

Sampling methods for Probit Regression and Application to GMRF

Mohammad Emtiyaz Khan

CS, UBC

Overview

- Probit Model
- Sampling methods : A&C (1993), H&H (2006)
- Comparison
- Modifications for GMRF

Probit Model

For $t = 1, \dots, T$, $y_t \in \{0, 1\}$, $\beta \in \mathcal{R}^d$,

$$(1) \quad Pr(y_t = 1|\beta) = \Phi(x_t\beta) \quad \text{where } \beta \sim \pi(\beta)$$

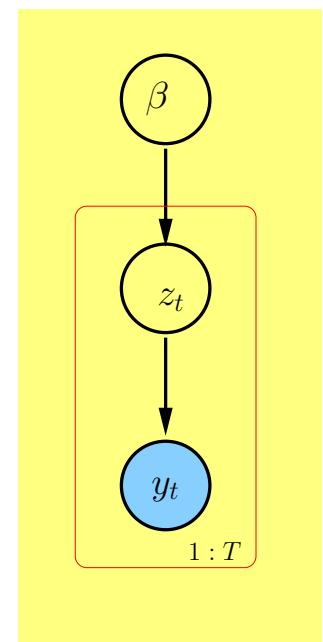
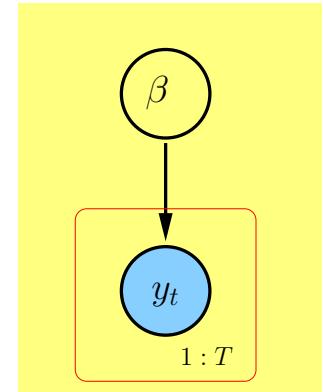
where $\Phi(u) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^u \exp(-u^2/2) du$.

Using Auxiliary variables, we can re-write:

$$(2) \quad y_t = \begin{cases} 1 & \text{if } z_t > 0 \\ 0 & \text{otherwise} \end{cases}$$

$$(3) \quad z_t = x_t\beta + \epsilon_t, \text{ where } \epsilon_t \sim \mathcal{N}(0, 1)$$

$$(4) \quad \beta \sim p(\beta)$$



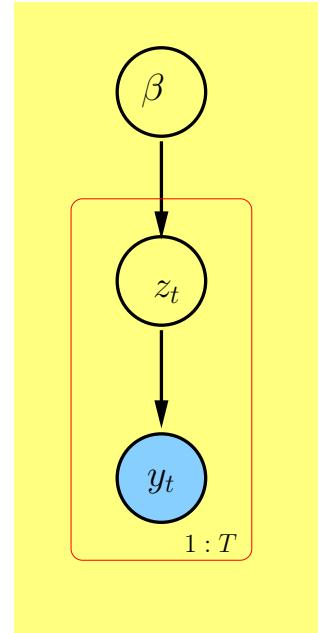
Gibbs Sampling I - Albert and Chib (1993)

If we assume prior $\beta \sim \mathcal{N}(b, v)$ and $X = [x'_1 x'_2 \dots x'_n]'$

$$(5) \quad p(\beta | z_{1:T}) \propto p(\beta) \prod_{t=1}^T p(z_t | \beta) = \mathcal{N}(\beta; B, V)$$

$$\begin{aligned} B &= V(v^{-1}b + X'z_{1:T}) \\ V &= (v^{-1} + X'X)^{-1} \end{aligned}$$

$$(6) \quad p(z_{1:T} | \beta, y_{1:T}) \propto \prod_{t=1}^T p(y_t | z_t) p(z_t | \beta) \\ = \begin{cases} \mathcal{N}(x_t \beta, 1) I(z_t \geq 0) & \text{if } y_t = 1 \\ \mathcal{N}(x_t \beta, 1) I(z_t < 0) & \text{otherwise} \end{cases}$$



Gibbs Sampling II - Holmes and Held (2006)

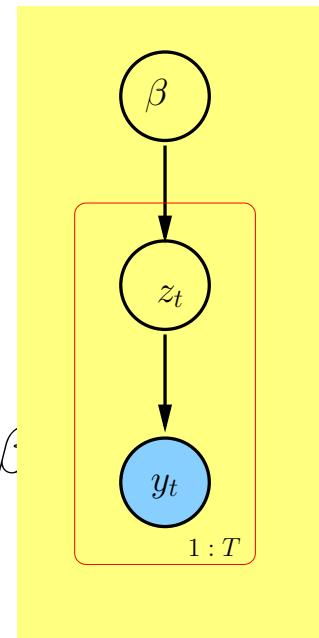
- An alternative - Update joint simultaneously:

$$p(\beta, z_{1:T} | y_{1:T}) = p(z_{1:T} | y_{1:T}) p(\beta | z_{1:T})$$

$$p(z_{1:T} | y_{1:T}) \propto p(y_{1:T} | z_{1:T}) \int_{\beta} p(z_{1:T} | \beta) p(\beta) d\beta$$

$$= p(y_{1:T} | z_{1:T}) f(z_{1:T}) \int_{\beta} p(\beta | z_{1:T}) d\beta$$

$$p(z_{1:T} | y_{1:T}) = I(z_{1:T}, y_{1:T}) \mathcal{N}(0, I_T + X V X')$$



- Sampling from a multivariate truncated Gaussian!
- However this can be simplified as described in HH(2006).

Our Work

- Implement A&C and H&H Gibbs sampler and compare.
- Modifications for GMRF.
- Modifications for polychotomous variables.
- Application to CGH microarray data.

Comparison: A&C and H&H

Mean of β

(189 Images (16x16), 300 iterations, $\beta \sim \mathcal{N}(0, I)$)

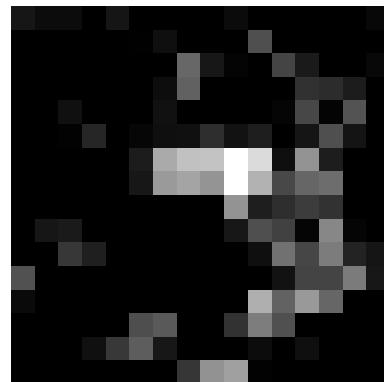
Template Images



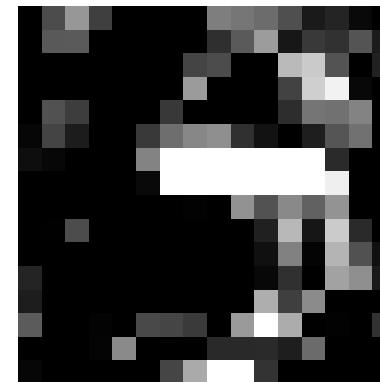
HH – MS code



AC – MTs code

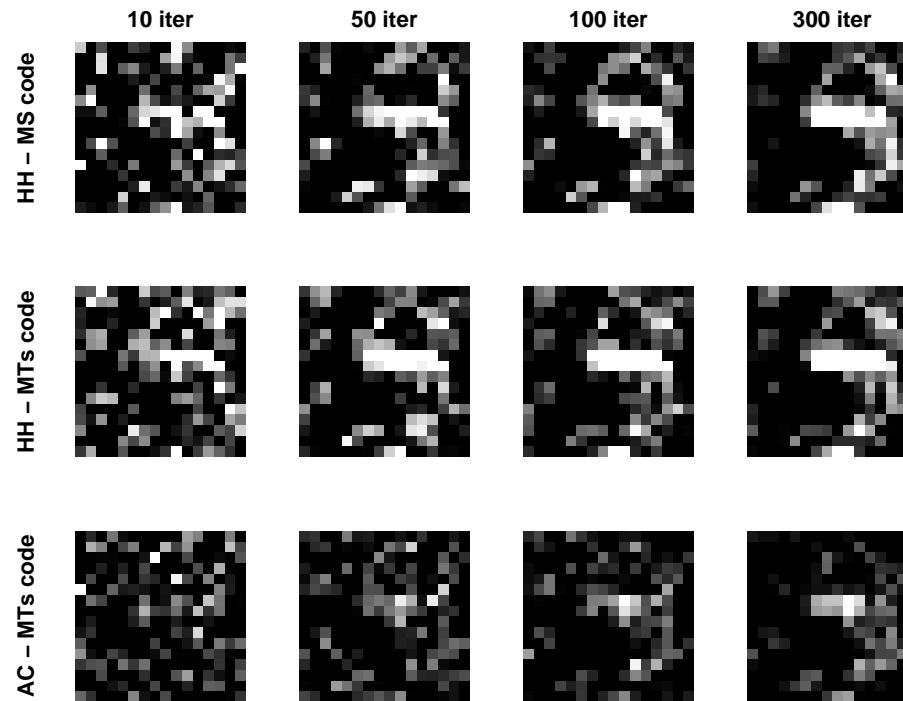


HH – MTs code

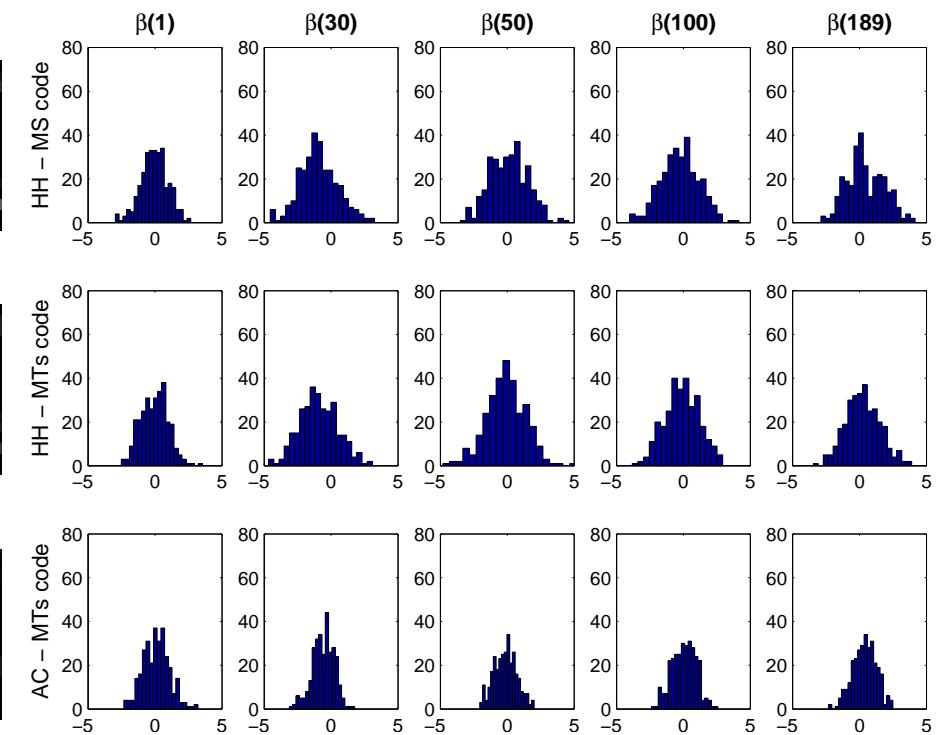


Comparison: A&C and H&H

Convergence



Distribution of β



Modification for GMRF

We extend these results to the case when β follows a Gaussian Markov Random Field (suggested by Kevin Murphy):

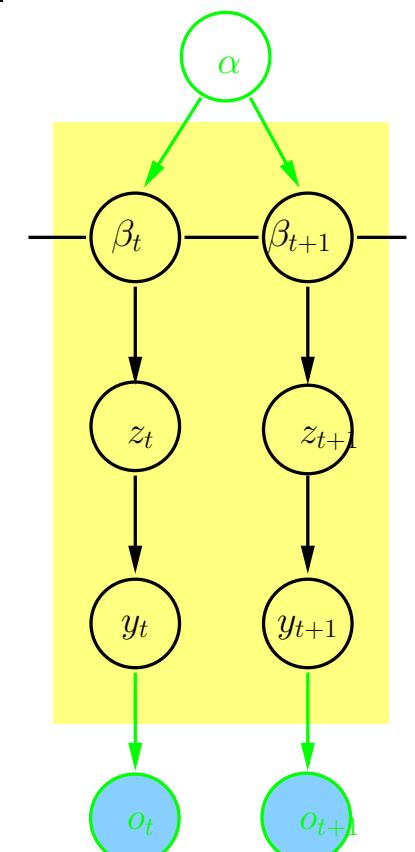
$$(7) \quad y_t = \begin{cases} 1 & \text{if } z_t > 0 \\ 0 & \text{otherwise} \end{cases}$$

$$(8) \quad z_t = \beta_t + \epsilon_t \quad \text{where } \epsilon_t \sim \mathcal{N}(0, 1)$$

$$(9) \quad \beta_{1:T} \sim \exp \left[-\frac{1}{2} \sum_{t=1}^{T-1} \frac{(\beta_t - \beta_{t-1})^2}{\alpha} \right]$$

Distribution of $\beta = \text{vec}(\beta_{1:T})$

$$(10) \quad \beta_{1:T} \propto \exp \left[-\frac{1}{2} \beta' W \beta \right]$$

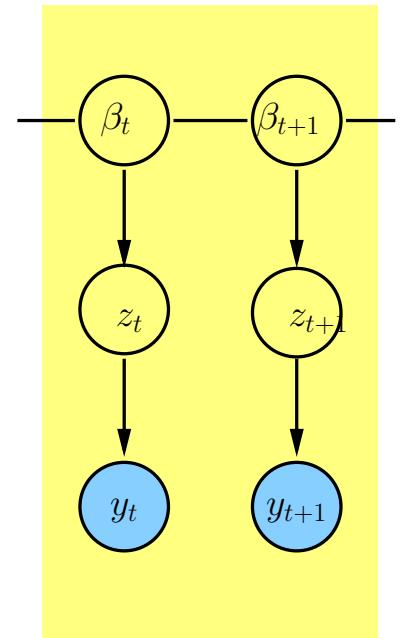


Modification for GMRF : Following A&C

If we assume prior $\beta_{1:T} \sim \mathcal{N}(0, v)$:

$$(11) \quad p(z_t | \beta_t, y_t) \propto \begin{cases} \mathcal{N}(x_t \beta_t, 1) I(z_t \geq 0) & \text{if } y_t = 1 \\ \mathcal{N}(x_t \beta_t, 1) I(z_t < 0) & \text{otherwise} \end{cases}$$

$$\begin{aligned} (12) \quad p(\beta_{1:T} | z_{1:T}) &\propto p(\beta_0) \prod_{t=1}^T p(z_t | \beta_t) p(\beta_t | \beta_{t-1}) \\ &= \mathcal{N}(\beta_{1:T}; B, V) \mathcal{N}(\beta_{1:T}; 0, W) \\ &= \mathcal{N}(\beta_{1:T}; B, V + W) \end{aligned}$$



Modification for GMRF : Following H&H

$$\begin{aligned} p(\beta_{1:T}, z_{1:T} | y_{1:T}) &= p(z_{1:T} | y_{1:T}) p(\beta_{1:T} | z_{1:T}) \\ p(z_{1:T} | y_{1:T}) &\propto p(y_{1:T} | z_{1:T}) \int_{\beta_{1:T}} p(z_{1:T} | \beta_{1:T}) p(\beta_{1:T}) d\beta_{1:T} \\ &= p(y_{1:T} | z_{1:T}) f(z_{1:T}) \int_{\beta_{1:T}} p(\beta_{1:T} | z_{1:T}) d\beta \\ p(z_{1:T} | y_{1:T}) &= I(z_{1:T}, y_{1:T}) \mathcal{N}(0, I_T + X V X') \end{aligned}$$

Summary

We presented

- A comparison two sampling methods for Probit models
- Suggested modifications for GMRF.

We wish to finish

- Modification for polychotomous variable model
- Application to CGH microarray data.

Acknowledgments: Thanks to Kevin Murphy, Mark Schmidt, and (of course) Arnaud.