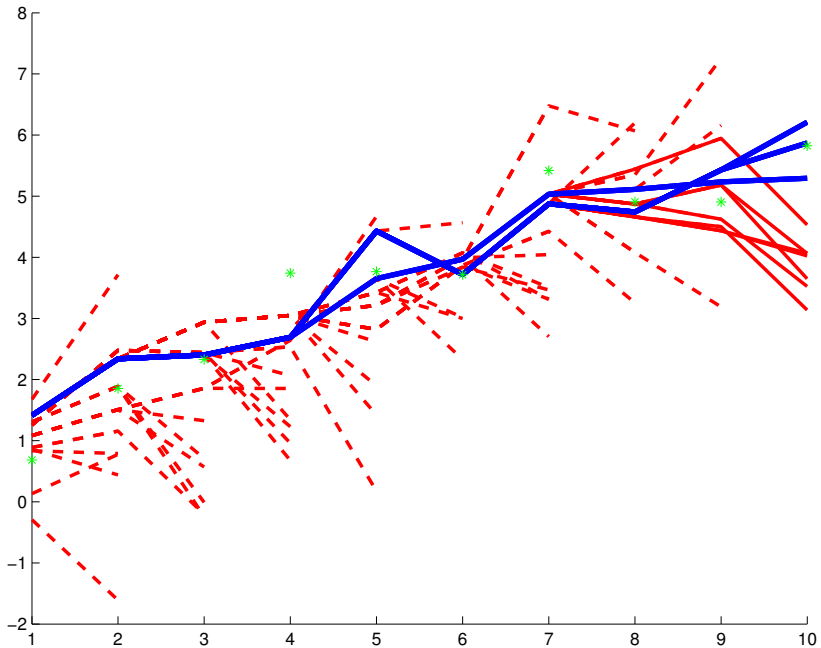
Iteration 8



Iteration 9



Iteration 10



Can be used to sample from *any* sequence of distributions:

- Given a sequence of *target* distributions, η_n , on $E_n \dots$,
- construct a synthetic sequence $\tilde{\eta}_n$ on spaces $\bigotimes_{p=1}^n E_p$
- by introducing Markov kernels, L_p from E_{p+1} to E_p :

$$\tilde{\eta}_n(x_{1:n}) = \eta_n(x_n) \prod_{p=1}^{n-1} L_p(x_{p+1}, x_p),$$

- These distributions
 - have the target distributions as final time marginals,
 - have the correct structure to employ SMC techniques.

- Given a sample $\{X_{1:n-1}^{(i)}\}_{i=1}^N$ targeting $\tilde{\eta}_{n-1}$,
- sample $X_n^{(i)} \sim K_n(X_{n-1}^{(i)}, \cdot)$,
- calculate

$$W_n(X_{1:n}^{(i)}) = \frac{\eta_n(X_n^{(i)})L_{n-1}(X_n^{(i)}, X_{n-1}^{(i)})}{\eta_{n-1}(X_{n-1}^{(i)})K_n(X_{n-1}^{(i)}, X_n^{(i)})}.$$

- Resample, yielding: $\{X_{1:n}^{(i)}\}_{i=1}^N$ targeting $\tilde{\eta}_n$.
- Hints that we'd like to use

$$L_{n-1}(x_n, x_{n-1}) = \frac{\eta_{n-1}(x_{n-1})K_n(x_{n-1}, x_n)}{\int \eta_{n-1}(x'_{n-1})K_n(x'_{n-1}, x_n)dx_{n-1}}.$$

- A model with:
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 - observed data, y .
- Aim to maximise Marginal likelihood

$$p(y|\theta) = \int p(x, y|\theta) dx$$

or posterior

$$p(\theta|y) \propto \int p(x, y|\theta)p(\theta) dx$$

- Using

$$\bar{\pi}_\gamma(\theta, x_{1:\gamma}|y) \propto p(\theta) \prod_{i=1}^{\gamma} p(x_i, y|\theta)$$

- Use a sequence of distributions $\eta_n = \bar{\pi}_{\gamma_n}$ for some $\{\gamma_n\}$.
- The MCMC approach (Doucet et al., 2002).
 - Requires slow “annealing”.
 - Separation between distributions is large.
 - Mixes poorly as γ increases.
- Using SMC has some substantial advantages:
 - Introducing bridging distributions, for $\gamma = \lfloor \gamma \rfloor + \langle \gamma \rangle$, of:

$$\bar{\pi}_{\gamma}(\theta, x_{1:\lfloor \gamma \rfloor+1} | y) \propto p(\theta) p(x_{\lfloor \gamma \rfloor+1}, y | \theta)^{\langle \gamma \rangle} \prod_{i=1}^{\lfloor \gamma \rfloor} p(x_i, y | \theta)$$

is straightforward.

- Population of samples improves robustness.
- It is less dependent upon mixing of K_{γ} .

- A generic SMC sampler can be written down directly...
- An easy case:
 - Sample from $p(x_t|y, \theta_{t-1})$ and $p(\theta_t|x_t, y)$.
 - Weight according to $p(y|\theta_{t-1})^{\gamma_t - \gamma_{t-1}}$.
- The general case:
 - Sample existing variables from a π_t -invariant kernel:

$$(\theta_t, X_{t,1:\gamma_{t-1}}) \sim K_t((\theta_{t-1}, X_{t-1}), \cdot).$$

- Sample new variables from an arbitrary proposal:

$$X_{t, [\gamma_{t-1}] + 1 : [\gamma_t]} \sim q(\cdot | \theta_t).$$

- Use combination of time-reversal and optimal auxiliary kernel.
 - Weight expression does not involve the marginal likelihood.

initialisation: $t = 1$:

sample $\left\{ \left(\theta_1^{(i)}, X_1^{(i)} \right) \stackrel{\text{iid}}{\sim} \nu \right\}_{i=1}^N$

calculate $W_1^{(i)} \propto \frac{\pi_{\gamma_1}(\theta_1^{(i)}, X_1^{(i)})}{\nu(\theta_1^{(i)}, X_1^{(i)})} \quad \left(\sum_{i=1}^N W_1^{(i)} = 1 \right)$

for $t = 2$ to T **do**

resample

sample $\left\{ \begin{array}{l} \left(\theta_t^{(i)}, X_{t,1:\lceil \gamma_{t-1} \rceil} \right) \sim K_{t-1} \left(\theta_{t-1}^{(i)}, X_{t-1}^{(i)}; \cdot \right) \\ \left\{ X_{t,j}^{(i)} \sim q(\cdot | \theta_t^{(i)}) \right\}_{j=\lceil \gamma_{t-1} \rceil+1}^{\lceil \gamma_t \rceil} \quad \text{if } \lceil \gamma_{t-1} \rceil < \lceil \gamma_t \rceil \\ X_{t,\lceil \gamma_t \rceil}^{(i)} \sim q_{\langle \gamma_t \rangle}(\cdot | \theta_t^{(i)}) \quad \text{if } \lceil \gamma_{t-1} \rceil < \lceil \gamma_t \rceil \neq \gamma_t \end{array} \right\}_{i=1}^N$

calculate

$$W_t^{(i)} \propto \frac{p(y, X_{t,\lceil \gamma_{t-1} \rceil} | \theta)^{1 \wedge \gamma_t - \lceil \gamma_{t-1} \rceil}}{p(y, X_{t,\lceil \gamma_{t-1} \rceil} | \theta)^{\langle \gamma_t \rangle}} \prod_{j=\lceil \gamma_{t-1} \rceil+1}^{\lceil \gamma_t \rceil} \frac{p(y, X_{t,j} | \theta_t)}{q(X_{t,j} | \theta_t)} \left(\frac{p(y, X_{t,\lceil \gamma_t \rceil} | \theta_t)^{\langle \gamma_t \rangle}}{q_{\langle \gamma_t \rangle}(X_{t,\lceil \gamma_t \rceil} | \theta_t)} \right)^I$$

with $I = \mathbb{I}(\lceil \gamma_t \rceil > \lceil \gamma_t \rceil \geq \lceil \gamma_{t-1} \rceil)$.

end for

Toy Example (using known marginal likelihood)

- Student t -distribution of unknown location parameter θ with $\nu = 0.05$.
- Four observations are available, $y = (-20, 1, 2, 3)$.
- Log likelihood is:

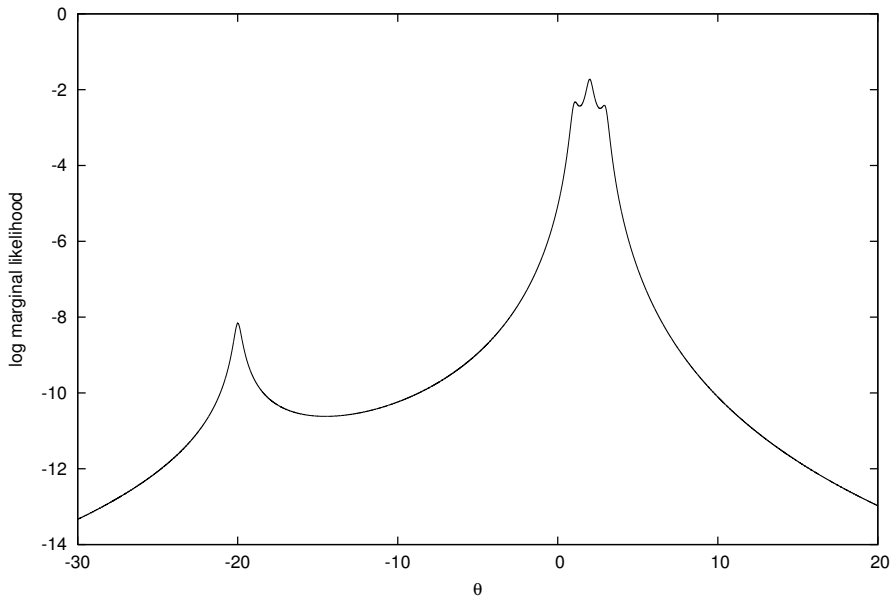
$$\log p(y|\theta) = -0.525 \sum_{i=1}^4 \log (0.05 + (y_i - \theta)^2).$$

- Global maximum is at 1.997.
- Local maxima at $\{-19.993, 1.086, 2.906\}$.
- Complete log likelihood ($X_i \sim \mathcal{G}a$):

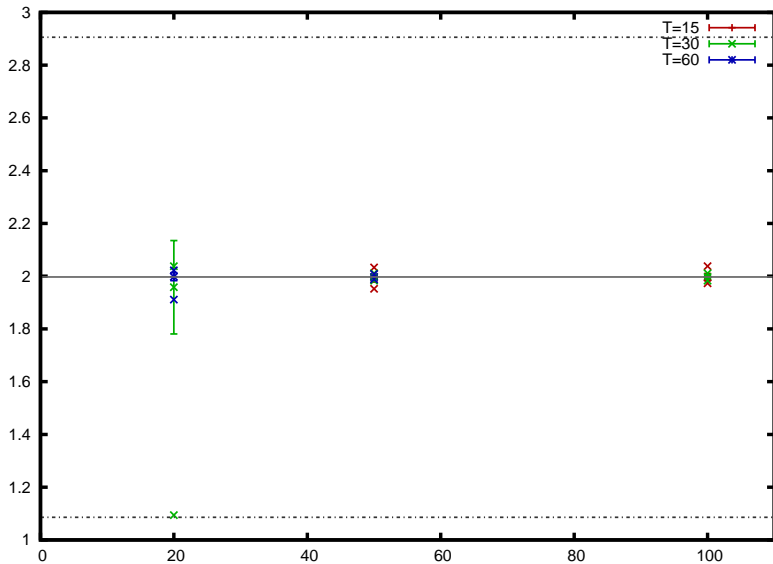
$$\log p(y, z|\theta) = - \sum_{i=1}^4 [0.475 \log x_i + 0.025x_i + 0.5x_i(y_i - \theta)^2].$$

Toy Example: Marginal Likelihood

Toy Example: Log Marginal Likelihood



Toy Example: SMC Method using Gibbs Kernels



- Likelihood $p(y|x, \omega, \mu, \sigma) = \mathcal{N}(y|\mu_x, \sigma_x^2)$.
- Marginal likelihood $p(y|\omega, \mu, \sigma) = \sum_{j=1}^3 \omega_j \mathcal{N}(y|\mu_j, \sigma_j^2)$.
- Diffuse conjugate priors were employed:

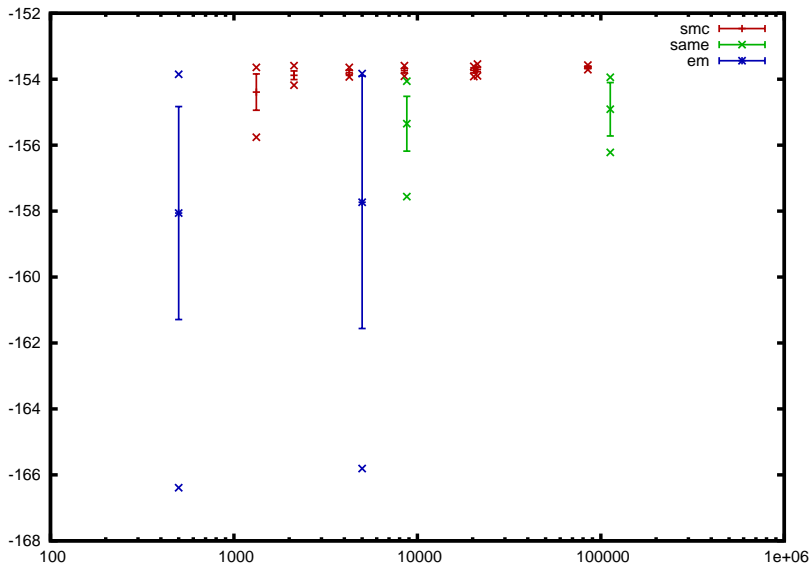
$$\omega \sim \mathcal{D}i(\delta)$$

$$\sigma_i^2 \sim \mathcal{IG}\left(\frac{\lambda_i + 3}{2}, \frac{\beta_i}{2}\right)$$

$$\mu_i | \sigma_i^2 \sim \mathcal{N}(\alpha_i, \sigma_i^2 / \lambda_i),$$

- **All full conditional distributions of interest are available.**
- **Marginal posterior can be calculated.**

3 Component GMM (Roeders Galaxy Data Set)



Pseudomarginal Monte Carlo

- The Pseudomarginal Method
- More of the SAME: multiple extensions of the space
- Example
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The Pseudomarginal Method

- Marginal MH-Acceptance Probability:

$$1 \wedge \frac{\pi(\theta')Q(\theta', \theta)}{\pi(\theta)Q(\theta, \theta')}$$

- But $\pi(\theta)$ isn't tractable: how about using:

$$1 \wedge \frac{\hat{\pi}(\theta')Q(\theta', \theta)}{\hat{\pi}(\theta)Q(\theta, \theta')}$$

where

$$\hat{\pi}(\theta) = \frac{1}{m} \sum_{i=1}^m \frac{\pi(\theta, X_i)}{q(X_i)} \quad X_i \stackrel{\text{iid}}{\sim} q$$

- Suggests two algorithms (Beaumont, 2003):
 - Monte Carlo within Metropolis
 - Grouped Independence Metropolis Hastings

- Extended Target (Andrieu & Roberts, 2009):

$$\begin{aligned}\tilde{\pi}(\theta, x_1, \dots, x_m) &= \sum_{j=1}^m \frac{1}{m} \pi(\theta, x_j) \prod_{k \neq j} q(x_k) \\ &= \frac{1}{m} \sum_{j=1}^m \frac{\pi(\theta, x_j)}{q(x_j)} \cdot \prod_{k=1}^m q(x_k) = \hat{\pi}(\theta) \prod_{k=1}^m q(x_k)\end{aligned}$$

- The acceptance probability becomes:

$$1 \wedge \frac{\tilde{\pi}(\theta', x'_1, \dots, x'_m) Q(\theta', \theta) \prod_{j=1}^m q(x_j)}{\tilde{\pi}(\theta, x_1, \dots, x_m) Q(\theta, \theta') \prod_{j=1}^m q(x'_j)} = 1 \wedge \frac{\hat{\pi}(\theta') Q(\theta', \theta)}{\hat{\pi}(\theta) Q(\theta, \theta')}$$

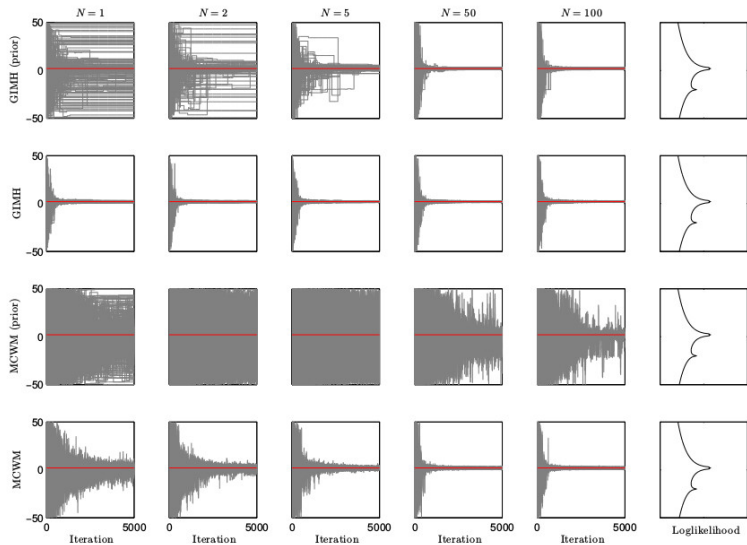
- NB MCWM is *not* exact... but perhaps we don't care.

- We'd like to target $\pi_\gamma(\theta|y) \propto p(\theta)p(y|\theta)^\gamma$.
- Why not use the pseudomarginal approach, considering instead:

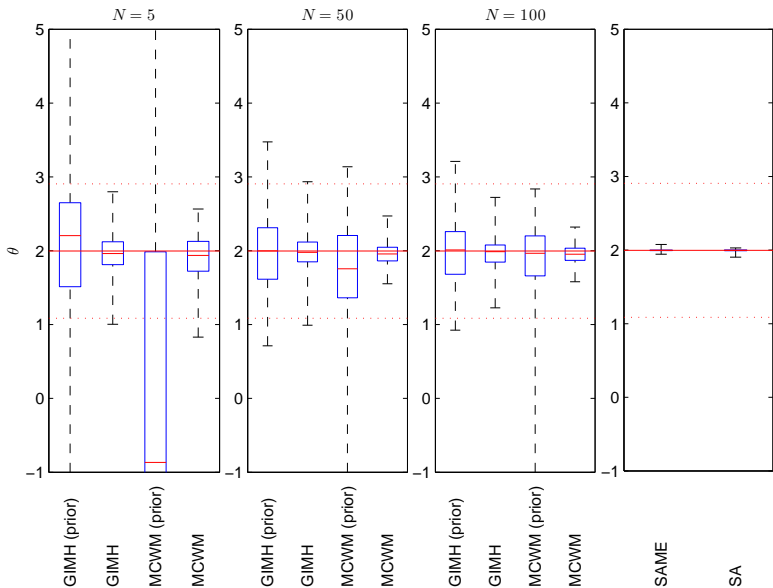
$$\tilde{\pi}_\gamma(\pi, x_{1:m}^1, \dots, x_{1:m}^\gamma) = p(\theta) \prod_{i=1}^{\gamma} \sum_{j=1}^m \frac{1}{m} \frac{p(x_j^i, y|\theta)}{q(x_j^i|\theta)} \prod_{k=1}^m q(x_k^i|\theta)$$

- Expect behaviour like simulated annealing for large m .

The Student t Model Revisited



Summary of 200 Runs



What about complicated latent variable structures?

- Actually, pseudomarginal algorithms are more flexible.
- We're especially interested in particle MCMC implementations (Andrieu et al., 2010):
 - Particle Marginal Metropolis-Hastings(PMMH)
 - MCWM variant of PMMH
 - Particle Gibbs (with ancestor sampling)
- State-space models are the real motivation for this methodology.
- Many other complex models could be addressed using this technique.

- Model:

$$X_t = AX_{t-1} + BU_t$$

$$Y_t = X_t + DV_t$$

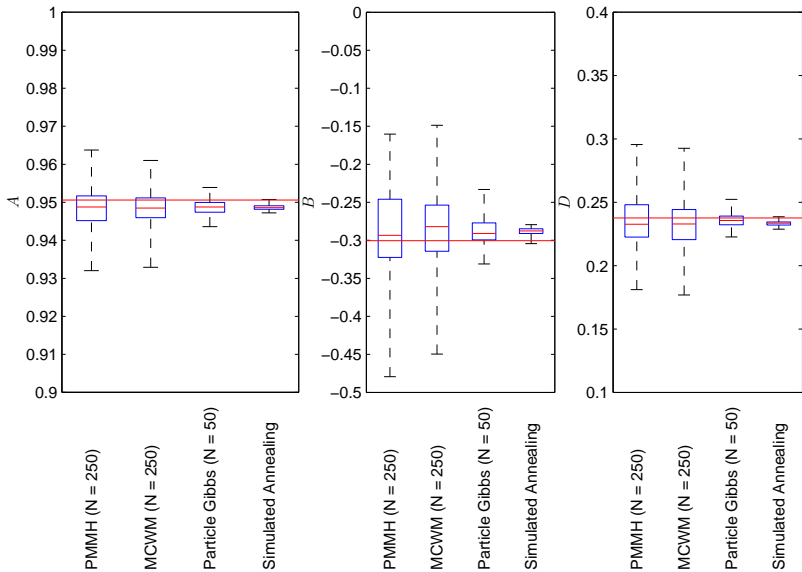
- Data 50 observations simulated using:

$$A = 0.9, B = 1, \text{ and } D = 1$$

- Algorithms

- The PMMH/MCWM algorithms use $N = 250$ particles;
- The PG algorithm (with ancestor sampling) uses $N = 50$ particles but attempts 100 static parameter updates per iteration.
- Inverse temperature increases linearly from 0.1 to 10.
- Final 1000 iterations $\gamma_t = 10$.
- Compare with exact marginal simulated annealing algorithm.

Summary of 100 Runs



A Simple Stochastic Volatility Model

- Model:

$$X_i = \alpha + \delta X_{i-1} + \sigma_u u_i \quad X_1 \sim \mathcal{N}(\mu_0, \sigma_0^2)$$

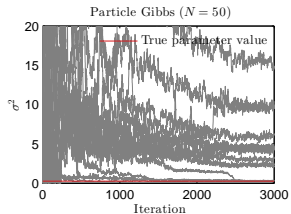
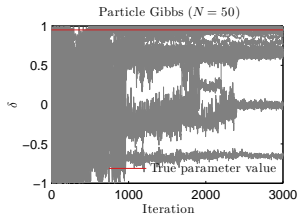
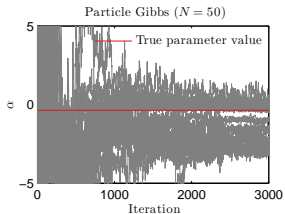
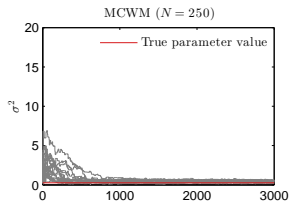
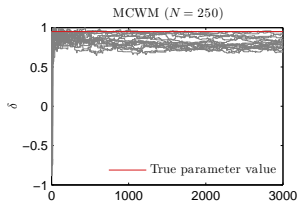
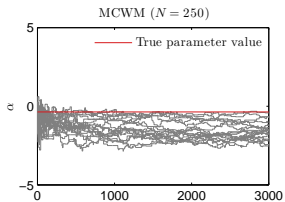
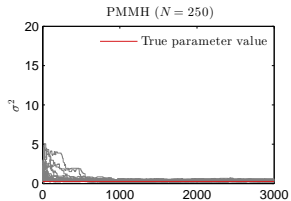
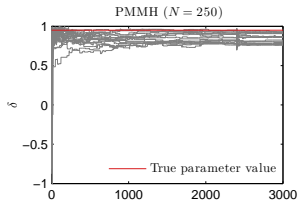
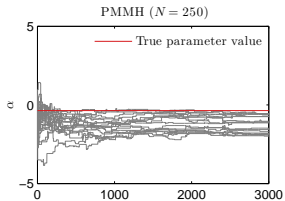
$$Y_i = \exp\left(\frac{X_i}{2}\right) \epsilon_i$$

where u_i and ϵ_i are uncorrelated standard normal random variables, and $\theta = (\alpha, \delta, \sigma_u)$.

- 200 Observations; simulated with $\delta = 0.95$, $\alpha = -0.363$ and $\sigma = 0.26$.
- Diffuse instrumental prior distributions:
 - $\delta \sim U(-1, 1)$
 - $\alpha \sim \mathcal{N}(0, 1)$
 - $\sigma^{-2} \sim \mathcal{Ga}(1, 0.1)$

are quickly forgotten.

- Inverse temperature increases linearly from 0.1 to 10.
- Final 500 iterations $\gamma_t = 10$.
- A more complex multi-factor model is also under investigation.



- Monte Carlo isn't just for calculating posterior expectations.
- SMC and Pseudomarginal methods are effective for ML and MAP estimation.
- Still work in progress. . .
- Scope for embedding Pseudomarginal target within SMC algorithm. . .
- and adaptation.

- [1] C. Andrieu and G. O. Roberts. The pseudo-marginal approach for efficient Monte Carlo computations. *Annals of Statistics*, 37(2):697–725, 2009.
- [2] C. Andrieu, A. Doucet, and R. Holenstein. Particle Markov chain Monte Carlo. *Journal of the Royal Statistical Society B*, 72(3):269–342, 2010.
- [3] M. Beaumont. Estimation of population growth or decline in genetically monitored populations. *Genetics*, 164(3):1139–1160, 2003.
- [4] P. Del Moral, A. Doucet, and A. Jasra. Sequential Monte Carlo samplers. *Journal of the Royal Statistical Society B*, 63(3):411–436, 2006.
- [5] A. Doucet, S. J. Godsill, and C. P. Robert. Marginal maximum a posteriori estimation using Markov chain Monte Carlo. *Statistics and Computing*, 12:77–84, 2002.
- [6] **A. Finke. *On Extended State-Space Constructions for Monte Carlo Methods*. Ph.D. thesis, University of Warwick, 2015. In preparation.**
- [7] **A. Finke and A. M. Johansen. *More of the SAME? Pseudomarginal methods for point estimation in latent variable models*. In preparation, 2015.**
- [8] C.-R. Hwang. Laplace’s method revisited: Weak convergence of probability measures. *Annals of Probability*, 8(6):1177–1182, December 1980.
- [9] A. M. Johansen, A. Doucet, and M. Davy. Maximum likelihood parameter estimation for latent models using sequential Monte Carlo. In *Proceedings of ICASSP*, volume III, pages 640–643. IEEE, May 2006.
- [10] **A. M. Johansen, A. Doucet, and M. Davy. Particle methods for maximum likelihood parameter estimation in latent variable models. *Statistics and Computing*, 18(1):47–57, March 2008.**