

Advanced Simulation - Lecture 10

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- Often we have various possible models for the same dataset.
- Sometimes there's an infinity of possible models!
- How to choose between models?

Green (1995), *Reversible Jump Markov chain Monte Carlo and Bayesian model determination*.

Motivation: Bayesian model choice

- Assume we have a collection of models \mathcal{M}_k for $k \in \mathcal{K}$.
- With data we can learn parameters given each model \mathcal{M}_k , but we can also learn about the models.
- Put a prior on models \mathcal{M}_k . Within each model, prior $p(\theta_k | \mathcal{M}_k)$ on the parameters.
- Joint posterior distribution of interest:

$$\pi(\mathcal{M}_k, \theta_k | y) = \pi(\mathcal{M}_k | y)\pi(\theta_k | y, \mathcal{M}_k)$$

which is defined on

$$\cup_{k \in \mathcal{K}} \{\mathcal{M}_k\} \times \Theta_k \equiv \cup_{k \in \mathcal{K}} \{k\} \times \Theta_k.$$

Polynomial regression

- Data $(x_i, y_i)_{i=1}^n$ where $(x_i, y_i) \in \mathbb{R} \times \mathbb{R}$.
- Polynomial regression model

$$\mathcal{M}_k : y = \underbrace{\sum_{j=0}^k \beta_j x^j}_{=f(x;\beta)} + \varepsilon, \quad \varepsilon \sim \mathcal{N}(0, \sigma^2).$$

- If k is too large then

$$f(x; \hat{\beta}) = \sum_{j=0}^k \hat{\beta}_j x^j$$

where $\hat{\beta} = (\hat{\beta}_0, \hat{\beta}_1, \dots, \hat{\beta}_k)$ is the MLE, will overfit.

Polynomial regression

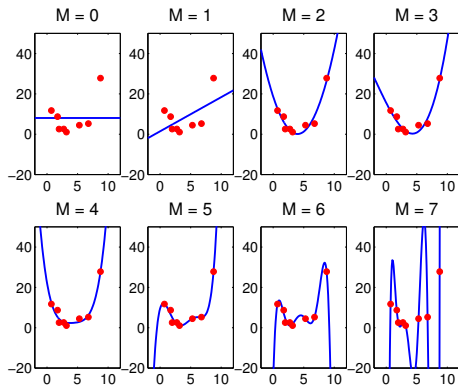


Figure: As order of the model $M = k$ increases, we overfit.

Bayesian polynomial regression

- We select $k \in \{0, \dots, M_{\max}\}$ and

$$\mathbb{P}(\mathcal{M}_k) = p_k = \frac{1}{M_{\max} + 1}$$

with $\Theta_k = \mathbb{R}^{k+1} \times \mathbb{R}^+$

$$p_k(\beta, \sigma^2) = \mathcal{N}(\beta; 0, \sigma^2 I_{k+1}) \mathcal{IG}(\sigma^2; 1, 1).$$

- In this case, we have analytic expression for

$$p_k(y_{1:n}) = \int_{\Theta_k} p_k(\beta, \sigma^2) \prod_{i=1}^n \mathcal{N}(y_i; f(x_i; \beta), \sigma^2) d\beta d\sigma^2.$$

- Bayesian model selection automatically prevents overfitting.

Bayesian Polynomial regression

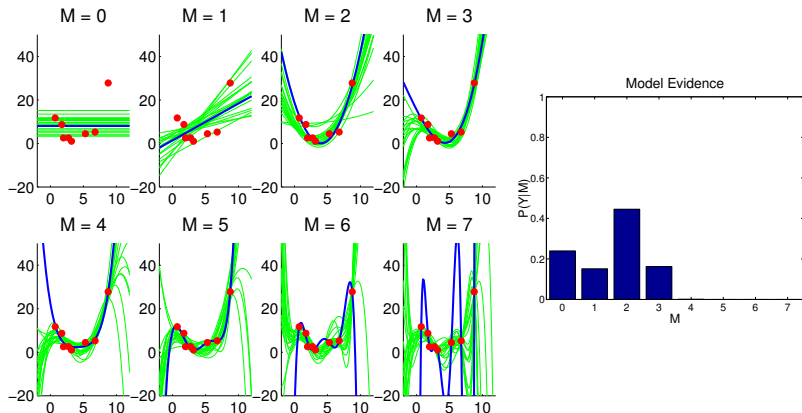


Figure: $f(x; \beta)$ for random draws from $p_M(\beta | y_{1:n})$ and evidence $p_M(y_{1:n})$.

Motivation: mixture models

- Assume the observations Y_1, \dots, Y_n come from

$$\sum_{k=1}^K p_k \mathcal{N}(\mu_k, \sigma_k^2)$$

with $\sum_{k=1}^K p_k = 1$. For any fixed K , the parameters to infer are $(p_1, \dots, p_{K-1}, \mu_1, \dots, \mu_K, \sigma_1^2, \dots, \sigma_K^2)$ of dimension $3K - 1$.

- But what about inference on K ?
- We can put a prior on K , e.g. a Poisson distribution.
- How do we get the posterior?

Sampling in transdimensional spaces

- Consider a collection of models \mathcal{M}_k , for $k \in \mathcal{K} \subset \mathbb{N}$.
- We want to design a Markov chain taking values in $\cup_{k \in \mathcal{K}} \{k\} \times \Theta_k$, with the correct joint posterior.
- Reversible jump MCMC is a generalized Metropolis-Hastings using a mixture of kernels.
- For each k , standard MH kernel from $\{k\} \times \Theta_k$ to $\{k\} \times \Theta_k$, i.e. standard within-model moves.
- How to move from $\{k\} \times \Theta_k$ to $\{k'\} \times \Theta_{k'}$?

Transdimensional moves

We can propose k' from $q(k' | k)$. Then we need to propose a move from Θ_k to $\Theta_{k'}$, of dimension d_k and $d_{k'}$.

- **dimension matching:** extend the spaces with auxiliary variables.
- Introduce $u_{k \rightarrow k'}$ and $u_{k' \rightarrow k}$ with distributions $\varphi_{k \rightarrow k'}$ and $\varphi_{k' \rightarrow k}$ respectively, and such that

$$d_k + \dim(u_{k \rightarrow k'}) = d_{k'} + \dim(u_{k' \rightarrow k}).$$

Transdimensional moves

- Given θ_k , we sample $u_{k \rightarrow k'} \sim \varphi_{k \rightarrow k'}$ and then apply a deterministic mapping to get

$$(\theta_{k'}, u_{k' \rightarrow k}) = G_{k \rightarrow k'}(\theta_k, u_{k \rightarrow k'}).$$

- The distributions φ are arbitrary and $G_{k \rightarrow k'}$ has to be a diffeomorphism.
- We now have our proposal from Θ_k to $\Theta_{k'}$. With what probability do we accept it?

Transdimensional moves

- Mimicking Metropolis-Hastings, given x we propose a point x' and accept or not with probability $\alpha(x \rightarrow x')$.
- We want P to be such that, for all A, B :

$$\int_{x, x' \in A \times B} \pi(dx) P(x \rightarrow dx') = \int_{x, x' \in A \times B} \pi(dx') P(x' \rightarrow dx)$$

or equivalently

$$\begin{aligned} & \int_{x, x' \in A \times B} \pi(dx) q(x \rightarrow dx') \alpha(x \rightarrow x') \\ &= \int_{x, x' \in A \times B} \pi(dx') q(x' \rightarrow dx) \alpha(x' \rightarrow x) \end{aligned}$$

Transdimensional moves

- Subtle point: $\pi(dx)P(x, dx')$ does not necessarily admit a density with respect to a standard measure.
- We cannot write e.g.

$$\pi(x)P(x, dx') = \pi(x)P(x, x')dx dx'$$

- However $\pi(dx)q(x, dx')$ can be assumed to be dominated and we write

$$\pi(x)q(x, dx') = \pi(x)q(x, x')dx dx'$$

Transdimensional moves

- First term is:

$$\int_{x,x' \in A \times B} \pi(x) q(x \rightarrow x') \alpha(x \rightarrow x') dx dx'$$

- Suppose we propose x' by sampling $u \sim \varphi$ and then taking $(x', u) = G(x, u)$ deterministically. We write $x'(x, u)$ and $u'(x, u)$.
- The expression becomes

$$\int_{x,x'(x,u) \in A \times B} \pi(x) \varphi(u) \alpha(x \rightarrow x'(x, u)) dx du$$

- What is the reverse transition from x' to x ? Sample $u' \sim \varphi'$ and take $(x, u) = G^{-1}(x', u')$.

Transdimensional moves

- Second term was:

$$\int_{x, x' \in A \times B} \pi(x') q(x' \rightarrow x) \alpha(x' \rightarrow x) dx dx'$$

- It becomes, with $(x, u) = G^{-1}(x', u')$:

$$\int_{x(x', u'), x' \in A \times B} \pi(x') \varphi'(u') \alpha(x' \rightarrow x(x', u')) dx' du'$$

Let us do a change of variable to get an integral with respect to $dx du$ instead of $dx' du'$:

$$\int \pi(x'(x, u)) \varphi'(u'(x, u)) \alpha(x'(x, u) \rightarrow x) \left| \frac{\partial G(x, u)}{\partial(x, u)} \right| dx du$$

- We see that the integrals are equal if

$$\begin{aligned} & \pi(x)\varphi(u)\alpha(x \rightarrow x'(x, u)) \\ &= \pi(x'(x, u))\varphi'(u'(x, u))\alpha(x'(x, u) \rightarrow x) \left| \frac{\partial G(x, u)}{\partial(x, u)} \right| \end{aligned}$$

- Thus we can see a valid choice of $\alpha(x \rightarrow x')$ in :

$$\alpha(x \rightarrow x') = \min \left(1, \frac{\pi(x')\varphi'(u')}{\pi(x)\varphi(u)} \left| \frac{\partial G(x, u)}{\partial(x, u)} \right| \right)$$

Transdimensional moves

We can now answer the initial question:

- How to move from $\{k\} \times \Theta_k$ to some other $\{k'\} \times \Theta_{k'}$? We start from some (k, θ_k) .
- Sample $k' \sim q(k \rightarrow k')$, then sample $u_{k \rightarrow k'}$ from $\varphi_{k \rightarrow k'}$.
- Compute deterministically $(\theta_{k'}, u_{k' \rightarrow k}) = G_{k \rightarrow k'}(\theta_k, u_{k \rightarrow k'})$.
- Compute

$$\alpha_{k \rightarrow k'} = \min \left(1, \frac{\pi(\theta_{k'}) \varphi_{k' \rightarrow k}(u_{k' \rightarrow k})}{\pi(\theta_k) \varphi_{k \rightarrow k'}(u_{k \rightarrow k'})} \frac{q(k' \rightarrow k)}{q(k \rightarrow k')} J_{k \rightarrow k'}(\theta_k, u_{k \rightarrow k'}) \right)$$

where

$$J_{k \rightarrow k'}(\theta_k, u_{k \rightarrow k'}) = \left| \frac{\partial G_{k \rightarrow k'}(\theta_k, u_{k \rightarrow k'})}{\partial(\theta_k, u_{k \rightarrow k'})} \right|.$$

Reversible Jump algorithm

- Starting with $(k^{(0)}, \theta^{(0)})$ iterate for $t = 1, 2, 3, \dots$
- With probability β , set $k^{(t)} = k^{(t-1)}$ and do one step of $K_{k^{(t)}}$ leaving $\pi(\theta_{k^{(t)}} | y, \mathcal{M}_{k^{(t)}})$ invariant.
- With probability $1 - \beta$, propose $k' \sim q(k' | k^{(t-1)})$.
 - Draw a random variable $u_{k^{(t-1)} \rightarrow k'} \sim \varphi_{k^{(t-1)} \rightarrow k'}$.
 - Apply the deterministic mapping $G_{k^{(t-1)} \rightarrow k'}$ to get θ', u' .
 - With “between-models” acceptance probability $a(\theta^{(t-1)} \rightarrow \theta')$:
accept, i.e. set $\theta^{(t)} = \theta', k^{(t)} = k'$,
otherwise reject, i.e. set $\theta^{(t)} = \theta^{(t-1)}, k^{(t)} = k^{(t-1)}$.

Toy example

- Two models, uniform prior on models $p(\mathcal{M}_1) = p(\mathcal{M}_2) = \frac{1}{2}$.
- In model \mathcal{M}_1 , $\theta \in \mathbb{R}$ and we can evaluate pointwise

$$\text{posterior}_1(\theta) \propto p(\theta | \mathcal{M}_1)\mathcal{L}(\theta | \mathcal{M}_1) = \exp\left(-\frac{1}{2}(\theta)^2\right)$$

- In model \mathcal{M}_2 , $\theta \in \mathbb{R}^2$ and we can evaluate pointwise

$$\text{posterior}_2(\theta) \propto p(\theta | \mathcal{M}_2)\mathcal{L}(\theta | \mathcal{M}_2) = \exp\left(-\frac{1}{2}(\theta_1)^2 - \frac{1}{2}(\theta_2)^2\right)$$

- In terms of model comparison, we should find

$$\begin{aligned}\frac{p(\mathcal{M}_2 | y)}{p(\mathcal{M}_1 | y)} &= \frac{p(y | \mathcal{M}_2)p(\mathcal{M}_2)}{p(y | \mathcal{M}_1)p(\mathcal{M}_1)} \\ &= \frac{\int_{\mathbb{R}^2} p(\theta | \mathcal{M}_2)\mathcal{L}(\theta | \mathcal{M}_2)d\theta}{\int_{\mathbb{R}} p(\theta | \mathcal{M}_1)\mathcal{L}(\theta | \mathcal{M}_1)d\theta} \times \frac{\frac{1}{2}}{\frac{1}{2}} \\ &= \frac{2\pi}{\sqrt{2\pi}} \\ &= \sqrt{2\pi} \approx 2.5066\end{aligned}$$

- In terms of parameters, in model \mathcal{M}_1 , $\theta \sim \mathcal{N}(0, 1)$ and in model \mathcal{M}_2 , $\theta \sim \mathcal{N}\left(\begin{pmatrix} 0 \\ 0 \end{pmatrix}, \begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix}\right)$.

Reversible Jump algorithm

We need to construct various Markov kernels.

- A Markov kernel “within \mathcal{M}_k ” for each model \mathcal{M}_k .

Toy example: introduce a Metropolis Hastings with random walk proposal, of variance σ^2 for model \mathcal{M}_1 and Σ for model \mathcal{M}_2 .

- A Markov kernel to move between models, i.e. for each pair k , proposing k' and proposing to move parameters of \mathcal{M}_k to parameters of $\mathcal{M}_{k'}$.

Toy example: introduce K_{12} moving a parameter $\theta \in \mathbb{R}$ to a parameter $(\theta_1, \theta_2) \in \mathbb{R}^2$, and introduce K_{21} moving a parameter $(\theta_1, \theta_2) \in \mathbb{R}^2$ to a parameter $\theta \in \mathbb{R}$.

Toy example

For K_{12} do the following.

- Sample u from $\mathcal{C}(0, 1)$, a standard Cauchy (*dimension matching*).
- Map deterministically $(\theta_1, \theta_2) = G_{1 \rightarrow 2}(\theta, u) = (\theta, u)$, with Jacobian equal to 1.
- Compute

$$\alpha_{1 \rightarrow 2} = \min \left(1, \frac{\exp(-0.5\theta^2 - 0.5u^2)}{\exp(-0.5\theta^2)\mathcal{C}(u; 0, 1)} \right)$$

Indeed the Jacobian is equal to 1, the priors on \mathcal{M}_1 and \mathcal{M}_2 are identical, and $q(k' | k) = q(k | k')$.

- Accept θ_1, θ_2 or stay at θ .

Toy example

For K_{21} do the following.

- Map deterministically $(\theta, u) = G_{2 \rightarrow 1}(\theta_1, \theta_2) = (\theta_1, \theta_2)$, with Jacobian equal to 1.
- Compute

$$\alpha_{2 \rightarrow 1} = \min \left(1, \frac{\exp(-0.5\theta_1^2) \mathcal{C}(\theta_2; 0, 1)}{\exp(-0.5\theta_1^2 - 0.5\theta_2^2)} \right)$$

Indeed the Jacobian is equal to 1, the priors on \mathcal{M}_1 and \mathcal{M}_2 are identical, and $q(k' | k) = q(k | k')$.

- Accept θ or stay at (θ_1, θ_2) .

Reversible Jump algorithm

- Introduce a probability of performing a “between-model” move at each step, say $\beta \in [0, 1]$.
- Given the current state of the chain k_t, θ_t at time t :
 - with probability β , between-model move: draw (k_{t+1}, θ_{t+1}) by drawing $k' \sim q(k' | k)$, dimension matching, deterministic mapping, RJ acceptance ratio. . .
 - with probability $1 - \beta$, within-model move: standard Metropolis-Hastings in the current model.

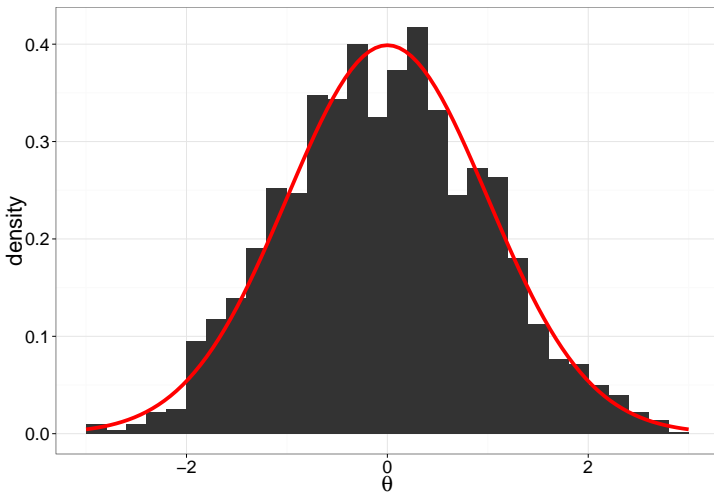


Figure: Parameter θ in model \mathcal{M}_1 .

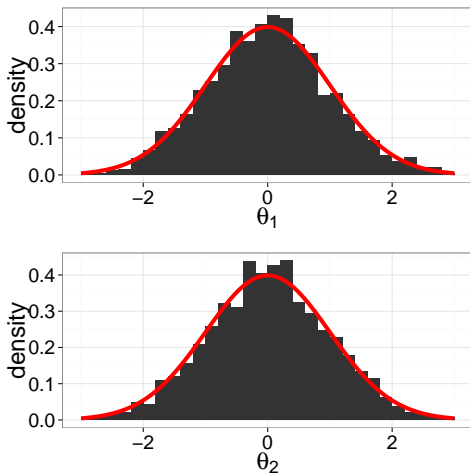


Figure: Parameter (θ_1, θ_2) in model \mathcal{M}_2 .

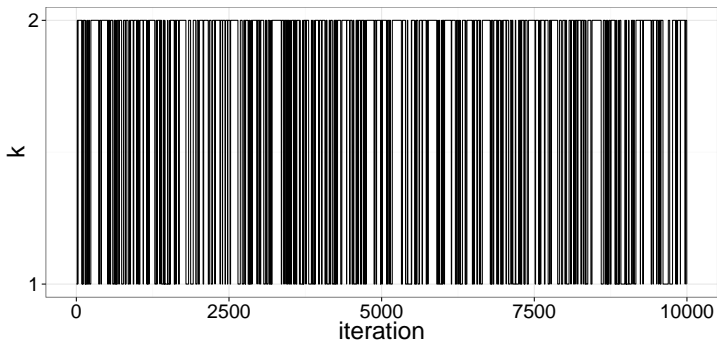


Figure: Model index k along iterations. Probability of accepting model jumps: $\approx 43.6\%$. The number of visits in \mathcal{M}_2 divided by the number of visits in \mathcal{M}_1 equals ≈ 2.39 , approximating the Bayes factor of ≈ 2.51 .

Results

If instead of $\mathcal{C}(0, 1)$ we use $\mathcal{N}(3, 1)$ for the dimension matching variable.

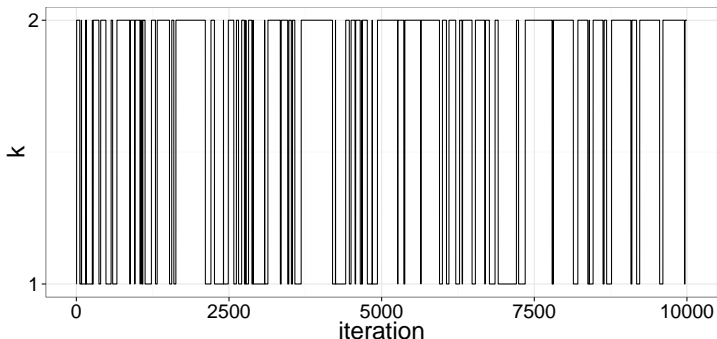


Figure: Model index k along iterations. Probability of accepting model jumps: $\approx 12.2\%$. Bayes factor approximated by ≈ 2.21 .

Results

If instead of $\mathcal{C}(0, 1)$ we use $\mathcal{N}(5, 1)$ for the dimension matching variable.

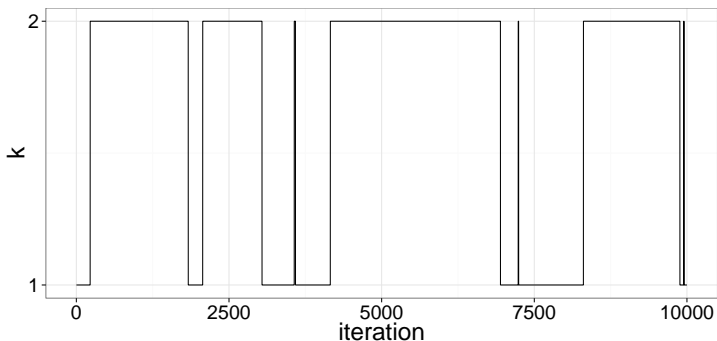


Figure: Model index k along iterations. Probability of accepting model jumps: $\approx 1.43\%$. Bayes factor approximated by ≈ 2.31 (not so bad!).

Reversible Jump algorithm: conclusion

- Probably the most ambitious MCMC algorithm, aiming at parameter estimation and model choice in one run.
- In general it's hard to design auxiliary variables for dimension matching and deterministic mappings such that the acceptance rate of between-model moves is decent.
- Transdimensional samplers constitute an on-going research area, see for instance:
Annealed Importance Sampling Reversible Jump MCMC Algorithms, by Karagiannis and Andrieu, 2013.