An Adaptive Sequential Monte Carlo Approach for Bayesian Model Comparison

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Outline

Bayesian model comparison

Sequential Monte Carlo Approach

Adaptive strategies

Some performance examples

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Summary

Bayesian model comparison

Basic formulas

Given a collection of (at most countable) models $\{M_k\}_{k \in \mathcal{K}}$,

$$\pi(M_k|\boldsymbol{y}) = \frac{\pi(M_k)p(\boldsymbol{y}|M_k)}{p(\boldsymbol{y})}$$
$$p(\boldsymbol{y}|M_k) = \int_{\Theta_k} \pi(\theta_k|M_k)p(\boldsymbol{y}|\theta_k, M_k) \, \mathrm{d}\theta_k$$

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Common approaches

Evaluate posterior model probabilities $\pi(M_k|\mathbf{y})$ directly Evaluate marginal likelihood $p(\mathbf{y}|M_k)$ individually Evaluate Bayes factors $\frac{p(\mathbf{y}|M_{k+1})}{p(\mathbf{y}|M_k)}$ sequentially

What do we want? Better estimates at less computational cost with less manual calibration

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Unbiased or almost unbiased Consistent

Smaller variance or smaller MSE if biased

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Less manual calibration

Generic approach Adaptive strategies

Initialization from $\eta_0(X)$ for $\pi_0(X)$

Iterate over intermediate distributions $\{\pi_t(X) = \gamma_t(X)/Z_t\}_{t=1}^T$

Terminiate at $\pi_T(X)$ and estimation

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Draw $\{X_0^{(i)}\}_{i=1}^N$ from η_0 Compute $\{W_0^{(i)}\}_{i=1}^N$

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Draw $\{X_0^{(i)}\}_{i=1}^N$ from η_0 Compute $\{W_0^{(i)}\}_{i=1}^N$

Resampling if necessary Draw $X_t^{(i)}$ from $K_t(X_{t-1}^{(i)}, \cdot)$ for i = 1, ..., NCompute incremental weights $\{\tilde{w}_t^{(i)}(X_{t-1}^{(i)}, X_t^{(i)})\}_{i=1}^N$ Compute normalized weights $\{W_t^{(i)}\}_{i=1}^N$

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 $\begin{aligned} \pi_t^N(\mathrm{d}\,x) &= \sum_{i=1}^N W_t^{(i)} \delta_{X_t^{(i)}}(\mathrm{d}\,x) \text{ approxiamte } \pi_t(\mathrm{d}\,x) \\ \frac{\hat{Z}_t}{Z_{t-1}} &= \sum_{i=1}^N W_{t-1}^{(i)} \tilde{w}_t^{(i)} \text{ estimates the ratio of normalizing constants recusively} \end{aligned}$

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Terminiate at $\pi_T(X)$ and estimation

$$\eta_0(\theta_0, M_0) = \pi_0(\theta_0, M_0) \propto \pi(M_0)\pi(\theta_0|M_0)$$

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 $\pi_t(\theta_t, M_t) \propto \pi(M_t)\pi(\theta_t|M_t)p(y|\theta_t, M_t)^{\alpha(t/T)}$ Markov kernel $K_t(X_{t-1}, \cdot)$ requires both within- and inter-model moves

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Estimate $\pi(M_k|\mathbf{y})$ using particle approximation to $\pi_T(\theta_T, M_T)$

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 $\begin{aligned} \pi_t(\theta_t) &\propto \pi(\theta_t | M_k) p(\boldsymbol{y} | \theta_t, M_k)^{\alpha(t/T)} \text{ or } \\ \pi_t(\theta_t) &\propto \pi(\theta_t | M_k) p(\boldsymbol{y}_{1:t} | \theta_t, M_k) \text{ (Chopin, 2002)} \\ \text{Markov kernel } K_t(X_{t-1}, \cdot) \text{ only within-model moves} \end{aligned}$

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Estimate $p(\mathbf{y}|M_k)$, the normalizing constant ratio Z_T/Z_0

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 $\pi_0(\theta_0) \propto \pi(\theta_0|M_k) p(y|\theta_0,M_k)$ $\eta_0(\theta_0): \text{ The particle system of the sampler iterating from model } M_{k-1} \text{ to } M_k.$

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$$\begin{split} \pi_t(\theta_t,M_t) \propto \pi_t(M_t) \pi(\theta_t|M_t) p(\pmb{y}|\theta_t,M_t) \\ \pi_t(M_t) &= \alpha(t/T) \end{split}$$
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Estimate Bayes factor $B_{k+1,k}$, the normalizing constant ratio Z_T/Z_0

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Sequential Monte Carlo – Path sampling estimator

Basic identity

Given a family of distributions $\{\pi_{\alpha}(x) = q_{\alpha}(x)/Z_{\alpha}\}_{\alpha \in [0,1]}$

$$\log\left(\frac{Z_1}{Z_0}\right) = \int_0^1 \mathbb{E}_{\alpha}\left[\frac{d\log q_{\alpha}(X)}{d\alpha}\right] \, \mathrm{d}\,\alpha$$

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Using SMC samples

Particle approximations of the expectations from SMC samplers Numerical integration to approximate the estimator Sequential Monte Carlo – Path sampling estimator

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Andrew Gelman and Xiao-Li Meng (1998). "Simulating normalizing constants: From importance sampling to bridge sampling to path sampling". In: *Statistical Science* 13.2, pp. 163–185

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Simple illustrative example: Gaussian mixture model

Model

Determine the number of components k, which define the model by

$$\begin{aligned} y_i | \theta_k &\sim \sum_{j=1}^k \omega_j \mathcal{N}(\mu_j, \lambda_j^{-1}) \qquad i = 1, \dots, n \\ \mu_j &\sim \mathcal{N}(\xi, \kappa^{-1}) \quad \lambda_j &\sim \mathcal{G}(\nu, \chi) \quad \omega_{1:k} \sim \mathcal{D}(\rho) \end{aligned}$$

Pierre Del Moral, Arnaud Doucet, and Ajay Jasra (2006). "Sequential Monte Carlo samplers". In: *Journal of Royal Statistical Society B* 68.3, pp. 411–436

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Simple illustrative example: Gaussian mixture model Comparison of estimates of Bayes factor $B_{5,4}$



Als & SMC: 1,000 particles, 100 time steps, $\alpha(t/T) = (t/T)^2$

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RJMCMC: 100,000 iterations

Purpose

Create a smooth sequence of distributions that reduces discrepancy between π_{t-1} and π_t

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Assumption

The criterion for adaptation can be calculated prior to the sampling of next iteration.

For example, $\tilde{w}_t(X_{t-1},X_t) \propto \pi_t(X_{t-1})/\pi_{t-1}(X_t)$ when $K_t(X_{t-1},\cdot)$ is π_t invariant

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Why does it matter?

"the variance of (\tilde{w}_t) will typically be high if the discrepancy between π_{t-1} and π_t is large *even if the kernel* K_t *mixes very well*" (Del Moral, Doucet, and Jasra, 2006)

Recall normalizing constants estimator relates directly to \tilde{w}_t

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Recall normalizing constants estimator relates directly to \tilde{w}_t

Desired effect of the adaptive strategy

Independent of resampling strategies – it is a property of the sequence of distributions

Using ESS (Jasra et al., 2008; Schäfer and Chopin, 2011)

$$\mathsf{ESS}_{t} = \frac{(\sum_{j=1}^{N} W_{t-1}^{(j)} \tilde{w}_{t}^{(j)})^{2}}{\sum_{j=1}^{N} (W_{t-1}^{(j)})^{2} (\tilde{w}_{t}^{(j)})^{2}}$$

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At time t - 1, find $\alpha(t/T)$ such that Ess_t equal a specific value

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At time t - 1, find $\alpha(t/T)$ such that Ess_t equal a specific value

Caveats

The sequence of distributions depends on the resampling strategies

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Simple illustrative example: Gaussian mixture model

Consider a SMC sampler on $\{\pi_t(\theta_t)\}_{t=0}^T$

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Problem

At each $\alpha(t/T)$, find the $\Delta \alpha(t/T) = \alpha((t+1)/T) - \alpha(t/T)$. How does $\Delta \alpha$ evolve along with α ?

Does the adaptive specification of the sequence of distributions, $\alpha(t/T)$ improve the normalizing constant estimator?

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Benchmark Comparison to $\alpha(t/T) = t/T$ Simple illustrative example: Gaussian mixture model Change of path sampling integrands ($\log p(y|\theta_t, M_k)$)



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Adaptive strategies – Specification of distributions Using ESS– Resampling every iteration (Schäfer and Chopin, 2011)



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Adaptive strategies – Specification of distributions Using ESS– Resampling only when ESS < N/2 (Jasra et al., 2008)



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A new approach: CESS- Conditional ESS

$$\mathsf{cess}_t = \frac{(\sum_{j=1}^N W_{t-1}^{(j)} \tilde{w}_t^{(j)})^2}{\sum_{j=1}^N \frac{1}{N} W_{t-1}^{(j)} (\tilde{w}_t^{(j)})^2}$$

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A new approach: cess- Conditional ess

$$\mathsf{CESS}_{t} = \frac{(\sum_{j=1}^{N} W_{t-1}^{(j)} \tilde{w}_{t}^{(j)})^{2}}{\sum_{j=1}^{N} \frac{1}{N} W_{t-1}^{(j)} (\tilde{w}_{t}^{(j)})^{2}}$$

Properties

Approximate the ESS as if resampling at time t - 1 without actually doing it Produce the same sequence regardless of resampling strategy

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Adaptive strategies – Specification of distributions Using CESS – Resampling every iteration



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Adaptive strategies – Specification of distributions Using CESS– Resampling every iteration



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Adaptive strategies – Specification of distributions Using CESS – Resampling only when ESS < N/2



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Adaptive strategies – Specification of distributions Using CESS – Resampling only when ESS < N/2



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Adaptive strategies – Calibrating RWM or MALA proposal scales

Estimating moments of parameters from particle approximations



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Performance: SMC vs AIS

Annealed importance resampling

sмс without resampling

Some argues SMC does not improve results for normalizing constant estimates

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Effects of resampling in estimating normalizing constants



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Performance: Path sampling using sмс vs Population-мсмс

Population-MCMC with path sampling estimator (Calderhead and Girolami, 2009)

Sampling parallel MCMC chains for $\pi(X_{0:T}) = \prod_{t=0}^{T} \pi_t(X_t)$, with local mixing and global swap/crossover moves

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Performance: Path sampling using sмс vs Population-мсмс

Population-MCMC with path sampling estimator (Calderhead and Girolami, 2009)

Sampling parallel MCMC chains for $\pi(X_{0:T}) = \prod_{t=0}^{T} \pi_t(X_t)$, with local mixing and global swap/crossover moves



sмс: Fix number of particles N = 1000; Population-мсмс: Fix number of iterations N = 10000

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Performance: Scalability on GPU parallelization Implemented with OpenCL on NVIDIA Quadro 2000



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Summary

Bayesian model comparison via Sequential Mote Carlo

 Can be used as drop-in replacement where conventional MCMC, RJMCMC, etc., were used

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- Requires minimal manual calibration
- Can provide better and more robust performance
- Can be parallelized straightforwardly

Thank You!

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