Using replica exchange Hamiltonian Monte Carlo and thermodynamic integration for comparison of dynamic rainfall-runoff models

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- 1. Introduction
- 2. Model development
- 3. Results

Introduction

- Hydrology is the study of the movement of water in the environment.
- Hydrologists develop different types of models to understand water dynamics.
- The Hydrologiska Byråns Vattenbalansavdelning (HBV) model and its variants are used in over 50 countries to model hydrological systems (Bergstrom, 2006).
- The HBV model links precipitation with hydrological catchment outflow. The calibrated models are used for flood prediction, water management etc.

- Today, Bayesian inference is widely used in hydrology for parameter identification (Marshall, Nott, & Sharma, 2005).
- However, Bayesian model selection criteria are not widely used, even though the model selection problem is equally important for practitioners.

- 1. Computational expense.
- 2. Poor robustness of algorithms for moderate dimensional problems.
- 3. Difficulty of implementing those algorithms.
 - Require gradients, not always available.
 - Technical or mathematical expertise.

- 1. Algorithmic
 - We introduce REHMC+TI, a combination of replica exchange (RE), Hamiltonian Monte Carlo (HMC) and thermodynamic integration (TI) for efficient and robust parameter and marginal likelihood estimation.
- 2. Statistical
 - We introduce formal posterior predictive checks for ODE-based models.
- 3. Methodological
 - Algorithms are implemented in the differentiable programming language TensorFlow Probability for flexible future use.

Data

Time series data from Magela Creek Australia



Figure 1: Plot of observed discharge and precipitation from 01-01-1980 to 31-03-1980.

- Magela Creek is a gauged catchment.
- The variable of interest is discharge (Q mmday⁻¹).
- The independent variables:
 - Precipitation (mmday⁻¹).
 - Actual evapotranspiration (mmday⁻¹).

Model development

We develop a system of ordinary differential equations to mimic the HBV model.

- P : Precipitation (L^3T^{-1})
- E_a : Actual evapotranspiration (L^3T^{-1})
- Q_1 : Discharge (L^3T^{-1})
- k_1 : outflow recession coefficient (T^{-1})
- $k_{1,2}$: inter-bucket recession coefficient (T^{-1})

$$\frac{\mathrm{d}V_1}{\mathrm{d}t} = P - E_a - k_{1,2}V_1 - Q_1$$
$$= P - E_a - k_{1,2}V_1 - k_1V_1.$$



Multi-bucket HBV model

- $V_t := \frac{\mathrm{d}V}{\mathrm{d}t}$ is the derivative of the state with respect to the time variable t.
- $\hat{V} \in \mathbb{R}^n$ are the initial conditions.

$$(V_{1})_{t} = P - E_{a} - k_{1}V_{1},$$

$$(V_{i})_{t} = k_{(i-1)(i)}V_{i-1} - k_{i}V_{i}, \ i = 2, \dots, n-1, n > 2,$$

$$(V_{n})_{t} = k_{(n-1)(n)}V_{n-1} - k_{n}V_{n},$$

$$V(t = 0) = \hat{V},$$

$$E_{a} = \frac{E_{p}}{V_{\max}}V_{1},$$

$$Q = \sum_{i=1}^{n} k_{i}V_{i}.$$



$$y = G_{\rm obs}G_{\rm sol}(\theta) + \eta,$$
$$\eta \sim \mathcal{N}(0, \sigma^2 I_p)$$

- $G_{sol}: \mathbb{R}^q \to X$ maps the parameter vector $\theta \in \mathbb{R}^q$, with q = 3n, to the total discharge $Q \in X$ through solving the ODE system.
- G_{obs} : Evaluates the total discharge at specific time points $\{t_1, \ldots, t_p\}$.
- η : noise, assumed Gaussian with covariance $\sigma^2 I_p \in \mathbb{R}^{p \times p}$ with I_p the identity matrix.

$$G := G_{\rm obs} G_{\rm sol}$$
$$y | \theta \sim \mathcal{N}(G(\theta), \sigma^2 I_p)$$



Figure 2: Schematic representation of likelihood construction

Theorem (Bayes theorem) $\underbrace{p(\theta_n|M_n, y)}_{\text{posterior}} = \frac{\underbrace{p(y|\theta_n, M_n)}_{p(y|M_n)} \underbrace{p(y|M_n)}_{\text{marginal (averaged) likelihood}} \\ = \frac{p(y|\theta_n, M_n)p(\theta_n|M_n)}{\int p(y|\theta_n, M_n)p(\theta_n|M_n)d\theta_n}.$

Bayesian model comparison

• The log Bayes factor log BF_{ij} is obtained by taking the ratio of the log marginal likelihoods of the *i*-th and *j*-th models

$$\log BF_{ij} = \log p(y|M_i) - \log p(y|M_j)$$

=
$$\log \int p(y|\theta_i, M_i) p(\theta_i|M_i) d\theta_i - \log \int p(y|\theta_j, M_j) p(\theta_j|M_j) d\theta_j.$$

• $\log BF_{ij} > 1$ is in favour of model *i*.

Computational aspects

There is usually no analytic solution for the marginal likelihood. Thus, we use sampling-based methods:

- 1. Thermodynamic integration: Robust method for marginal likelihood estimation that does not require *a priori* choice of bridge/importance distribution.
- 2. Hamiltonian Monte Carlo: Scales better in high dimensions even when parameters show strong correlations.
- 3. Replica exchange: Accelerates chain mixing and can handle multimodality, which is inherent in ODE based models.

leading to Replica Exchange Hamiltonian Monte Carlo (REHMC).

We first define the *power posterior* which continuously connects the prior and posterior through the inverse temperature parameter β

$$p(y|\beta) = \int [p(y|\theta)]^{\beta} \pi(\theta) \, \mathrm{d}\theta, \qquad 0 \le \beta \le 1$$

Taking the logarithm and differentiating gives

$$\begin{aligned} \frac{\partial}{\partial\beta} \log p(y|\beta) &= \frac{1}{p(y|\beta)} \frac{\partial}{\partial\beta} p(y|\beta) \\ &= \frac{1}{p(y|\beta)} \int \frac{\partial}{\partial\beta} \left[p(y|\theta) \right]^{\beta} \pi(\theta) \, \mathrm{d}\theta \end{aligned}$$

Thermodynamic integration

Further simplifying with the identity $f'(x) = a^x \log a \iff f(x) = a^x$ gives

$$\frac{\partial}{\partial\beta}\log p(y|\beta) = \frac{1}{p(y|\beta)} \int \left[p(y|\theta)\right]^{\beta}\log p(y|\theta)\pi(\theta) \ d\theta$$
$$= \int \frac{\left[p(y|\theta)\right]^{\beta}\pi(\theta)}{p(y|\beta)}\log p(y|\theta) \ d\theta$$
$$= \mathbb{E}_{p(\theta|y,\beta)}[\log p(y|\theta)].$$

giving the final result:

$$\log p(y) = \int_0^1 \mathbb{E}_{p(\theta|y,\beta)}[\log p(y|\theta)] \ d\beta,$$

The trapezoidal + Monte Carlo estimate of the log marginal likelihood can then be written

$$\log p(y) \approx \sum_{j=1}^{N} \frac{(\beta_j - \beta_{j-1})}{2} \left[\frac{1}{S} \sum_{i=1}^{S} \log p(y|\theta_i, \beta_j) + \frac{1}{S} \sum_{i=1}^{S} \log p(y|\theta_i, \beta_{j-1}) \right],$$

where ${\cal N}$ is the number of integration points and ${\cal S}$ are the number of Monte Carlo samples.

Replica exchange Hamiltonian Monte Carlo (REHMC)



Effectiveness of REHMC



(c) Target distribution.

Results

- Two sets of experiments:
 - 1. Data generated from the model M_2 with the least number of parameters
 - 2. Data generated from model M_3 with the highest number of parameters.
- The precipitation and potential evapotranspiration are from the Magala Creek dataset.
- We assigned lognormal priors to all model parameters except $\sigma^2 \sim IG(\alpha, \beta)$.
- The parameters associated with upper buckets are assigned priors with faster timescales (runoff processes vs storage processes).

log marginal likelihood

• Experiment 1, M_2 is the data generating model.

Table 1: log marginal likelihood

M_2	M_3	M_4	
201.336	194.722	179.406	

• Experiment 2 M_3 is the data generating model.

Table 2: log marginal likelihood

M_2	M_3	M_4
74.815	158.716	152.581

Based on the BF interpretation table by (Kass & Raftery, 1995) we have decisive evidence in favour of the data generating models.

Posterior predictive checks



Figure 3: Graphical posterior predictive check

Real discharge data

Table 3: Results using real discharge data

	M ₂ (95% CI)	$M_3(95\%$ CI)	M4(95% CI)
k_1	1.281(0.893, 1.708)	1.255(0.851, 1.674)	1.298(0.863, 1.752)
k_2	1.506(0.860, 2.180)	1.863(1.142, 2.703)	2.060(1.227, 2.849)
k_3	-	1.342(0.719, 1.997)	1.408(0.775, 2.107)
k_4	-	-	1.072(0.574, 1.559)
$k_{1,2}$	1.182(0.788, 1.638)	2.296(0.589, 1.310)	2.292(1.325, 3.327)
$k_{2,3}$	-	0.711(0.481, 0.978)	0.731(0.451, 1.008)
$k_{3,4}$	-	-	0.828(0.541, 1.170)
\hat{V}_1	1.066(0.026, 2.776)	1.235(0.027, 3.336)	1.190(0.029, 3.154)
\hat{V}_2	0.821(0.061, 1.902)	1.077(0.061, 2.871)	0.997(0.059, 2.557)
\hat{V}_3	-	1.220(1.181, 0.029)	1.152(0.029 , 3.228)
\hat{V}_4	-	-	1.138(0.029, 3.066)
V_{\max}	0.841(0.585, 1.109)	0.941(0.635, 1.251)	0.842(0.595, 1.150)
σ^2	7.591(6.661, 8.787)	7.623(6.620, 8.697)	7.633(6.602, 8.750)
$\log p(y M)$	-388.826	-386.716	-388.978

- 1. We have introduced a modelling framework in hydrology that consists of parameter estimation, model selection, and posterior predictive checks.
- 2. We have illustrated using the gradient-based sampler REHMC that the marginal log-likelihood can be estimated efficiently for ODE-type models.
- 3. Our framework can be used as an efficient alternative to widely used gradient-free samplers.
- 4. The entire framework has been implemented in the differentiable programming language TensorFlow Probability.

Thank you

References

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