

STT 465

Gibbs Sampler

Gibbs Sampler

- ⇒ So far we have discussed models where the joint posterior distribution have a closed form.
- ⇒ However in most models, the joint posterior distribution does not have a closed form.
- ⇒ In those cases we can use Monte Carlo Markov Chain algorithms to draw samples from the joint posterior without explicitly drawing from that distribution.
- ⇒ Distributions we have discussed so far

<u>Prior</u>	$p(\theta_1, \theta_2)$
<u>Joint posterior</u>	$p(\theta_1, \theta_2 y)$
<u>Fully conditional distribution</u>	$p(\theta_1 y, \theta_2) ; p(\theta_2 y, \theta_1)$

- ⇒ In Gibbs sampler we draw samples iteratively from fully conditional distributions.

Gibbs Sampler

⇒ Target Posterior Density: $p(\theta_1, \theta_2, \dots, \theta_p \mid y)$

⇒ Fully Conditionals $p(\theta_j \mid y, ELSE) = p(\theta_j \mid y, \theta_{-j})$

⇒ Algorithm

- Initialize parameters (use values that have no-zero prior prob.)

- Iteratively sample from

$$\theta_1^s \sim p(\theta_1 \mid y, \theta_2^{s-1}, \dots, \theta_p^{s-1})$$

$$\theta_2^s \sim p(\theta_2 \mid y, \theta_1^s, \theta_3^{s-1}, \dots, \theta_p^{s-1})$$

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$$\theta_p^s \sim p(\theta_p \mid y, \theta_1^s, \theta_2^s, \dots, \theta_{p-1}^s)$$

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⇒ The algorithm generates a dependence sequence of samples of

$$\theta^s = [\theta_1^s, \theta_2^s, \dots, \theta_p^s]$$

⇒ After convergence these samples can be regarded as draws from the joint posterior density.

⇒ Given the way these samples were generated, the sequence is not an IID sequence.

⇒ Samples are not independent.

⇒ Markov Chain Property. However, the sequence has the property that

$$p(\theta^s, \theta^{s-1}, \theta^{s-2}, \dots, \theta^0 | y) = p(\theta^s | \theta^{s-1}, y) \times p(\theta^{s-1} | \theta^{s-2}, y) \times \dots \times p(\theta^1 | \theta^0, y)$$

⇒ Since θ^s can be regarded as a sample from the posterior distribution

⇒ $g(\theta^s)$ can be regarded as a sample from $p(g(\theta) | y)$

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⇒ Diagnostics

- ⇒ Some of the original samples may need to be discarded (burn-in)
- ⇒ Due to auto-correlation, we may need to thin (i.e., take x samples out of q)
- ⇒ Due to auto-correlation, MC errors cannot be computed using the formulas we used so far.

⇒ Diagnostics:

- ⇒ Trace plot
- ⇒ Auto-correlation between samples
- ⇒ Time-series MC error
- ⇒ Effective number of samples