Markov Random Fields with Applications to M-reps Models

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Outline:

□ Background;

Definition and properties of MRF;

□ Computation;

□ MRF m-reps models.

Markov Random Fields

Model a large collection of random variables with complex dependency relationships among them.

Markov Random Fields

- A model based approach;
- Has been applied to a variety of problems:
 - Speech recognition
 - Natural language processing
 - Coding
 - Image analysis
 - Neural networks
 - Artificial intelligence

• Usually used within the Bayesian framework.

The Bayesian Paradigm

X = space of the unknown variables,
e.g. labels;
Y = space of data (observations),
e.g. intensity values;
Given an observation y∈Y, want to make inference about x∈X.

The Bayesian Paradigm

Prior P_X : probability distribution on X;Likelihood $P_{Y|X}$: conditional distribution of Ygiven X;

Statistical inference is based on the *posterior* distribution $P_{X|Y} \propto P_X \cdot P_{Y|X}$.

The Prior Distribution

- Describes our assumption or knowledge about the model;
- X is usually a high dimensional space. P_X describes the joint distribution of a large number of random variables;
- How do we define P_X?

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Assumptions

- $X = \{X_s\}_{s \in S}$, where each X_s is a random variable; *S* is an index set and is finite;
- There is a common state space *𝔅*: X_s∈*𝔅* for all s ∈ *𝔅*; | *𝔅* | is finite;
- Let $\Omega = \{ \omega = (x_{s_1}, ..., x_{s_N}) : x_{s_i} \in \mathcal{R}, 1 \le i \le N \}$ be the set of all possible *configurations*.

Dependency Graph

A simple undirected graph $\mathcal{G} = (S, \mathcal{N})$:

- *S* is the set of sites (vertices);
- N={N_s}_{s∈S} is the neighborhood structure (the set of edges). The neighbors of s are those sites that are connected to s by an edge;
- Let *C* denote the set of cliques completely connected subgraphs of *G*, including singletons.

Markov Random Field: Definition

P is an Markov random field on Ω with respect to $\mathcal{G} = (S, \mathcal{N})$ if (1) $P(X=\omega) > 0$ for all $\omega \in \Omega$; (2) $P(X_s=x_s | X_r=x_r, r \neq s)$ $= P(X_s=x_s | X_r=x_r, r \in \mathcal{N}_s)$ (local characteristics)

The local characteristics uniquely determines a joint distribution.

Examples of MRF: Nearest Neighbor Systems

• 1st order Markov chain $\{X_0, X_1, ..., X_n, ...\}$: $P(X_{n+1} = x_{n+1} | X_n = x_n, X_{n-1} = x_{n-1}, ..., X_0 = x_0)$ $= P(X_{n+1} = x_{n+1} | X_n = x_n)$

• 4-neighbor lattice system: $P(X_{i,j} | \text{ all other random variables}) = P(X_{i,j} | X_{i-1,,j}, X_{i+1,j}, X_{i,,j-1}, X_{i,,j+1})$

Gibbs Field

P is Gibbs on Ω with respect to $\mathcal{G} = (S, \mathcal{N})$ if $P(\omega) = 1/Z \cdot \exp\{-H(\omega) / T\},$ where

- Z is a normalizing constant (*partition function*);
- H is the *energy*. $H(\omega) = \sum_{C \in \mathcal{C}} U_C(\omega)$. \mathcal{C} is the set of cliques for \mathcal{G} . { $U_C \ge 0$ } are called the *potentials*;
- $U_{C}(\omega)$ depends only on those x_{s} of ω for which $s \in C$;
- T is a parameter (*temperature*).

The Hammersley-Clifford Theorem

P is an MRF with respect to **G** if and only if P is a Gibbs distribution with respect to **G**.

Advantage of Using the Gibbs Form

- The Gibbs form explicitly specifies the joint distribution;
- Local characteristics (conditional probabilities) can be easily formulated from the Gibbs form;
- The potentials can be learned from training data (see later slides) .

Examples: Nearest Neighbor Systems (cont.)

• 1-D :

$$H(\{x_i\}) = \sum U_i(x_i) + \sum U_{(i,i+1)}(x_i, x_{i+1})$$

• 2-D :

The most general form of the energy is $H(\{x_{i,j}\}) = \sum U_{\{(i,j)\}} (x_{i,j}) + \sum U_{\{(i,j), (i+1,j)\}} (x_{i,j}, x_{i+1,j}) + \sum U_{\{(i,j), (i,j+1)\}} (x_{i,j}, x_{i,j+1})$

Important Properties of MRF

• Markov property:

Let A, B, C \subset S. If every path from $a \in A$ to $c \in C$ meets some $b \in B$, then X_A and X_C are conditionally independent given X_B . *Can still model complicated dependencies!*

• Maximum entropy property:

The family $P_{\lambda}(\omega) = 1/Z_{\lambda} \exp\{-\Sigma_c \lambda_c U_c(\omega)\}$ are the maximum entropy models with fixed values for $E(U_c(\omega)) = U_c^*$ (average energy).

Learning by the ME Principle

- Choose a set of (local) features;
- Obtain empirical distribution of the features from training set;
- Learn the potentials by the ME principle.
- Example: ME distribution with specified mean and variance yields a Gaussian distribution.

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Computation Methods

• Dynamic programming

 the basic idea behind a lot of different algorithms, e.g. forward-backward, parsing, Viterbi, sum-product, belief propagation, etc.;

- relatively fast;
- does not work for all MRF's.
- Stochastic relaxation

General Computation Problems

- a) Sample from a Gibbs distribution;
- b) Find minimum energy;
- c) Compute expected values;
- d) Test model and estimate parameters.
- Among them, a) is the most basic problem.

Direct sampling from a Gibbs field $P(x) = Z^{-1} \exp(-H(x))$, $x \in X$, is usually not feasible because

- the underlying space X is huge;
- the partition function Z is intractable.

Stochastic Sampling Algorithms

Design a Markov chain with state space Ω whose equilibrium distribution is the desired Gibbs distribution.

Examples:

- Metropolis -Hastings algorithms: based on having "elementary" Markov chains;
- Gibbs sampler: based on using local characteristics.

Temperature in Gibbs Distribution

Any Gibbs field P can be put in a family $\{P_T\}$ with parameter T = temperature: $P_T(x) = 1/Z_T P(x)^{1/T}$ $= 1/Z_T \cdot \exp\{-E(x)/T\},$ $- \text{ as } T \to \infty, P_T \to \text{ uniform distribution;}$ $- \text{ as } T \to 0, P_T \to \delta_{\text{ mode}(P)}.$

Simulated Annealing

- Goal: find the global minimum energy (*ground state*), e.g. MAP estimates.
- Algorithm:
 - choose a cooling scheme $T_1 > T_2 > ... \rightarrow 0$;
 - generate a Markov chain $\{X^{(n)}\}\}$ on Ω where $X^{(n)} \rightarrow X^{(n+1)}$ is a transition by P^{Tn} ; the transition probabilities are specified by the Metropolis/Gibbs sampler;
 - If one cools at a *very slow* pace, then X⁽ⁿ⁾ converges in probability to the mode of P.

Simulated Annealing (cont.)

- Advantages:
 - guaranteed to find global minima (in principle), as opposed to greedy algorithms;
 - works for any Gibbs fields;
- Disadvantages:
 - convergence is very slow;
 - stopping rule is not clear;
 - hard to analyze;

Markov Chain Monte Carlo

- Goal: compute $E_{p}(f)$ for a function f on Ω .
- Traditional Monte Carlo: sample uniformly from Ω and average w.r.t. P $E_P(f(X)) \approx \sum_k f(X_k) P(X_k) / \sum_k P(X_k)$
- MCMC: sample from P and average uniformly



Summary of the Theory

- MRF provides a general framework for studying complex random systems;
- Computation is usually complicated;
- How can we do better?
 - data driven methods in computation;
 - better design of MRF, e.g. hierarchical MRF modes (HMF, HMRF, etc.);
 - other approximations, e.g. mean field, continuous stochastic processes.

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M-reps Models

- Multiscale shape models. Each scale k is described by a set of primitives {z^k_i};
- Object intrinsic coordinates provide correspondences among object population;
- Can easily describe both *global* and *local* variations, as well as inter-object relations.

MRF M-reps Models

- The probability distribution on the shape space is given by P({z^k_i});
- Markov assumption: $P(\mathbf{z}_{i}^{k} | \text{ all other primitives at all scales } \leq k)$

 $= P(\mathbf{z}_{i}^{k} | \mathcal{N}(\mathbf{z}_{i}^{k}), \mathcal{G}(\mathbf{z}_{i}^{k}))$

• If \mathbf{z}^k denotes scale k, then a multiscale MRF mreps model can be written as a Markov chain $P(\mathbf{z}^1, \dots, \mathbf{z}^n) = P(\mathbf{z}^1) \cdot P(\mathbf{z}^2 | \mathbf{z}^1) \cdots P(\mathbf{z}^n | \mathbf{z}^{n-1})$

MRF M-reps Models

- By the H-C theorem, the model has "two-sided" Markov property, i.e. $P(\mathbf{z}^k | \text{ all other scales}) = P(\mathbf{z}^k | \mathbf{z}^{k-1}, \mathbf{z}^{k+1})$, or equivalently, $P(\mathbf{z}^k_i | \text{ all other primitives at all scales})$
 - $= P(\mathbf{z}_{i}^{k} | \mathcal{N}(\mathbf{z}_{i}^{k}), \mathcal{P}(\mathbf{z}_{i}^{k}), \mathcal{P}(\mathbf{z}_{i}^{k}))$
- Use residues (differences) as features;
- The basic problem is how to specify the conditional probabilities $P_k = P(\mathbf{z}^k | \mathbf{z}^{k-1})$

The Boundary Level: MRF Model

- Primitives: $z_i = \tau_i$, the (normalized) displacement along the normal direction at point i;
- Neighborhood structure: nearest 4neighbors;
- The Gibbs distribution thus involves potentials of the form $A_i(\tau_i)$ and $B_{ij}(\tau_i, \tau_j)$, where i and j are 4-neighbors.

The Boundary Level: MRF Model

- Further assumptions:
 - Potentials have the same function form;
 - Gaussian (quadratic potentials);
- The joint distribution of $\{\tau_i\}$ has density

$$P(\{\tau_i\}) = \frac{1}{Z} \exp\left\{-\frac{1}{2\sigma_1^2} \sum_{i} s_i \tau_i^2 - \frac{1}{2\sigma_2^2} \sum_{\langle i,j \rangle} w_{ij} (\tau_i - \tau_j)^2\right\}$$

• σ_1 , σ_2 are parameters; $\{s_i\}$ and $\{w_{ij}\}$ are fixed from the previous stage.

The Boundary Level: Conditional Distribution

$$P(\{\tau_i\}) = \frac{1}{Z} \exp\left\{-\frac{1}{2\sigma_1^2} \sum_{i} s_i \tau_i^2 - \frac{1}{2\sigma_2^2} \sum_{\langle i,j \rangle} w_{ij} (\tau_i - \tau_j)^2\right\}$$

The log of the conditional probability density of τ_i is essentially

$$-\frac{s_{i}\tau_{i}^{2}}{2\sigma_{1}^{2}}-\frac{\sum_{j\in\mathcal{N}(i)}w_{ij}}{2\sigma_{2}^{2}}(\tau_{i}-\sum_{j\in\mathcal{N}(i)}\frac{w_{ij}}{\sum_{j\in\mathcal{N}(i)}w_{ij}}\tau_{j})^{2}$$

<u>Interpretation</u>: penalizes large τ_i and large deviation from "predicted τ " by neighbors.

The Boundary Level: Prior Model Learning

- The parameters σ₁, σ₂ can be learned from training data, using maximum likelihood estimates or other criteria;
- Other choices of model:
 - position-dependent parameters;
 - non-Gaussian models, maximum entropy learning.

The Atom Level

- Primitive: $z_i = A_i = (x_i, R_i, r_i)$, describing position **x**, local frame \mathcal{F} , and radius r of atom i. $z_i \in \mathbb{R}^3 \times SO(3) \times \mathbb{R}^+$;
- With 4-neighbor structure, the Gibbs distribution contains potentials of the form f_i (A_i) and g_{ij}(A_i, A_j) for neighboring atoms;
- Need a metric to describe difference between atoms ...

Atom Distance

Define a metric on atoms (or $\mathbf{R}^3 \times SO(3) \times \mathbf{R}^+$) by $d(\mathbf{A}_i, \mathbf{A}_j) = \sqrt{\alpha_E d_E^2(\mathbf{x}_i, \mathbf{x}_j) + \alpha_R d_R^2(\mathbf{R}_i, \mathbf{R}_j) + \alpha_r d_r^2(\mathbf{r}_i, \mathbf{r}_j)}$

where

- d_E is Euclidean distance in \mathbb{R}^3 ; - d_R is the Riemannian distance in SO(3); - d_r is the log-distance in \mathbb{R}^+ : $d(r_1, r_2) = |log(r_1/r_2)|$; - α_E , α_R , α_r are appropriate weights.

The Atom Level: MRF Model

 Let ΔA_i denote the "difference" between A_i and A'_i, where A'_i is the corresponding atom at the previous scale. In other words,

 $\Delta \mathbf{A}_{i} = (\Delta \mathbf{x}_{i}, \Delta \mathbf{R}_{i}, \Delta \mathbf{r}_{i}) = ((\mathbf{x}_{i} - \mathbf{x'}_{i}) / \mathbf{r'}_{i}, (\mathbf{R'}_{i})^{-1} \mathbf{R}_{i}, \mathbf{r}_{i} / \mathbf{r'}_{i}).$

• Prior model (quadratic potentials):

$$P(\{A_i\} | \{A'_i\}) = \frac{1}{Z} \exp\left\{-\sum_{i} \frac{s_i}{2\sigma_i^2} d^2(A_i, A'_i)\right\}$$

$$-\sum_{\langle i,j\rangle}\frac{W_{ij}}{2\sigma_{ij}^{2}}d^{2}(\Delta \mathbf{A}_{i},\Delta \mathbf{A}_{j})\}$$

The Atom Level: Conditional Distribution

 $P(\mathbf{A}_{i}) \propto \exp\left\{-\frac{\mathbf{s}_{i}}{2\sigma_{i}^{2}}\left[\alpha_{E} d_{E}^{2}(\Delta \mathbf{x}_{i}, 0) + \alpha_{R} d_{R}^{2}(\Delta \mathbf{R}_{i}, \mathbf{I}) + \alpha_{r} d_{r}^{2}(\Delta \mathbf{r}_{i}, 1)\right]\right\}$

 $-\sum_{j\in\mathcal{N}(i)}\frac{W_{ij}}{2\sigma_{ij}^{2}}\left[\alpha_{E}d_{E}^{2}(\Delta\mathbf{x}_{i},\Delta\mathbf{x}_{j})+\alpha_{R}d_{R}^{2}(\Delta\mathbf{R}_{i},\Delta\mathbf{R}_{j})+\alpha_{r}d_{r}^{2}(\Delta\mathbf{r}_{i},\Delta\mathbf{r}_{j})\right]\right\}$

- σ_i , σ_{ii} are trainable parameters of the model;
- The density is with respect to the Haar measure on the product space R³×SO(3)×R⁺;
- <u>Interpretation</u>: penalties on being away from "parent atom" and "neighbor mean";

MRF M-reps Models

- Similar MRF models can be designed for all other scale levels, using appropriate parent and neighbor terms;
- The full joint distribution is a probability measure on the shape space, with a relatively small number of parameters;
- The model is trainable (parametric vs. nonparametric).

References

- For Markov random fields:
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- For deformable m-reps models:
 - Joshi, Pizer, et al, TMI 2002;
 - Pizer, Joshi, et al, IJCV 2002.