

STA 250 Class Note - Oct. 21

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I. Metropolis-Within-Gibbs

- Allows you to sample from a high-dimensional distribution using a sequence of lower dimensional distributions.
 - Generalize Gibbs Sampler in two ways:
 - If we cannot sample directly (exactly), then you can use MH.
 - Sample sub-blocks of parameters, not necessarily full conditionals.
i.e $P(\theta_1, \theta_2, \phi | \vec{y}) \rightarrow$ sample $P(\theta_1, \theta_2 | \phi, \vec{y}) \rightarrow$ sample $P(\phi | \theta_1, \theta_2, \vec{y})$
- [**Note:** Idea for homework: 2 strategies – (a) Sampling from $P(\vec{\beta} | \vec{y})$ using MH and a multivariate proposal for $\vec{\beta}$. (b) For $j = 1, \dots, p$ sample β_j from $P(\beta_j | \beta_{[-j]}, \vec{y})$.]
- When to use non-symmetric proposals?
 - If θ has compact support, then a non-symmetric proposal can be used to respect the boundaries.
 - example:
 $X \sim_{iid} Poi(\lambda)$, $i = 1, \dots, n$, prior $\lambda \sim I_{[0, \infty]}$.
Use MH proposal $\theta^* \sim TN(\theta^{(t)}, \nu^2, [0, \infty)) \implies$ No longer symmetric!!

II. Checking Your MCMC Codes

For one dataset, how to know if we reach convergence?

- (i) Trace plot, "by eye".
- (ii) Use ESS (effective sample size) to gauge roughly "how well" a chain converges (i.e., the equivalent number of independent samples).
- (iii) Other diagnostics.
 - Gelman-Rubin (Multiple chains)
 - Heidelberger
 - Geweke

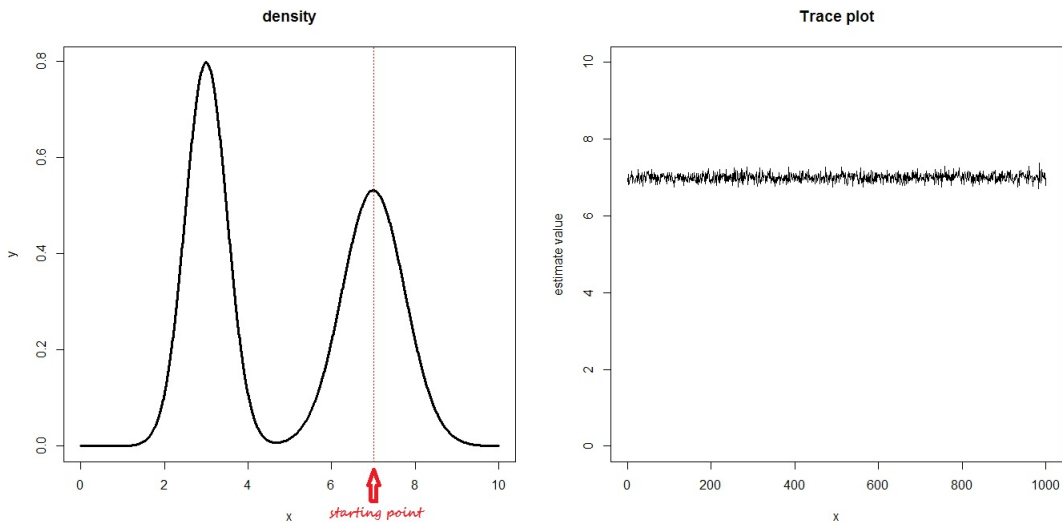
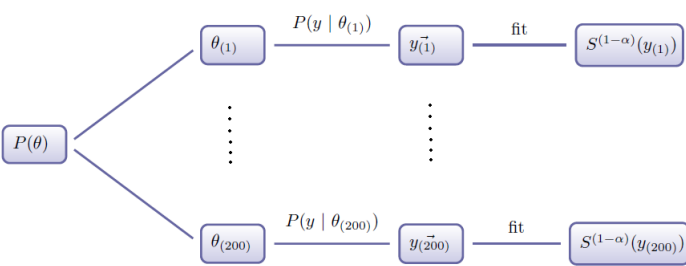


Figure 1: If our starting point is $x = 7$, then our Markov Chain will walk around $x = 7$ with beautiful trace plot. In this case, although we pass three tests above, we can not conclude our MCMC converges.

The other choice for checking convergence: Fortunately, we can use **simulation studies** to check everything is working.

Idea:

1. Simulate $\theta_{(j)}$ from the prior $P(\theta)$.
2. Simulate a dataset $y_{(j)}$ from the model $P(y | \theta_{(j)})$.
3. Sample from the posterior for data set $y_{(j)}$.
4. Find $100(1 - \alpha)\%$ central credible interval for θ from dataset $y_{(j)}$.



5. Record Yes/No whether the interval contained $\theta_{(j)}$.
6. Check roughly $100(1 - \alpha)\%$ of intervals contain their own specific $\theta_{(j)}$.

Why does it work?

$$\begin{aligned}\int P(\theta) \int I_{\{\theta \in S^{1-\alpha}(y)\}} P(\vec{y} | \theta) dy d\theta &= \int \int I_{\{\theta \in S^{1-\alpha}(y)\}} P(\theta) P(\vec{y} | \theta) dy d\theta \\ &= \int \int I_{\{\theta \in S^{1-\alpha}(y)\}} P(\vec{y}) P(\theta | \vec{y}) d\theta dy \\ &= \int P(\vec{y}) \left(\int I_{\{\theta \in S^{1-\alpha}(y)\}} P(\theta | \vec{y}) d\theta \right) dy \\ &= \int P(\vec{y}) (1 - \alpha) dy \\ &= 1 - \alpha\end{aligned}$$

$$\begin{cases} \text{Bayes:} & \int P(\theta) \int I_{\{\theta \in S^{1-\alpha}(y)\}} P(\vec{y} | \theta) dy d\theta = 1 - \alpha \\ \text{Frequentist:} & \int I_{\{\theta \in C^{1-\alpha}(y)\}} P(\vec{y} | \theta) dy = 1 - \alpha, \quad \forall \theta \end{cases}$$

Example

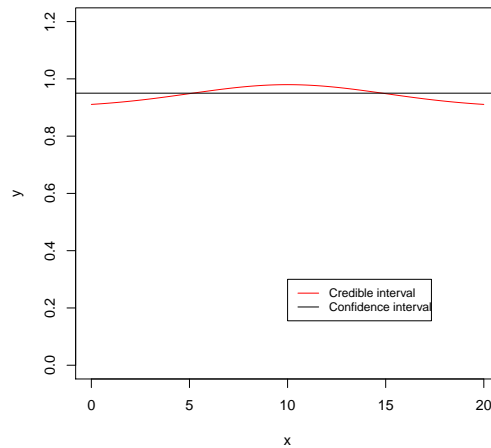


Figure 2: 95% credible interval v.s 95% C.I.

III. Posterior Predictive Checking

Validation simulation checks you can sample from the posterior under your model. It doesn't tell you if your model is a good fit to the data.

Idea:

Having fit your model to the data, you know roughly what θ is, so if you simulated from

$P(y | \theta)$, the simulated data should look "similar" to the real data.

Formally:

$$\begin{aligned}
 P(\tilde{y} | \vec{y}) &= \int P(\tilde{y}, \theta | \vec{y}) d\theta \\
 &= \int P(\tilde{y} | \theta) P(\theta | \vec{y}) d\theta
 \end{aligned}$$

Recipe:

- Sample $\theta^{(t)}$ from the posterior $P(\theta | \vec{y})$.
- For each $\theta^{(t)}$, sample a new dataset from $P(y | \theta^{(t)})$.
- We now have m predictive dataset and one real dataset.
- We can take univariate summary statistics of each dataset and the real dataset and compare. [**Note:** We could use min, mean, median, max.]

Example:

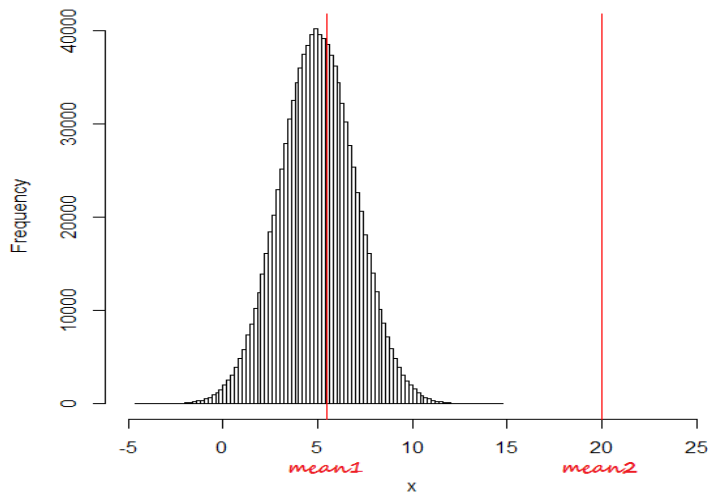


Figure 3: If the real mean equals mean1, then we will conclude we have a good fit. If the real mean equals mean2, then we will conclude it's not a good fit.

Posterior predictive p-value:

$$2 \min\{P(T(y^*) > T(\vec{y})), P(T(y^*) < T(\vec{y}))\}$$

Heuristically, if P is small, then your model is not a good fit to the data.