MSA101/MVE187 2022 Lecture 11 Missing data / augmented data Hamiltonian MCMC

Petter Mostad

Chalmers University

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Review: Bayesian framework

- Prediction variable Y_{pred} , data Y_{data} , parameter θ .
- Specify a complete model by specifying prior $\pi(\theta)$, likelihood $\pi(Y_{data} \mid \theta)$, and prediction distribution $\pi(Y_{pred} \mid \theta)$.
- Derive the posterior $\pi(\theta \mid Y_{data})$.
- Make predictions using

$$\pi(Y_{pred} \mid Y_{data}) = \int \pi(Y_{pred} \mid \theta, Y_{data}) \pi(\theta \mid Y_{data}) \, d\theta$$

(in some cases $\pi(Y_{pred} \mid \theta, Y_{data}) = \pi(Y_{pred} \mid \theta))$.

If we cannot approximate the integral, we may instead simulate from

$$\pi(Y_{pred}, \theta \mid Y_{data})$$

and simply use the part of the simulated vectors that represent Y_{pred} .

Heart transplant example:

- Data: Exposures e_i and deaths y_i at i = 1,...,94 hospitals.
 Parameters: Mortality rates λ₁,..., λ₉₄ at the 94 hospitals.
 Underlying parameters μ, α. Prediction: E.g., prob that λ₁ < λ₂.
- Model: $y_i \sim \text{Poisson}(e_i \lambda_i)$, $\lambda_i \sim \text{Gamma}(\alpha, \alpha/\mu)$, $\mu \propto_{\mu} 1/\mu$, $\alpha \propto_{\alpha} 1/\alpha$.
- Draw the model as a graph!
- More examples:
 - The examples of conjugacy we started the course with
 - Second assignment!

Review: A hierarchical example

Data x_1, \ldots, x_8 and y_1, \ldots, y_6 are organized into groups, and we want to predict a value z_1 in a third group. We assume a model

$x_1,, x_8$	\sim	$Normal(\mu_1, au_1^{-1})$
y_1,\ldots,y_6	\sim	$Normal(\mu_2, au_1^{-1})$
<i>z</i> ₁	\sim	$Normal(\mu_3, au_1^{-1})$
μ_1, μ_2, μ_3	\sim	$Normal(10, au_0^{-1})$
$ au_0$	\sim	Gamma(1,4)
$ au_1$	\sim	Gamma(7, 3)

Draw the model!

- We simulate from the conditional distribution of all unknowns τ₀, τ₁, μ₁, μ₂, μ₃, z₁ after fixing the data values x₁,..., x₈, y₁,..., y₆.
- We then use the simulated values for z_1 to make predictions.
- For many hierarchical models, Gibbs sampling is a good way to simulate from the conditional distribution: See next overhead.

- The joint density for all variables in a hierarchical model is a product with one factor for each node in its graph.
- To find a function proportional to the posterior density for Y_{pred}, θ given Y_{data}, simply fix in this function all values corresponding to the data Y_{data}.
- To simulate from the resulting posterior density using Gibbs sampling, the conditional distribution for each unknown variable given all the others must be found.
- A function proportional to each such conditional density can be found by fixing all the other variables, and removing constant factors.
- The resulting functions will only have terms from original factors corresponding to edges in or out of the variable in the graph!

Inference for hierarchical models

- 1. Write down the joint probability of all variables $(Y_{data}, Y_{pred}, \text{ and } \theta)$ as a product over all the nodes in the hierarchical model.
- 2. If possible, try to analytically marginalize over some of the components of θ .
- 3. In the expression from (1), fix all values in Y_{data} , to obtain a function proportional to the joint posterior Y_{pred} , $\theta \mid Y_{data}$. (Remove constant factors).
- 4. Generate an (approximate) sample from the remaining variables; the dimensions corresponding to Y_{pred} represent an approximate sample for your prediction.
- It may be convenient to use a Metropolis Hastings with symmetric proposals, or:
- Many or all of the conditional densities needed for a Gibbs sampler can often be found easily.
- If some conditional densities cannot be found easily, one may for these use a standard Metropolis Hastings proposal function (and acceptance probability).

Review: Conditional distributions for the example

The conditional distributions become (prove yourself!)

$$\begin{array}{rcl} \mu_{1} \mid x_{1}, \dots, x_{8}, \tau_{1}, \tau_{0} & \sim & \operatorname{Normal}\left(\frac{10\tau_{0} + 8\overline{x}\tau_{1}}{\tau_{0} + 8\tau_{1}}, \frac{1}{\tau_{0} + 8\tau_{1}}\right) \\ \mu_{2} \mid y_{1}, \dots, y_{6}, \tau_{1}, \tau_{0} & \sim & \operatorname{Normal}\left(\frac{10\tau_{0} + 6\overline{y}\tau_{1}}{\tau_{0} + 6\tau_{1}}, \frac{1}{\tau_{0} + 6\tau_{1}}\right) \\ \mu_{3} \mid z_{1}, \tau_{1}, \tau_{0} & \sim & \operatorname{Normal}\left(\frac{10\tau_{0} + z_{1}\tau_{1}}{\tau_{0} + \tau_{1}}, \frac{1}{\tau_{0} + \tau_{1}}\right) \\ \tau_{0} \mid \mu_{1}, \mu_{2}, \mu_{3} & \sim & \operatorname{Gamma}\left(1 + \frac{3}{2}, 4 + \frac{1}{2}\sum_{i=1}^{3}(\mu_{i} - 10)^{2}\right) \\ \tau_{1} \mid \mu_{1}, \mu_{2}, \mu_{3}, x_{1} \dots x_{8}, y_{1} \dots y_{6}, z_{1} & \sim & \operatorname{Gamma}\left(7 + \frac{15}{2}, 3 + \frac{1}{2}\sum_{i=1}^{8}(x_{i} - \mu_{1})^{2} \\ + \frac{1}{2}\sum_{i=1}^{6}(y_{i} - \mu_{2})^{2} + \frac{1}{2}(z_{1} - \mu_{3})^{2}\right) \\ z_{1} \mid \mu_{3}, \tau_{1} & \sim & \operatorname{Normal}(\mu_{3}, \tau_{1}^{-1}) \end{array}$$

Missing data / augmented data

- Assume some data values are *censored*: You don't know them exactly, only that they are (for example) above some threshold. How to deal with this?
- Example application: Survival analysis. You want to know how long people live after some event. But some people are still alive at the end of the study (or they died from other causes).
- We want to learn about density f(· | θ) from sample where x₁,..., x_k are observed values and c₁,..., c_n are observations that the corresponding x_i is greater than some a_i. The likelihood becomes

$$\pi(x_1,\ldots,x_k,c_1,\ldots,c_n\mid\theta)=\prod_{i=1}^k f(x_i\mid\theta)\prod_{i=1}^n (1-F(a_i\mid\theta))$$

where $F(\cdot \mid \theta)$ is the cumulative distribution function.

- > You may simulate from the posterior for θ using for example random walk MH.
- ALTERNATIVELY: You may add to the model variables representing the censored values, and simulate these together with the unknown θ.

- In many classical statistical methods, missing data may present a problem.
- The standard Bayesian answer in such cases: Add to the model random variables representing the unobserved values, and simulate them together with parameters and other variables of interest.
- This solves the problem in theory, but may of course sometimes be difficult in practice.

Example: Augmented data

Example (7.7. in RC): In a genetics problem, one wants to know how close two genes are on the chromosome, measured by a parameter θ. Given n individuals, the number of individuals x₁, x₂, x₃, x₄ in each of 4 categories will be multinomially distributed accoring to

$$(x_1, x_2, x_3, x_4) \mid \theta \sim \mathsf{Multinomial}\left(n, \frac{1}{2} + \frac{\theta}{4}, \frac{1}{4}(1-\theta), \frac{1}{4}(1-\theta), \frac{\theta}{4}\right)$$

Given a prior on θ, how do you simulate from the posterior?
The likelihood for θ makes necessary approximate or numerical simulation:

$$\pi(x_1,\ldots,x_4\mid heta) \propto_ heta \left(rac{1}{2}+rac{ heta}{4}
ight)^{x_1} \left(rac{1}{4}(1- heta)
ight)^{x_2} \left(rac{1}{4}(1- heta)
ight)^{x_3} \left(rac{ heta}{4}
ight)^{x_4}$$

• We extend the data (x_1, x_2, x_3, x_4) with a latent variable z, so that

$$(z, x_1-z, x_2, x_3, x_4) \mid heta \sim \mathsf{Multinomial}\left(n; rac{1}{2}, rac{ heta}{4}, rac{1}{4}(1- heta), rac{1}{4}(1- heta), rac{ heta}{4}
ight)$$

The likelihood becomes

$$\pi(z, x_1, \dots, x_4 \mid heta) \propto_{ heta} heta^{x_1 - z + x_4} (1 - heta)^{x_2 + x_3}$$

- Note that, with the augmented data (z, x₁, x₂, x₃, x₄), the likelihood has the Beta family of densities as conjugate priors! Assume, for example, θ ~ Beta(α, β).
- You can now use Gibbs sampling to sample from the distribution π(z, θ | x₁,..., x₄):

•
$$\theta \mid z, x_1, x_2, x_3, x_4 \sim \text{Beta}(\alpha + x_1 - z + x_4, \beta + x_2 + x_3).$$

$$\triangleright z \mid \theta, x_1, x_2, x_3, x_4 \sim \text{Binomial}\left(x_1, \frac{\frac{1}{2}}{\frac{1}{2} + \frac{\theta}{4}}\right).$$

Exercise: Derive the Binomal distribution for *z* above.

- We have looked at several ideas for constructing good proposal densities. Somehow, they take into account the properties of the target density.
- Can one construct general methods that "automatically" learns about the target density and makes good proposals based on that?
- Several methods exist that do this; they have varying degrees of success with good convergence.
- We will look at one quite popular and clever method: Hamiltonian Monte Carlo.

Hamiltonian Monte Carlo: Idea

We are given a posterior density $\pi(q) \propto_q \exp(-U(q))$ for vectors $q = (q_1, \ldots, q_d)$. We want to find a smart proposal function that utilizes U:

- Look at U(q) as some kind of "potential energy" for a particle that can move between different q's.
- If the particle moves so that it looses potential energy, it gains kinetic energy, i.e., it moves faster.
- If the particle moves in this way, it will move faster in the direction of higher density for $\pi(q)$.
- Idea: As a proposal function, randomly generate a direction and a speed for the particle to move from the current q. Then let the particle move according to dynamics above for time period s.
- Below, we use pairs (p, q) of particle momentum $p = (p_1, \ldots, p_d)$ and particle position q, moving the particle so that the total energy

$$H(p,q) = U(q) + rac{1}{2}\sum_{i=1}^d rac{p_i^2}{\sigma_i^2}$$

is kept constant (where we may set $\sigma_i^2 = m$ where *m* is the "mass").

Assume $\pi(x)$ is a density and T_1 , T_2 are invertible transformations on the set of possible x values satisfying for all x

$$\pi(T_1(x)) = \pi(x)$$
 $\pi(T_2(x)) = \pi(x)$ $T_1(T_2(x)) = T_2^{-1}(T_1^{-1}(x)).$

Then the deterministic M-H proposal function $T_2(T_1(x))$ has acceptance probability 1.

Proof: We have symmetry

$$T_2(T_1(T_2(T_1(x)))) = T_2(T_2^{-1}(T_1^{-1}(T_1(x)))) = x$$

and invariance of the density

$$\pi(T_2(T_1(x))) = \pi(T_1(x)) = \pi(x).$$

Hamiltonian dynamics

Given a "Hamiltonian" function H(p,q), with $p,q \in \mathbb{R}^d$, $H(p,q) \in \mathbb{R}$.

- A particle $p : \mathbb{R} \to \mathbb{R}^{2d}$, with p(t) = (p, q), is said to have "position" q and "momentum" p at time t.
- The particle follows Hamiltonian dynamics if, for i = 1, ..., d,

$$\frac{dq_i}{dt} = \frac{\partial H}{\partial p_i}$$
 and $\frac{dp_i}{dt} = -\frac{\partial H}{\partial q_i}$. (1)

- After time s, the particle starting at position q with momentum p will have position q* and momentum p*. So the solution to Equations 1 defines a mapping T₁ sending (p, q) to (p*, q*).
- We have $H(T_1(p,q)) = H(p,q)$ because

$$\frac{dH}{dt} = \sum_{i=1}^{d} \left(\frac{dq_i}{dt} \frac{\partial H}{\partial q_i} + \frac{dp_i}{dt} \frac{\partial H}{\partial p_i} \right) = \sum_{i=1}^{d} \left(\frac{\partial H}{\partial p_i} \frac{\partial H}{\partial q_i} - \frac{\partial H}{\partial q_i} \frac{\partial H}{\partial p_i} \right) = 0.$$

If H(−p, q) = H(p, q) for all (p, q) then Equations 1 are unchanged if we simultaneously switch the signs of t and p. Defining T₂(p, q) = (−p, q) we get T₁(T₂(p, q)) = T₂⁻¹T₁⁻¹(p, q). Assume $\pi(p,q) \propto \exp(-H(p,q))$ with H(-p,q) = H(p,q) as above.

- ► A Metropolis Hastings algorithm alternating between the proposals
 - simulate p using $\pi(p \mid q) \propto \exp(-H(p,q))$
 - compute $T_2(T_1(p,q))$

and always accepting will provide an approximate sample from $\pi(p,q)$ provided the Markov chain is ergodic.

- If H(p, q) = V(p) + U(q) with V(−p) = V(p) then p and q are independent, and π(p | q) ∝ exp(−V(p)) is simulated independently in each iteration.
- ▶ By throwing away the part of the sample concerning p we get an approximate sample from $\pi(q) \propto \exp(-U(q))$.

Hamiltonian Monte Carlo algorithm

- ▶ Start with a density $\pi(q) \propto \exp(-U(q))$ you want to simulate from.
- Define $V(p) = \frac{1}{2} \sum_{i=1}^{d} p_i^2 / \sigma_i^2$ for some $\sigma_1, \ldots, \sigma_d$.
- Find an initial value $q^{(0)}$ for q.
- For each iteration, given $q^{(j)}$:
 - Simulate $p_i \sim \text{Normal}(0, \sigma_i^2)$
 - Compute $(p^*, q^*) = T_1(p, q^{(j)})$
 - Set $q^{(j+1)} = q^*$ and throw away p^* .

It "just" remains to find an efficient way to compute T₁(p, q^(j)).
 Note that

$$\frac{dq_i}{dt} = \frac{\partial H}{\partial p_i} \qquad \text{becomes} \qquad \frac{dq_i}{dt} = \frac{p_i}{\sigma_i^2}$$
$$\frac{dp_i}{dt} = -\frac{\partial H}{\partial q_i}. \qquad \text{becomes} \qquad \frac{dp_i}{dt} = -\frac{\partial U}{\partial q_i}.$$

and

The Leapfrog algorithm: A numerical approximation of T_1

▶ For simplicity set all $\sigma_i = 1$ and use vector notation: We need that

$$rac{dq}{dt}=p$$
 and $rac{dp}{dt}=-
abla U(q)$

- Let $q_0, q_1, q_2, \ldots, q_n$ be the values of q along the particle path at times $0, \frac{s}{n}1, \frac{s}{n}2, \ldots, \frac{s}{n}n = s$, respectively.
- ▶ Let $p_0, p_1, p_2, \ldots, p_{n+1}$ be the values of p along the particle path at times $0, \frac{s}{n}(1-\frac{1}{2}), \frac{s}{n}(2-\frac{1}{2}), \ldots, \frac{s}{n}(n-\frac{1}{2}), s$, respectively.
- Approximate $\frac{dq}{dt} = p$ with

$$\frac{q_{j+1}-q_j}{s/n} = p_{j+1}$$
 $j = 0, \dots, n-1.$

• Approximate $\frac{dp}{dt} = -\nabla U(q)$ with

$$\frac{p_{j+1}-p_j}{s/n}=-\nabla U(q_i) \qquad j=1,\ldots,n-1.$$

while using half stepsize for j = 0 and j = n.

► We get $p_1 = p_0 - (s/2n)\nabla U(q_0)$, $p_{n+1} = p_n - (s/2n)\nabla U(q_n)$, and $q_{j+1} = q_j + (s/n)p_{j+1}$ $p_{j+1} = p_j - (s/n)\nabla U(q_j)$

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- Note: *n* computations of the gradient ∇U must be done: Possible? Time consuming?
- Note: As this is an approximation, we only have that H(p^{*}, q^{*}) ≈ H(p, q). But this is no problem, as we can compute and use the standard acceptance probability for Metropolis Hastings proposals.
- Note: You must still check that the Markov chain is Ergodic: In practice, that the algorithm can reach any q from any q.
- Can give great fast convergence in the cases where the gradient of the logged density is easily available and computable.
- For more information see for example Neal (2011) "MCMC Using Hamiltonian Dynamics".