

Markov random fields II

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Outline:

1. Auto-logistic (Ising) and auto-Poisson models
2. Estimation for Ising model
3. Bayesian Image analysis
4. Gibbs sampler (MCMC algorithm)
5. Phase-transition for Ising model

Brooks vs. Hammersley-Clifford

Given a set of (allegedly) full conditionals we can use either Brooks or H-C to identify candidate for a joint (unnormalized) density $p(\cdot)$. In both cases we need to check that $p(\cdot)$ can be normalized and that it is consistent with the given full conditionals.

On disadvantage of Brooks is that it in principle yields $n!$ solutions (possible non-uniqueness) and it does not inform on the form of $p(\cdot)$.

For H-C, we can construct the interaction functions using the full conditionals in a systematic way following the proof of 1. \Rightarrow 2. For given y these interaction functions and hence $p(\cdot)$ are uniquely determined by the full conditionals. Moreover, we can easily check that the constructed interaction functions are consistent with the full conditionals since

$$p_i(x_i|x_{-i}) \propto \frac{p_i(x_i|x_{-i})}{p_i(y_i|x_{-i})} = \frac{p(x)}{p(x_{-i}, y_i)} = \prod_{C:i \in C} \phi(x_C)$$

Gaussian MRF

Consider graph $G = (V, E)$ and full conditionals

$$p_i(x_i | x_{-i}) \propto \exp\left(-\frac{1}{2\kappa_i}(x_i - \mu_i + \sum_{l \sim i} \beta_{il}(x_l - \mu_l))^2\right)$$

where $\beta_{ij}/\kappa_i = \beta_{ji}/\kappa_j$.

Letting $y = (\mu_l)_{l \in V}$, we have following H-C that

$$\phi_\emptyset = p(\mu) \text{ (to be determined)} \quad \phi_i(x_i) = \exp\left(-\frac{1}{2\kappa_i}(x_i - \mu_i)^2\right)$$

$$\phi_{\{i,j\}}(x_i, x_j) = \exp\left(-\frac{\beta_{ij}}{\kappa_i}(x_i - \mu_i)(x_j - \mu_j)\right)$$

and $\phi_C(x_C) = 1$ for $\#C > 2$.

Note $\phi_{\{i,j\}}$ covers both pairs (i, j) and (j, i) .

How ?

Start with $\phi_\emptyset = p(\mu)$ (which we do not know yet).

By construction in proof of H-C:

$$\phi_i(x_i) = \frac{p(x_i, \mu_{-i})}{p(\mu)} = \frac{p_i(x_i|\mu_{-i})}{p_i(\mu_i|\mu_{-i})} = \exp\left(-\frac{1}{2\kappa_i}(x_i - \mu_i)^2\right)$$

Note that this implies $p(x_i, \mu_{-i}) = \phi_i(x_i)p(\mu)$.

Next,

$$\begin{aligned}\phi_{\{i,j\}}(x_i, x_j) &= \frac{p(x_i, x_j, \mu_{-\{i,j\}})}{p(\mu)\phi_i(x_i)\phi_j(x_j)} = \frac{p(x_i, x_j, \mu_{-\{i,j\}})}{p(x_i, \mu_{-i})\phi_j(x_j)} = \\ &= \frac{p_j(x_j|x_i, \mu_{-\{i,j\}})}{p_j(\mu_j|x_i, \mu_{-\{i,j\}})\phi_j(x_j)} = \exp\left(-\frac{\beta_{ji}}{\kappa_j}(x_i - \mu_i)(x_j - \mu_j)\right)\end{aligned}$$

Proceeding in the same way, we obtain $\phi_C(x_C) = 1$ for all C of cardinality $\#C > 2$ (of course we only need to consider C that are cliques with respect to G)

Considering the constructed $p(x)/p(\mu)$ and letting $Q_{ij} = \beta_{ij}/\kappa_i$ (with $\beta_{ii} = 1$) we see that this is the unnormalized density of $N(\mu, Q^{-1})$ provided Q is positive definite.

If Q is positive definite we can conclude

$$\phi_\emptyset = p(\mu) = (2\pi)^{-\#V/2} |Q|^{1/2}$$

For a Gaussian vector $X = (X_I)_{I \in V}$ we know that the full conditional of X_i only depends on those X_j for which $Q_{ij} \neq 0$.

Hence, X is a MRF with respect to $G \Leftrightarrow Q_{ij} = Q_{ji}$ differs from zero only if $\{i, j\} \in E$.

Note that the above is an example that ϕ_C can be equal to one also for C that is in fact a clique.

Auto-logistic model

Consider 2D rectangular $L \times K$ lattice V with horizontal/vertical neighbours. Only possible cliques are then singletons or pairs of horizontal or vertical neighbours.

Consider stochastic vector X on $\{0, 1\}^V$ with

$$p_i(x_i | x_{-i}) = \frac{\exp(\alpha x_i + \beta \sum_{j \in N_i} x_i x_j)}{1 + \exp(\alpha + \beta \sum_{j \in N_i} x_j)}$$

Note $p_i(1 | x_{-i})$ corresponds to logistic regression with covariate given by number of neighbouring 1's.

Following construction in proof of Hammersley-Clifford with $y = (0, \dots, 0)$ we obtain

$$\phi_i(x_i) = \exp(\alpha x_i) \quad \phi_{\{i,j\}}(x_{\{i,j\}}) = \exp(\beta x_i x_j)$$

We do not need to consider C with $\#C > 2$ since such a C can not be a clique.

Unknown normalizing constant

Hence joint density is

$$p(x) = p(0) \exp\left(\alpha \sum_{l \in V} x_l + \beta \sum_{\{i,j\} \in E} x_i x_j\right)$$

Sum defining

$$p(0) = \left[\sum_{x \in \{0,1\}^V} \exp\left(\alpha \sum_{l \in V} x_l + \beta \sum_{\{i,j\} \in E} x_i x_j\right) \right]^{-1}$$

has 2^{LK} terms !

Finite but in general impossible to compute exactly.

Hence we only know $p(\cdot)$ up to proportionality.

Boundary conditions

- ▶ free boundary: pixels at edges have only 2 or 3 neighbours
- ▶ fixed boundary: we condition on fixed values of boundary pixels. Then all interior “random” pixels have 4 neighbours
- ▶ toroidal (similar to circulant): edge pixels neighbours of pixels on opposite edge. E.g. pixel $(1, j)$ becomes neighbour of pixel (L, j) . Hence all pixels have 4 neighbours.

Symmetric case

Suppose all pixels have 4 neighbours (fixed or toroidal boundary).

If $\sum_{j \in \mathcal{N}_i} = 2$ we may want $p_i(0|x_{-i}) = p_i(1|x_{-i})$.

This is achieved with $\alpha = -2\beta$.

Ising model

Autologistic model is another name for the very famous Ising model (from statistical physics). In statistical physics 0, 1 are replaced by $-1, 1$ representing “spins” of elementary particles in piece of iron.

An equivalent form is

$$p(x) \propto \exp(\tilde{\alpha} \sum_{I \in V} x_I + \tilde{\beta} \sum_{\{i,j\} \in E} 1[x_i = x_j]) \quad (1)$$

That is, with $\tilde{\beta} > 0$, the model assigns large probabilities to x with many neighbours of equal value.

If $x_i \in \{0, 1\}$ and all pixels have four neighbours then (1) is equivalent to auto-logistic with $\alpha = \tilde{\alpha} - 4\tilde{\beta}$ and $\beta = 2\tilde{\beta}$.

Simulation of MRF

Gaussian MRF: use sparse matrix Cholesky decomposition of precision matrix.

General MRF: Markov chain Monte Carlo. Here we consider the so-called Gibbs sampler

Gibbs sampler

Idea: generate Markov chain X^1, X^2, \dots so X^n converges to the distribution p of X .

Reasonable requirement: p is invariant distribution for Markov chain. That is, if $X^i \sim p$ then also $X^{i+1} \sim p$. This is implied by reversibility:

$$P(X^i \in A, X^{i+1} \in B) = P(X^i \in B, X^{i+1} \in A) \quad \text{when } X^i \sim p$$

(set B equal to sample space S of X . Then reversibility implies $P(X^i \in A) = P(X^{i+1} \in A)$)

Gibbs sampler update: given $X^i = x^i$ pick I in V and let $X^{i+1} = (X_{-I}^i, Y_I)$ where Y_I is sampled from conditional distribution of $X_I | X_{-I} = x_{-I}^i$.

I can be chosen at random in V or we can run through V in a systematic order.

Gibbs update is reversible

Let $S = \prod_{I \in V} S_I$ be sample space of X .

$$P(X^i \in A, X^{i+1} \in B) = \int_S \int_{S_I} 1[(x_{-I}, y_I) \in B, x \in A] p_I(y_I | x_{-I}) dy_I p(x) dx$$

Moreover, using a change of variable,

$$\begin{aligned} P(X^i \in B, X^{i+1} \in A) &= \int_S \int_{S_I} 1[(x_{-I}, y_I) \in A, x \in B] p_I(y_I | x_{-I}) dy_I p(x) dx \\ &= \int_S \int_{S_I} 1[x \in A, (x_{-I}, y_I) \in B] p_I(x_I | x_{-I}) dx_I p(x_{-I}, y_I) dx_{-I} dy_I \end{aligned}$$

These two integrals are equal since

$$p(x) p_I(y_I | x_{-I}) = p(y_I, x_{-I}) p_I(x_I | x_{-I})$$

Under weak regularity conditions one can show that the Gibbs sampler Markov chain converges to $p(\cdot)$.

I.e. X^1, X^2, \dots serves as a random sample of (dependent) observations from $p(\cdot)$.

Estimation

Suppose we have observed realization of auto-logistic model.

Likelihood is

$$p(x; \alpha, \beta) = p(0; \alpha; \beta) \exp\left(\alpha \sum_{i \in V} x_i + \beta \sum_{i \sim j} x_i x_j\right)$$

Problem: normalizing constant

$$c(\alpha, \beta) = [p(0; \alpha, \beta)]^{-1} = \sum_{x \in V^{\{0,1\}}} \exp\left(\alpha \sum_{i \in V} x_i + \beta \sum_{i \sim j} x_i x_j\right)$$

can not be evaluated exactly and is difficult to approximate numerically.

Besag's pseudo-likelihood

Likelihood function for auto-logistic is intractable due to unknown normalizing constant

Julian Besag suggested to maximize the pseudo-likelihood (product of full conditionals)

$$PL(\alpha, \beta) = \prod_{i \in V} p_i(x_i | x_{-i}; \alpha, \beta)$$

Not likelihood except if X_i 's independent.

Score of log pseudo-likelihood is an unbiased estimating function

$$\mathbb{E} \frac{d}{d\alpha d\beta} \log p_i(X_i | X_{-i}; \alpha, \beta) = 0$$

(Bartlett identity) and one can show that PL estimates are asymptotically normal.

Computationally straightforward - formally equivalent to logistic regression.

Bayesian image analysis

Consider a pixel image $X = (X_i)_{i \in V}$ where X_i represents the color/intensity for pixel i .

Suppose we observe “dirty” image Y where

$$Y_i = X_i + \epsilon_i$$

where ϵ_i represents independent zero-mean noise terms with some density $\epsilon_i \sim f$.

We want to reconstruct X given observation y of Y !

Idea behind Bayesian image analysis: represent prior beliefs about X using a probability distribution and infer X using posterior distribution $X|Y = y$.

Pixel values continuous

Suppose $X_i \in \mathbb{R}$. We believe neighbouring pixel values are similar. We might model this using Gaussian MRF introduced in previous lecture. I.e. with $\mu_i = \mu$ and $\kappa_i = \kappa$,

$$X_i | X_{-i} = x_{-i} \sim N\left(\mu + \frac{1}{\#N_i + \tau} \sum_{k \sim i} (x_k - \mu), \frac{\kappa}{\#N_i + \tau}\right)$$

That is the conditional mean of X_i is essentially μ corrected with average deviations for neighbours. Joint density is of form

$$p(x) \propto \exp\left[-\frac{1}{2}(x - \mu)^T(Q + \tau I)(x - \mu)\right]$$

Assume $\epsilon_j \sim N(0, \sigma^2)$. Posterior is

$$p(x|y) \propto p(y|x)p(x) \propto \exp\left[-\frac{1}{2\sigma^2} \sum_{i \in V} (y_i - x_i)^2\right] p(x) \quad (2)$$

which is again a Gaussian MRF.

Posterior is known exactly (we can evaluate normalizing constant).

Note also: posterior Gaussian MRF is well-defined also with $\tau = 0$ in which case it does not depend on μ .

This is nice since we then do not need to specify μ .

Image segmentation

Image consists of two types (e.g. black or white) homogeneous regions. We may take $X_i \in \{0, 1\}$ with 0 for black and 1 for white.

Homogeneity: most neighbouring pixel values are of the same type
 \Rightarrow use Ising model as prior !

Assume again Gaussian noise. Then posterior is

$$p(x|y) \propto \exp\left(-\frac{1}{2\sigma^2} \sum_{i \in V} (y_i - x_i)^2 + \tilde{\alpha} \sum_{i \in V} x_i + \tilde{\beta} \sum_{i \sim j} 1[x_i = x_j]\right)$$

Again MRF distribution !

This time normalizing constant intractable but we can at least simulate posterior using Gibbs sampler.

We may want to use symmetric prior with $\tilde{\alpha} = 0$.

Contingency tables and graphical models

Consider a K -way contingency table given by combinations of K factors where the k th factor has values in set S_k .

For example 3 factors Smoker $S_1 = \{yes, no\}$, lung cancer $S_2 = \{yes, no\}$, Age $S_3 = \{young, middle, old\}$.

Consider an individual/object which is classified according to random values of these factors - leads to discrete random vector X that takes value $x = (x_1, \dots, x_K)$ if factor l takes the value x_l . E.g. outcome could be $(yes, no, middle)$ if person is middle-aged smoker without lung cancer.

Let

$$p(x) = P(X = x)$$

for x in sample space $S = \prod_{k=1}^K S_k$. E.g. $p(yes, no, middle)$ is probability of above outcome.

Suppose we have n individuals with vectors X_1, \dots, X_n . Let N_x denote the number of individuals with $X_i = x$.

We can model vector of numbers $N = (N_x)_{x \in S}$ of individuals for each combination x of factor levels using a multinomial distribution $N \sim \text{multinomial}(n, (p(x))_{x \in S})$.

Imposing a MRF structure on probability $p(x)$ allows us to study conditional independence properties of various factors. E.g. is smoking conditionally independent of lung cancer given age ? (OK, not true :))

Conditional independence structure can be visualized via accompanying graph where vertices represent factors.

Phase transition

Ernst Ising proposed his model as a model for ferromagnets. The spins represent orientations of iron-atoms. If majority of spins either $+$ or $-$ then the piece of iron is a magnet.

Consider the model with $\tilde{\alpha} = 0$ (no preference for either $+$ or $-$)

In one dimension, the Ising model is a Markov chain. According to the central limit theorem $M = \frac{1}{\sqrt{n}} \sum_{i \in V} x_i$ will converge to a zero mean normal distribution. I.e. distribution centered on configurations with roughly equal numbers of $+$ and $-$.

In two or more dimensions the picture is completely different. There exists a critical value $\tilde{\beta}_c \approx 0.88$ so that for $\tilde{\beta} < \tilde{\beta}_c$, the distribution of M is unimodal, while for $\tilde{\beta} > \tilde{\beta}_c$, the distribution is bi-modal ! I.e. either majority of $+$ or majority of $-$!

You can observe this by simulation: run a Gibbs sampler for large number of iterations starting from a random starting point (X_i^1 + or - with probability 0.5 each and initial spins independent).

For super critical $\tilde{\beta} > 0.88$ the Markov chain will end up in configurations dominated by either + or -. And once in a configuration with majority of + it takes a (very) long time to move to a configuration with a majority of - (and vice versa).

If β sub critical roughly equal amount of + and -

Exercises

1. Identify the ϕ_C functions for the auto-logistic model (following proof of the Hammersley-Clifford theorem, use $y = (0, 0, \dots, 0)$).
2. Use Brook's lemma to identify $p(\cdot)$ for the auto-logistic model. Does the result depend on the order of the factorization ?
3. Show that (1) is equivalent to the auto-logistic model in the case where all pixels have 4 neighbours

Hint: $1[x_i = x_j] = x_i x_j + (1 - x_i)(1 - x_j)$ when $x_i, x_j \in \{0, 1\}$.

4. Auto-Poisson: suppose $X_i | X_{-i} = x_{-i}$ is Poisson with mean $\exp(\alpha + \beta \sum_{j \in N_i} x_j)$ with neighbourhood structure as for the auto-logistic. Find the joint distribution of X . Show that it is well-defined when $\beta \leq 0$ (meaning $\sum_{x \in \mathcal{S}} h(x) < \infty$) but not ($\sum_{x \in \mathcal{S}} h(x) = \infty$) when $\beta > 0$ and $h(\cdot)$ denotes the unnormalized simultaneous density.

Exercises continued

5. How can you simulate a Gaussian MRF when the Cholesky decomposition $Q = LL^T$ has been obtained for the precision matrix ?
6. Show that the posterior distribution (2) is a Gaussian MRF. Also show that the posterior does not depend on μ when $\tau = 0$.

Hint: if $Z \sim N_n(\xi, K^{-1})$, then

$$p(z) \propto \exp\left(-\frac{1}{2}z^T K z + z^T K \xi\right).$$

7. Implement and run Gibbs sampler for the Ising model (1) with $x_i \in \{0, 1\}$. Use fixed boundary with all boundary pixels equal to 1. Consider the symmetric case $\tilde{\alpha} = 0$ and values of $\tilde{\beta} = 0.4, 0.7, 0.9$. What do you observe ? (some code available on webpage).

Exercises continued

8. 8.1 Show that the score function of pseudo-likelihood is unbiased.
- 8.2 Implement pseudo-likelihood for auto-logistic model when a fixed boundary condition is used (use R-procedure `glm`) (some code available on webpage).
- 8.3 Estimate α and β from the data set `isingdata.txt` using fixed boundary condition.

The data was generated from (1) with $\tilde{\alpha} = 0$ and $\tilde{\beta} = 0.4$. Do your estimates of α and β seem reasonable compared to this ?

9. The data set `imageAnoisy.txt` contains a binary (black/white) image corrupted by iid normal noise with mean zero and standard deviation 0.25. You can read and view it using

```
temp=as.matrix(read.table("imageAnoisy.txt"))
```

 and

```
image(temp)
```

. Adapt the previously constructed Gibbs sampler to sample from the posterior distribution when the Ising model is used as a prior. Use toroidal edge correction and try out different $\tilde{\beta}$ values.

Exercises continued

10. Consider the posterior distribution in exercise 6 with $\tau = 0$. Show that the posterior mean is

$$\hat{x} = \frac{1}{\sigma^2} (Q + \frac{1}{\sigma^2} I)^{-1} y$$

Compute the posterior mean based on the image data from previous exercise ($\sigma^2 = 0.25$). Try out varying values of κ .

Hint: use the sketch code `bayesian_GMRF.R`. Explain what is going on. Note moreover that $x = K^{-1}y \Leftrightarrow Kx = y$. If K is positive definite, $K = U^T U$ for an upper triangular U . Thus we can solve $Kx = y \Leftrightarrow U^T Ux = y$ in two steps involving first U^T and next U . Each step is computationally efficient because U and U^T are triangular matrices.