Bayesian Inference for DCMs

Will Penny

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Likelihood

We consider Bayesian estimation of nonlinear models of the form

$$y = g(heta, m) + e$$

where $g(\theta)$ is some nonlinear function, and *e* is zero mean additive Gaussian noise with covariance C_y . The likelihood of the data is therefore

$$p(y|\theta, \lambda, m) = N(y; g(\theta, m), C_y)$$

The error covariances are assumed to decompose into terms of the form

$$C_y^{-1} = \sum_i \exp(\lambda_i) Q_i$$

where Q_i are known precision basis functions and λ are hyperparameters.

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Priors

We allow Gaussian priors over model parameters

$$p(heta|m) = \mathsf{N}(heta; \mu_ heta, oldsymbol{C}_ heta)$$

where the prior mean and covariance are assumed known.

The hyperparameters are constrained by the prior

$$p(\lambda | m) = \mathsf{N}(\lambda; \mu_{\lambda}, C_{\lambda})$$

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VL Posteriors

The Variational Laplace (VL) algorithm assumes an approximate posterior density of the following factorised form

$$\begin{array}{lll} q(\theta,\lambda|y,m) &=& q(\theta|y,m)q(\lambda|y,m) \\ q(\theta|y,m) &=& \mathsf{N}(\theta;m_{\theta},S_{\theta}) \\ q(\lambda|y,m) &=& \mathsf{N}(\lambda;m_{\lambda},S_{\lambda}) \end{array}$$

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Energies

The above distributions allow one to write down an expression for the joint log likelihood of the data, parameters and hyperparameters

 $L(\theta, \lambda) = \log[p(y|\theta, \lambda, m)p(\theta|m)p(\lambda|m)]$

The approximate posteriors are estimated by minimising the Kullback-Liebler (KL) divergence between the true posterior and these approximate posteriors. This is implemented by maximising the following variational energies

$$I(\theta) = \int L(\theta, \lambda) q(\lambda)$$
$$I(\lambda) = \int L(\theta, \lambda) q(\theta)$$

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Gradient Ascent

This maximisation is effected by first computing the gradient and curvature of the variational energies at the current parameter estimate, $m_{\theta}(old)$. For example, for the parameters we have

$$j_{ heta}(i) = rac{dI(heta)}{d heta(i)}$$

 $H_{ heta}(i,j) = rac{d^2I(heta)}{d heta(i)d heta(j)}$

where *i* and *j* index the *i*th and *j*th parameters, j_{θ} is the gradient vector and H_{θ} is the curvature matrix. The estimate for the posterior mean is then given by

 $m_{ heta}(\textit{new}) = m_{ heta}(\textit{old}) + \Delta m_{ heta}$

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Adaptive Step Size

The change is given by

$$\Delta m_{\theta} = [\exp(vH_{\theta}) - I] H_{\theta}^{-1} j_{\theta}$$

This last expression implements a 'temporal regularisation' with parameter *v*. In the limit $v \to \infty$ the update reduces to

$$\Delta m_{ heta} = -H_{ heta}^{-1} j_{ heta}$$

which is equivalent to a Newton update. This implements a step in the direction of the gradient with a step size given by the inverse curvature. Big steps are taken in regions where the gradient changes slowly (low curvature).

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Likelihood

$$y(t) = -60 + V_a[1 - \exp(-t/\tau)] + e(t)$$



 $V_a = 30, \tau = 8, \exp(\lambda) = 1$

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Prior Landscape

A plot of log $p(\theta)$



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Samples from Prior

The true model parameters are unlikely apriori

$$V_{a}=$$
 30, $au=$ 8



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Posterior Landscape

A plot of $\log[p(y|\theta)p(\theta)]$



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Metropolis-Hastings

MH creates as series of random points $(\theta(1), \theta(2), ...)$ whose distribution converges to the target distribution of interest. For us, this is the posterior density $p(\theta|y)$. Each sequence can be considered a random walk whose stationary distribution is $p(\theta|y)$.

MH makes use of a proposal density $q(\theta'; \theta)$ which is dependent on the current state vector θ . For symmetric q(such as a Gaussian) samples from the posterior density can be generated as follows.

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MH update

First, start at some point $\theta(0)$ in parameter space. Then generate a proposal θ' using the density q. This proposal is then accepted according to the standard Metropolis-Hastings procedure.

That is, with probability min(1, r) where

$$r = rac{p(y| heta')p(heta')}{p(y| heta)p(heta)}$$

If the step is accepted we set $\theta(n + 1) = \theta'$. If it is rejected we set $\theta(n + 1) = \theta(n)$.

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Adaptive proposal density

We use a zero mean Gaussian proposal density with covariance C_s . This covariance is initialised to

$$C_s = \sigma C_{\theta}$$

where C_{θ} is the prior covariance and $\sigma = 1$.

We then use a three stage procedure comprising (i) scaling, (ii) tuning and (iii) sampling steps in which the scaling and tuning stages are used to optimize the proposal covariance C_s .

The first two stages are regarded as a burn-in phase and samples from this period are later discarded. At the end of this C_s is fixed and sampling proper begins.

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Scaling

The proposal covariance is given by

$$C_{s} = \sigma C_{\theta}$$

In the scaling step σ is adjusted as follows.

If the acceptance rate, as measured over the last $n_s = 100$ proposals, is less than 20 per cent then σ is halved.

If the acceptance rate is greater than 40 per cent σ is doubled.

Otherwise, σ remains unchanged.

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Tuning

The tuning step makes use of adaptive estimation of a covariance matrix C_{tune} based on a Robbins-Monro update.

At the beginning of the tuning stage we set $C_{tune} = C_s$. We then update according to

$$\mu_t = \mu_{t-1} + \frac{1}{n_t} (x_t - \mu_t)$$

$$\Delta C_{tune} = \frac{1}{n_t} [(x_t - \mu_t) (x_t - \mu_t)^T - C_{tune} (t-1)]$$
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where n_t is the number of elapsed iterations in the tuning period. At the end of tuning set $C_s = C_{tune}$.

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MH Samples



64,000 samples from MH posterior

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VL Samples

3.5 3 2.5 og t 2 1.5 1 0.5 2.5 3 3.5 Δ log V

64,000 samples from VL posterior

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Likelihood

Nonlinear oscillator with a = 0.2, b = 0.2, c = 3.

$$\dot{v} = c[v - \frac{1}{3}v^3 + r]$$
$$\dot{r} = -\frac{1}{c}[v - a + br]$$

We have noise level $exp(\lambda) = 10$.



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Priors

A plot of log $p(\theta)$



 $\mu_{ heta} = [-0.69, -0.69]^T, C_{ heta} = diag([1/8, 1/8]);$ $\mu_{\lambda} = 0, C_{\lambda} = 1$

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Priors

1.5 Ω 0.5 0<u>`</u>0 0.5 1.5 а

True value a = 0.2, b = 0.2 is apriori unlikely

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Posterior

A plot of $\log[p(y|\theta)p(\theta)]$



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VL optimisation I

Global maxima



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MH - Scaling

Init: [-0.2, -0.2]. Then 1000 samples



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MH - Tuning

1000 samples



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MH - Sampling

2000 samples



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Pat Benatar Interlude

Happy Birthday Jean !



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Model Evidence

The model evidence is not straightforward to compute, since this computation involves integrating out the dependence on model parameters

$$p(y|m) = \int p(y|\theta, m)p(\theta|m)d\theta.$$

Once computed two models can be compared via the Bayes factor

$$B_{12}=\frac{p(y|m_1)}{p(y|m_2)}$$

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Free Energy

The free energy is composed of sum squared precision weighted prediction errors and Occam factors

$$F = -\frac{1}{2} e_{y}^{T} C_{y}^{-1} e_{y} - \frac{1}{2} \log |C_{y}| - \frac{N_{y}}{2} \log 2\pi \qquad (8)$$

$$= -\frac{1}{2} e_{\theta}^{T} C_{\theta}^{-1} e_{\theta} - \frac{1}{2} \log \frac{|C_{\theta}|}{|S_{\theta}|}$$

$$= -\frac{1}{2} e_{\lambda}^{T} C_{\lambda}^{-1} e_{\lambda} - \frac{1}{2} \log \frac{|C_{\lambda}|}{|S_{\lambda}|}$$

where prediction errors are the difference between what is expected and what is observed

$$egin{array}{rcl} e_y &=& y-g(m_ heta) \ e_ heta &=& m_ heta-\mu_ heta \ e_\lambda &=& m_\lambda-\mu_\lambda \end{array}$$

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Prior Arithmetic Mean

The simplest approximation to the model evidence

$$p(y|m) = \int p(y|\theta, m)p(\theta|m)d\theta.$$

is the Prior Arithmetic Mean

$$p_{PAM}(y|m) = \frac{1}{S}\sum_{s=1}^{S}p(y|\theta_s,m)$$

where the samples θ_s are drawn from the prior density.

A problem with this estimate is that most samples from the prior will have low likelihood. A large number of samples will therefore be required to ensure that high likelihood regions of parameter space will be included in the average. Bayesian Inference for DCMs

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Posterior Harmonic Mean

A second option is the Posterior Harmonic Mean

$$p_{PHM}(y|m) = \left[rac{1}{S}\sum_{s=1}^{S}rac{1}{p(y| heta_s,m)}
ight]^{-1}$$

where samples are drawn from the posterior (eg. through MH sampling).

A problem with the PHM is that the largest contributions come from low likelihood samples which results in a high-variance estimator.

Both PAM and PHM can be motivated from the perspective of importance sampling.

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Savage-Dickey

For models 1 and 2 having common parameters θ_1 and model 2 having additional parameters θ_2 , then if

 $p(\theta_1|m_2) = p(\theta_1|m_1)$

the Bayes factor is given by

$$B_{12} = \frac{p(\theta_2 = 0|y, m_2)}{p(\theta_2 = 0|m_2)}$$



Here
$$B_{12} = 0.9$$
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Thermodynamic Integration

We define inverse 'temperatures' β_k such that

$$0 = \beta_0 < \beta_1 < .. < \beta_{k-1} < \beta_K = 1$$

For example

$$\beta_k = \left(\frac{k}{K}\right)^5$$

We also define

$$f_k(\theta) = p(y|\theta, m)^{\beta_k} p(\theta|m)$$

Sample from *k*th chain using MH with prob

$$r = \frac{f_k(\theta'_k)}{f_k(\theta_k)}$$

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Thermodynamic Integration

We can define the normalising constants

$$z_k = \int f_k(heta) d heta$$

where $z_0 = 1$ and $z_K = p(y|m)$. Now

$$\log p(y|m) = \log z_K - \log z_0$$

We can write this as

$$\log p(y|m) = \int_0^1 \frac{d \log z(\beta)}{d\beta} d\beta$$

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Thermodynamic Integration

The log evidence can therefore be approximated as

$$\log p_{TI}(y|m) = \sum_{k=1}^{K-1} (\beta_{k+1} - \beta_k) \left(\frac{E_{k+1} + E_k}{2} \right)$$

where

$$E_k = rac{1}{N_k} \sum_{s=1}^{N_k} \log p(y| heta_{ks})$$

where θ_{ks} is the *s*th sample from the *k*th chain.

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Free Energy Sampling Prior Arithmetic Mean Posterior Harmonic Mean Savage-Dickey

Thermodynamic Integration

DCM for fMRI

Conclusions

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Synthetic fMRI example

Design matrix from Henson et al. Regression coefficients from responsive voxel in occipital cortex. Data was generated from a 12-regressor model with SNR=0.2. We then fitted 12-regressor and 9-regressor models. This was repeated 25 times.



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Log Bayes factors

For these linear Gaussian models the free energy defaults to the exact model evidence. Bayes factors are therefore exact. This also holds for Savage-Dickey. The average true logBF was 3.45 in favour of the 12-regressor model.

The boxplots show estimated minus true logBF for each approach:



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Auditory DCMs





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DCM for fMRI

Chumbley et al (2007) 175,000 burn samples +...



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What Bayes factor results might look like !

Estimated logBF - *F_{diff}* for each approach:



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Conclusions

VL is good. Care with $p(\theta), p(\lambda)$.

Bottleneck for sampling methods is speed of function evaluation. For DCMs we can generate 2-20 samples/second per core. This equates to 7200 to 72000 samples per hour.

May be worth looking at Metropolis Adjusted Langevin Algorithms (MALA) - basically a stochastic VL with MH step.

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