Machine Learning and Graphical Models

(Lecture I)

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Outline

- A brief overview of Machine Learning
- Graphical Models
 - Representation
 - Inference
 - Learning



Definition of Machine Learning:

Learning from experiences.

"A computer program is said to learn from experience E with respect to some class of tasks T and performance measure P, if its performance at tasks in T, as measured by P, improves with experience E."



- Tom Mitchell

"Classical" Machine Learning Tasks:

- Classification: $f: \mathbb{R}^n \to \{-1,1\}$
 - spam filter, face recognition, ...
- Regression $f: \mathbb{R}^n \to \mathbb{R}$

$$f: \mathbf{R}^n \to \mathbf{R}$$

Hook's law, Kepler's law,...



Ranking

$$f: \mathbf{R}^n \to \mathbf{R}$$

- Search engine
- Probability (Distribution) Estimation

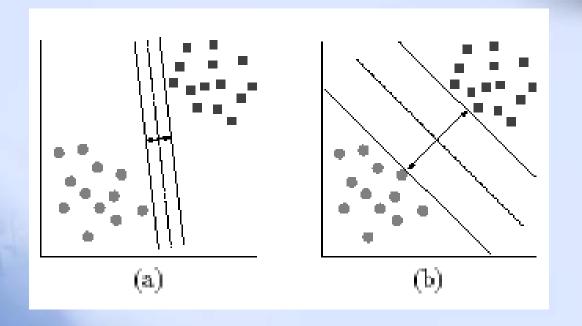
"Classical" Machine Learning Algorithms

- Classification
 - SVM
 - Boosting
 - Random Forest
 - Bagging
 - (Deep) Neural Networks
- Regression
 - Lasso
 - Boosting



Support Vector Machines (SVMs)

• SVM: the large l_2/l_2 margin classifier





SVM: hinge loss minimization + regularization

Boosting

- Boosting: (implicit) large l_1/l_{∞} margin classifier
- Boosting: exp loss minimization (+ regularization)



"Classical" Machine Learning Theories

VC theory

Capacity of the hypothesis space

- PAC-theory
- Margin theory
- Empirical Processes
- PAC-Bayes theory
- Regularization

Confidence

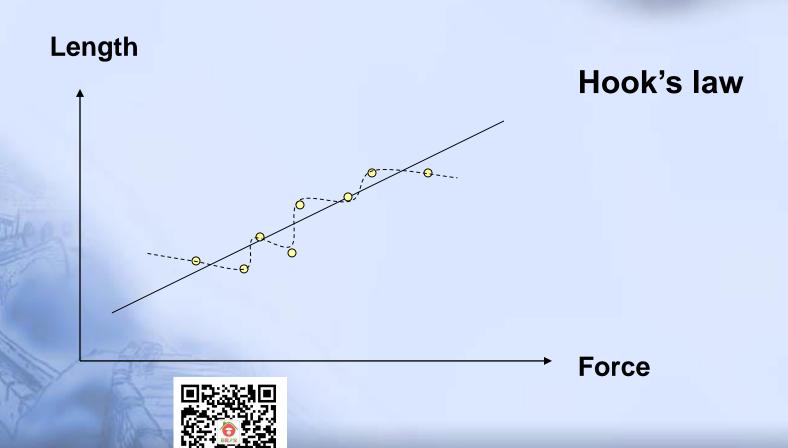
Capacity

PAC in Bayes framework

Capacity, smoothness



ML theories: Quantification of Occam's Razor



- Comparison of "Classical" Machine Learning Theories
 - Regularization:
 - Bayesian optimality
 - Only asymptotic (convergence, rate, non-uniform)
 - VC/PAC, Margin, PAC-Bayes,...
 - Relative optimality (optimal in a hypothesis space)
 - Non-asymptotic (finite sample bounds)



- Limitations of the "Classical" ML
 - Representation
 - Euclidean representation for input.
 - Simple representation for output.

How to represent STRUCTURES in data?



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Chapter I: Representation



Probabilistic Graphical Models: What and Why

- PGMs:
 - A model for joint probability distribution over random variables.
 - Represent dependencies and independencies between the random variables.
- Why is probability distribution important?
 - Genes and diseases, and everything
- Why PGM was invented by computer scientist, why not the statisticians?



Two types of PGMs

- Directed graph: Bayesian Networks (BNs).
- Undirected graph: Markov Random Fields (MRFs)



Bayesian Networks (BNs)



Example 1:

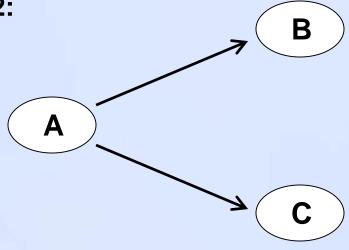


$$P(ABC) = P(A)P(B|A)P(C|B)$$

Given B, C and A are independent

Note: Dependency vs. Causality



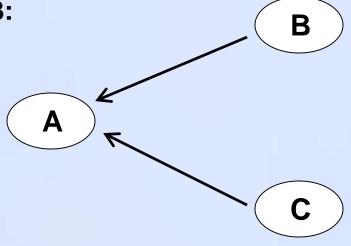




$$P(ABC) = P(A)P(B|A)P(C|A)$$

Given A, B and C are independent



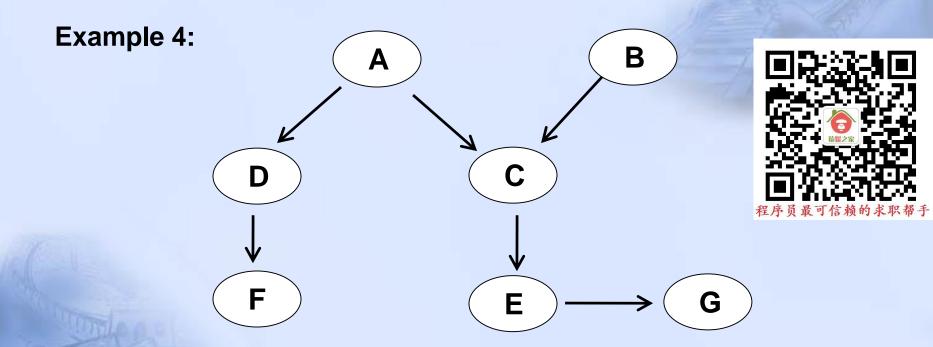




$$P(ABC) = P(B)P(C)P(A \mid BC)$$

B and C are independent;

But given A, B and C are NOT independent



P(ABCDEFG)

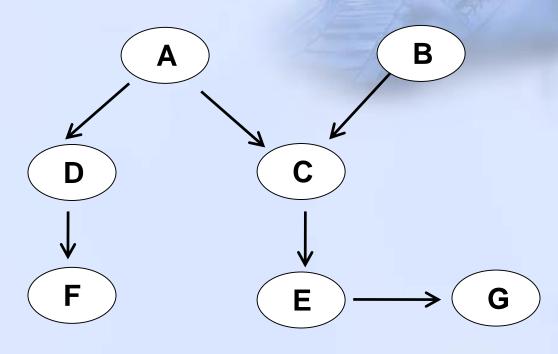
 $= P(A)P(B)P(C \mid AB)P(D \mid A)P(E \mid C)P(F \mid D)P(G \mid E)$

P(ABCDEFG)

= P(A)P(B)P(C | AB)P(D | A)P(E | C)P(F | D)P(G | E)

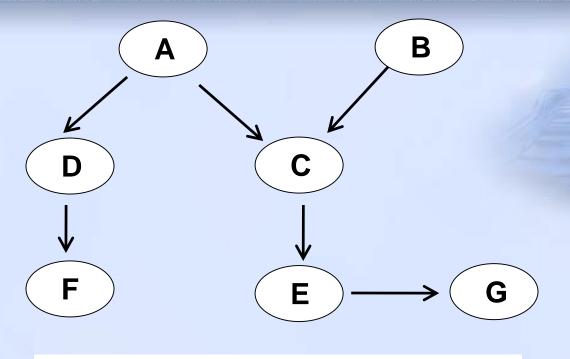
Learning:

 Find a factorization rule according to previous examples.



Factorization:

$$P(X_1, \dots, X_n) = \prod_i P(X_i | Pa(X_i))$$





$$P(X_1, \dots, X_n) = \prod_i P(X_i | Pa(X_i))$$

The graph must be acyclic!

BN must be DAG

Definition (Factorize according to a DAG):

A probability distribution P is said to be factorized according to a directed acyclic graph G if

$$P(X_1, \dots, X_n) = \prod_i P(X_i | Pa(X_i))$$

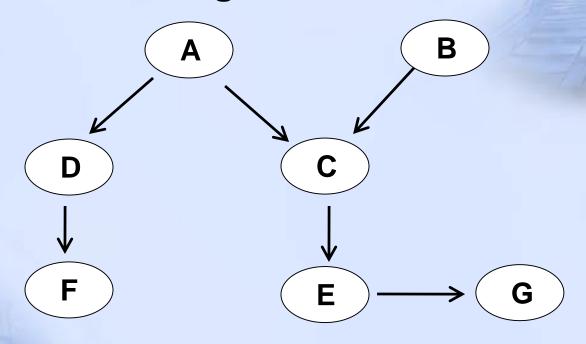


Definition (Bayesian Network):

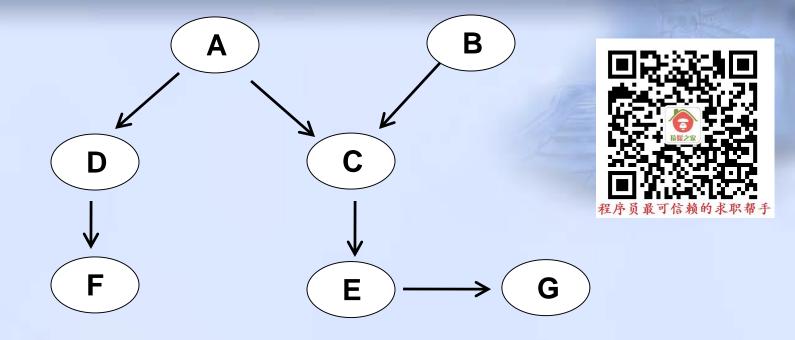
A Bayesian network is a pair (P, G) of a probability distribution and a DAG, where P is factorized according to G, and all the "local" conditional probability distributions are given.

$$P(X_1, \dots, X_n) = \prod_i P(X_i | Pa(X_i))$$

 Given the factorization, which variables are independent of C, given C's Parents A and B?



P(ABCDEFG)= $P(A)P(B)P(C \mid AB)P(D \mid A)P(E \mid C)P(F \mid D)P(G \mid E)$ D, F



$$P(ABCDEFG) = P(A)P(B)P(C \mid AB)P(D \mid A)P(E \mid C)P(F \mid D)P(G \mid E)$$

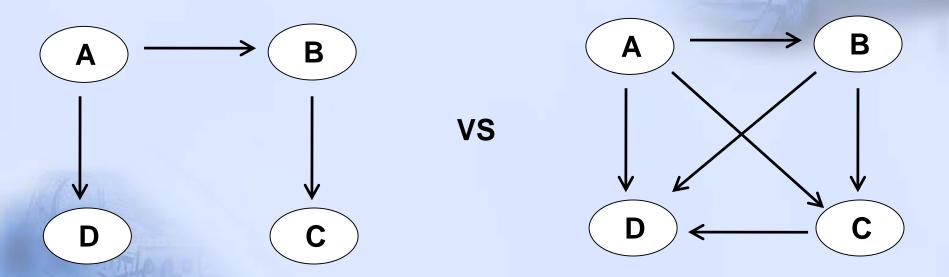
$$P(CD \mid AB) = P(C \mid AB)P(D \mid AB)$$

$$P(CF \mid AB) = P(C \mid AB)P(F \mid AB)$$

• Question: Let C be a node (random variable) in a BN. Which nodes (random variables) are independent of C, given C's Parents?

Theorem (Local Markov property for BN): For any node C (random variable) in a BN, all nodes that are not descendents of C are independent of C, given C's parents.

Sparse vs. Dense



- What is the joint pdf of the right BN?
- Is there any independence in the right BN?

• Question: Given a BN=(P, G), can you determine, for three arbitrary sets of random variables X={...}, Y={...}, and Z={...}, whether the following conditional independency hold?



Definition (active trail in BN)

Let x, y be two nodes and Z be a set of nodes. A path between x and y are said to be an active trial given Z, if the followings are true:

- 1) Let $x \Leftrightarrow ... \Leftrightarrow x' \to m \leftarrow y' \Leftrightarrow ... \Leftrightarrow y$ be the path, then m or one of its descendants is not in Z. That is, whenever there is a "**v-structure**" in the path, the middle node or one of its descendants is in Z;
- No other node along the path is in Z.

Definition (D-separation in BN)

Let X, Y, Z be three sets of nodes. X and Y are said to be D-separated by Z if for every node x in X and every y in Y, and every path between x and y, the path is not an active trial given Z.

Theorem (Informal)

The independencies in a BN are exactly those characterized by D-separation.

Theorem

For any BN (P,G), and arbitrary sets of nodes X, Y, Z. If X and Y are D-separated by Z in G, then

$$P(X,Y/Z) = P(X/Z) P(Y/Z)$$

Theorem

For any DAG G, and any sets of nodes X, Y, Z. If X and Y are not D-separated by Z in G, then there must exist a probability distribution P which factorize according to G, but

$$P(X,Y|Z) \neq P(X|Z) P(Y|Z)$$

The Representation Limit of BN

> Is there a BN that has precisely these independencies?

$$A \perp B \mid C, D;$$
 $C \perp D \mid A, B;$

Not every distribution can be represented by a BN satisfying exactly all the independencies!

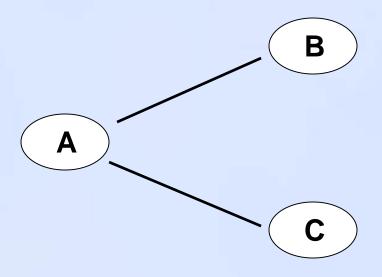
- To sum up:
 - BN represents joint probability distributions that can factorize according to:

$$P(X_1, \dots, X_n) = \prod_i P(X_i | Pa(X_i))$$

- The local independencies in BN are characterized by parents and non-descendants.
- The global independencies in BN are characterized by D-separation.



How MRFs Represent Joint pdfs:

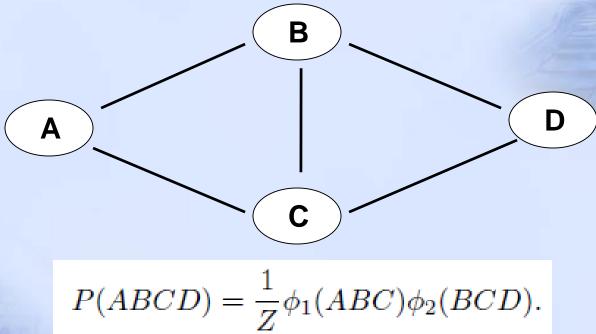


factors

$$P(ABC) = \frac{1}{Z}\phi_1(AB)\phi_2(AC).$$

Partition function

How MRFs Represent Joint pdfs:



- Factors correspond to maximal cliques.
- The joint distribution is the product of all factors normalized by the partition function.

Formal Definition (MRFs):

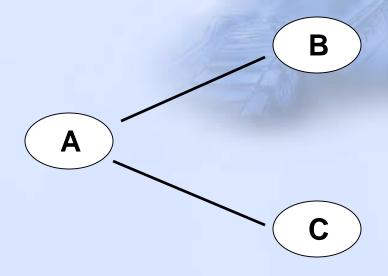
A Markov network is a pair (P, G), where G is an undirected graph and P factorizes according to G, i.e., P has the form

$$P(X_1,...,X_n) = \frac{1}{Z}\widetilde{P}(X_1,...,X_n) = \frac{1}{Z}\prod_i \phi_i(C_i)$$

where each C_i is a (maximal) clique in G.

The independence in MRF:

$$P(ABC) = \frac{1}{Z}\phi_1(AB)\phi_2(AC).$$

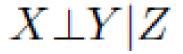


Easy to see:

$$B \perp C \mid A$$

$$P(BC|A) = P(B|A)P(C|A)$$

• Question: Given a MRF=(P, G), can you determine, for three arbitrary sets of random variables X={...}, Y={...}, and Z={...}, whether the following conditional independency hold?



Definition (Separation)

Let X, Y, Z be three sets of nodes in an undirected graph G. X and Y are said to be separated by Z if for every node x in X and every y in Y, and every path between x and y, there is a node in the path that belongs to Z.

Theorem (Informal)

All independencies in MRF are characterized by separation.

Theorem

For any MRF (P,G), and arbitrary sets of nodes X, Y, Z. If X and Y are separated by Z in G, then

$$P(X,Y/Z) = P(X/Z) P(Y/Z)$$

Theorem

For any undirected graph G, and any sets of nodes X, Y, Z. If X and Y are not separated by Z in G, then there must exist a probability distribution P which factorize according to G, but

$$P(X,Y|Z) \neq P(X|Z) P(Y|Z)$$

Machine Learning and Graphical Models

(Lecture II)

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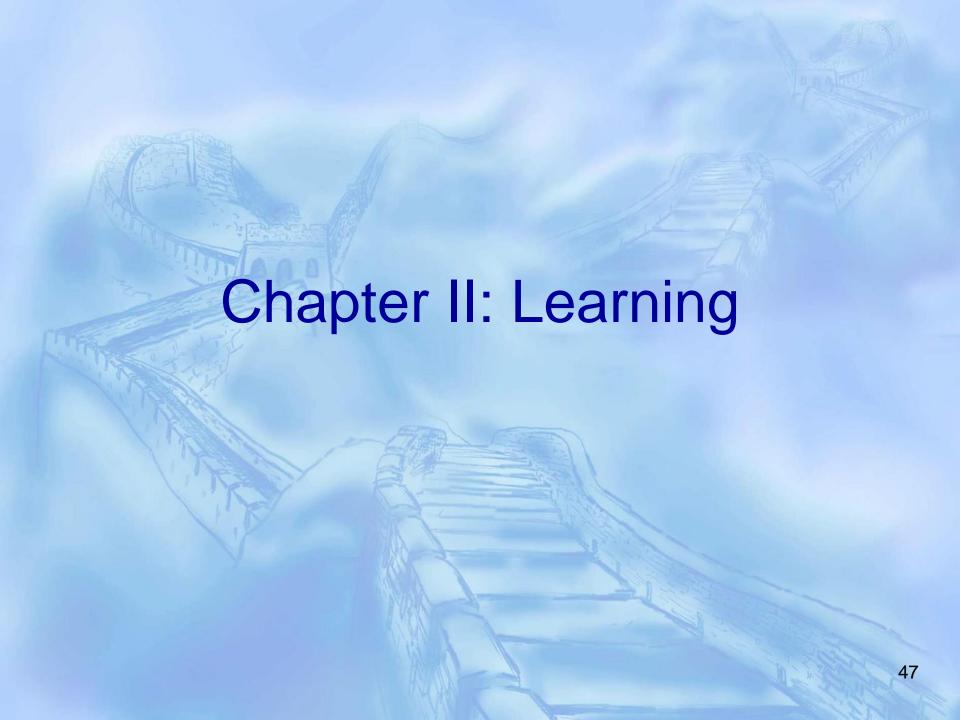
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- A brief overview of Machine Learning
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- Learning Graphical Models:
 - Definition of Bayesian Networks:

A Bayesian network is a pair (P, G) of a probability distribution and a DAG, where P is factorized according to G, and all the "local" conditional probability distributions are given.

$$P(X_1, \dots, X_n) = \prod_i P(X_i | Pa(X_i))$$

- Learning Bayesian Networks
 - BN = Structure (graph) + Local conditional distribution
 - Learning BN:
 - How to learn distribution's <u>parameters</u> from data
 - Relatively simple, standard parameter estimation.
 - How to learn structure from data
 - Very difficult! Why?

- Structure learning:
 - Structure (graph): edges between nodes.
 - Is the structure of a BN learnable?



 Note: the edge A —> C exists or not equals to whether the following equation holds strictly.

$$P(AC \mid B) = P(A \mid B)P(C \mid B)$$

Useful structure learning methods:

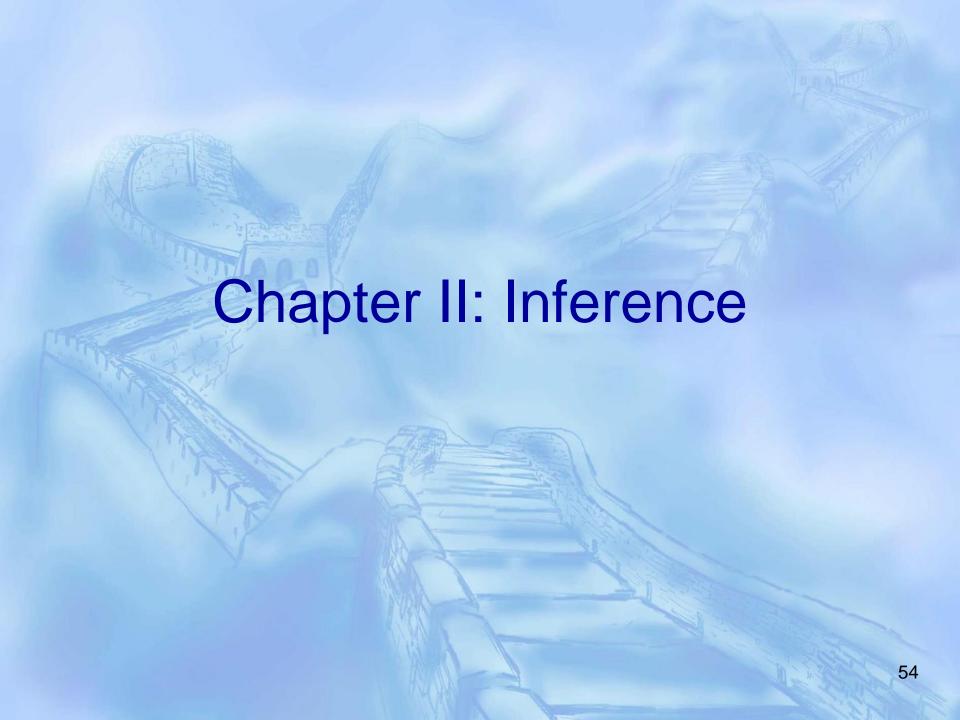
- Constraint-based structure learning.
 - Using hypothesis test to obtain independencies.
 - Construct the graph.
- Score-based structure learning: Penalizing "dense" graph
 - Likelihood scores
 - BIC
 - MDL
 - AIC

Open Problems

Robust Learning of Structures?

Outline

- A brief overview of Machine Learning
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- What is inference in GM
- The hardness of inference in GM
- Exact inference algorithms
- Approximate inference algorithms
- Future research directions



 Input: a graph and the local conditional distributions.

- Goal: two types of inference
 - Conditional probability

$$Pr(X = x | E = e)$$

MAP inference

$$\max_{x} \Pr(X = x \mid E = e)$$



- Exact inference in GM is hard
 - Decision version of exact inference is NPC.
 - Exact inference is #P complete.

- Approximate inference in GM is hard
 - \mathcal{E} -approximate inference is **NP**-hard for every \mathcal{E}

• Thm.1: Decide Pr(X = x) > 0 is **NP** complete. Proof: Reduction from 3SAT.

Thm.2: Compute Pr(X = x) is **#P** complete. Proof: Use above reduction from #3SAT. A Levin reduction, certificates are one-to-one.

Thm.3: For every $\varepsilon > 0$, compute an ε -approximate of $\Pr(X = x)$ is **NP**-hard.

Proof: \hat{p} is an ε -approximation of $\Pr(X = x)$ means that

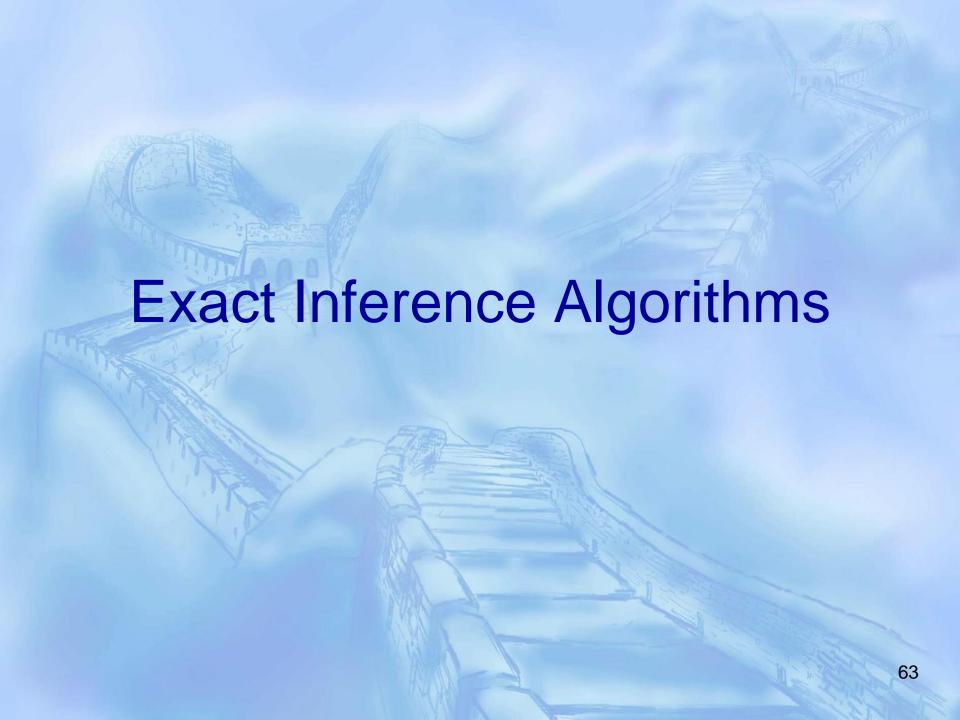
$$\frac{\Pr(X=x)}{1+\varepsilon} \le \hat{p} \le \Pr(X=x)(1+\varepsilon)$$

Clearly, if one has an ε -approximation of Pr(X = x), one can solve the **NP**C problem Pr(X = x) > 0.

Remark: For absolute approximate it's still NP-hard.

Thm.4: Exact and approximate MAP inference are all hard.

 Conclusion: The worst-case complexity of the inferences, both exact and approximate are NP-hard.



- The relation between BN and MRF
 - From BN to MRF (factors)

BN:
$$P(X_1,\ldots,X_n) = \prod_i P(X_i|Pa(X_i))$$

MRF:
$$P(X_1,...,X_n) = \frac{1}{Z} \widetilde{P}(X_1,...,X_n) = \frac{1}{Z} \prod_i \phi_i(C_i)$$

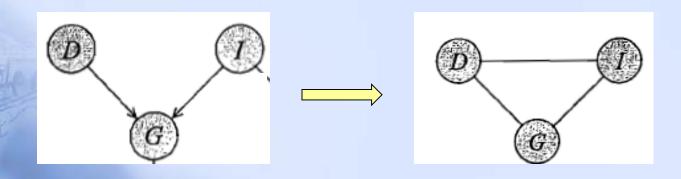
BN --> MRF:
$$\phi_i(X_i, Pa(X_i)) = P(X_i | Pa(X_i))$$

$$Z = 1$$

- From BN to MRF (graphs)
 - Moral graph (联姻图)

Delete the directions for edges;

Connecting the parents.



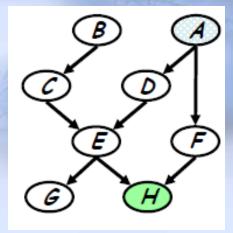
- Exact inference algorithms:
 - The variable elimination algorithm.
 - Belief propagation: the sum-product algorithm.
 - Belief update.

- Why study exact inference algorithms?
 - Gain intuition and insight for developing useful approximate algorithms.

The variable elimination algorithm

query:
$$Pr(A = a | H = h)$$

distribution:

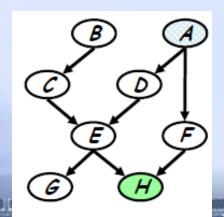


$$P(A)P(B)P(C \mid B)P(D \mid A)P(E \mid C,D)P(F \mid A)P(H \mid E,F)P(G \mid E)$$

solve:

$$= \sum_{b,c,d,e,f,g} P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(h|e,f)P(g|e)$$

Variable elimination----dynamic programming



$$\phi(a) \sum_{b} \phi(b) \sum_{c} \phi(c,b) \sum_{d} \phi(d,a) \sum_{e} \phi(e,c,d) \sum_{f} \phi(f,a) \phi(h,e,f) \sum_{g} \phi(g,e)$$

Step 1:
$$\delta(e) = \sum_{g} \phi(g, e) = 1$$
 (g,e)

Step 2:
$$\delta(a,e) = \sum_{f} \phi(f,a)\phi(h,e,f)(\delta(e))$$
 (a,e,f)

$$\phi(a) \sum_{b} \phi(b) \sum_{c} \phi(c,b) \sum_{d} \phi(d,a) \sum_{e} \phi(e,c,d) \delta(a,e)$$

$$\phi(a) \sum_{b} \phi(b) \sum_{c} \phi(c,b) \sum_{d} \phi(d,a) \sum_{e} \phi(e,c,d) \delta(a,e)$$

Step 3:

$$\delta(a,c,d) = \sum \phi(e,c,d)\delta(a,e)$$

(a,c,d,e)



$$\phi(a)\sum_{b}\phi(b)\sum_{c}\phi(c,b)\sum_{d}\phi(d,a)\delta(a,c,d)$$

Step 4:

$$\delta(a,c) = \sum_{d} \phi(d,a) \delta(a,c,d)$$

(a,c,d)

$$\phi(a)\sum_{b}\phi(b)\sum_{c}\phi(c,b)\delta(a,c)$$

$$\phi(a)\sum_{b}\phi(b)\sum_{c}\phi(c,b)\delta(a,c)$$

$$\delta(a,b) = \sum_{c} \phi(c,b) \delta(a,c)$$

(a,b,c)

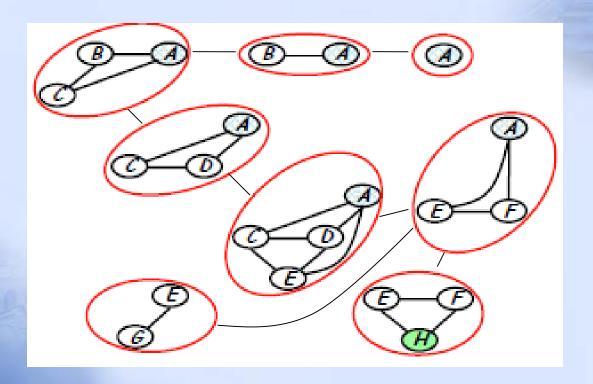
$$\phi(a)\sum_{b}\phi(b)\delta(a,b)$$

$$\delta(a) = \sum_{b} \phi(b) \delta(a,b)$$

(a,b)

$$\phi(a)\delta(a)$$

- Message Passing: the SP algorithm
 - Variable elimination induced clique tree



Family preserving property: factors scope

Running intersection property!

Variable elimination as message passing on clique tree:

• General message passing from clique C_i to C_j

$$\delta_{i \to j} = \sum_{C_i \setminus S_{ij}} \psi_i \prod_{k \in Nb(i) \setminus \{j\}} \delta_{k \to i}$$

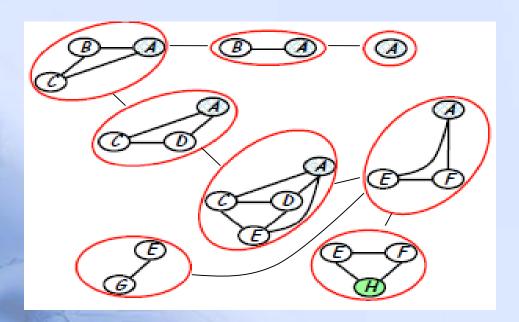
$$S_{ij} = C_i \cap C_j$$

 ψ_i : original terms (potential) of C_i

- The Sum-Product belief propagation algorithm:
 - Construct a clique tree that satisfies the family preserving and running intersection properties.
 - Choose clique where the query variable lies in as the root.
 - Message passing from the leaves.
 - After all messages arrive at the root, sum over all other variables in the root clique other than the query variable.

Thm: The Sum-Product algorithm always gives the correct answer for any clique tree!

- What if we want to compute k queries?
 - Message passing k times?
 - No! Message can be reused.
 - For each edge in the clique tree, twice is enough, one for each direction (up-down)



 After message passing on all edges in two directions, each clique has a belief (joint probability).

• Belief:
$$\beta_i = \psi_i \prod_{k \in Nb(i)} \delta_{k \to i}$$

The system must satisfy the calibration property:

$$\sum_{C_i \setminus S_{ij}} \beta_i = \sum_{C_j \setminus S_{ij}} \beta_j$$

Why calibration holds?

$$\beta_i = \widetilde{P}(C_i) \propto P(C_i)$$

Beliefs are marginal probabilities on the cliques!

$$\sum_{C_i \setminus S_{ij}} \beta_i = \sum_{C_j \setminus S_{ij}} \beta_j = \widetilde{P}(S_{ij}) \propto P(S_{ij})$$

Agree on the marginal probability of S_{ij} !

- Belief update algorithm:
 - Message passing:

$$\delta_{i \to j} = \sum_{C_i \setminus S_{ij}} \psi_i \prod_{k \in Nb(i) \setminus \{j\}} \delta_{k \to i}$$

• Belief: $\beta_i = \psi_i \prod_{k \in Nb(i)} \delta_{k \to i}$

• So
$$\delta_{i o j} = rac{\displaystyle\sum_{C_i \setminus S_{ij}} eta_i}{\delta_{j o i}}$$

Propagating beliefs from C_i to C_j

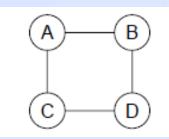
Belief update algorithm:

- Construct clique tree
- Initialize: $\beta_i = \psi_i$ $\mu_{i \to j} = 1$
- Update: $t_{i \to j} \leftarrow \sum_{C_i \setminus S_{ij}} \beta_i$

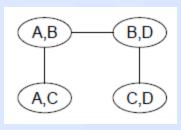
$$\beta_j \leftarrow \frac{\beta_j t_{i \to j}}{\mu_{ij}}$$

$$\mu_{ij} \leftarrow t_{i \rightarrow j}$$

- How to construct clique tree?
 - Graph:

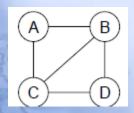


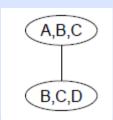
Clique tree?



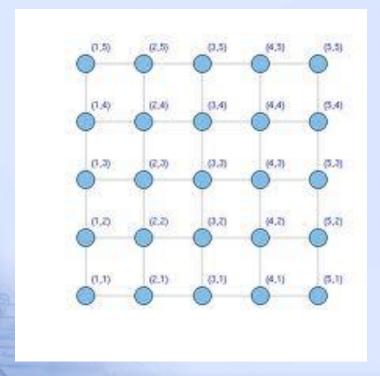
No!

- The graph cannot have a 4-loop (or larger)!
- Solution: triangulation and chordal graph.



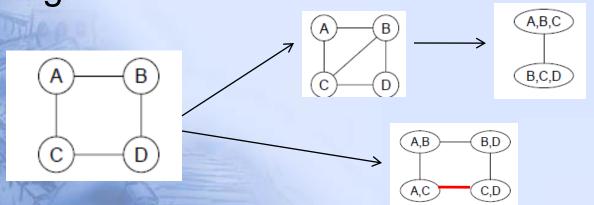


What if the graph is an Ising model,
 Triangulation and chordal graph induce clique trees that have large width and the computational complexity is very high.



- Construction of Clique Trees:
 - Variable Elimination;
 - BN to Moral graph, then to chordal graph, then find all the maximum cliques, and then use the maximum spanning tree algorithm to construct the edges of the clique tree.

- Reflection of the Sum-Product Belief Propagation algorithm:
 - The algorithm itself does require the clique tree has to be a "tree".
 - What if we construct a general "clique graph" instead of a "clique tree" and run the SP algorithm?

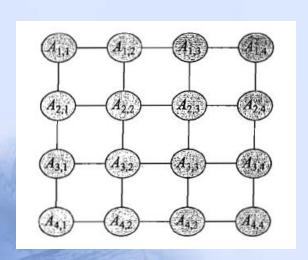


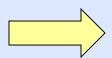
Clique tree

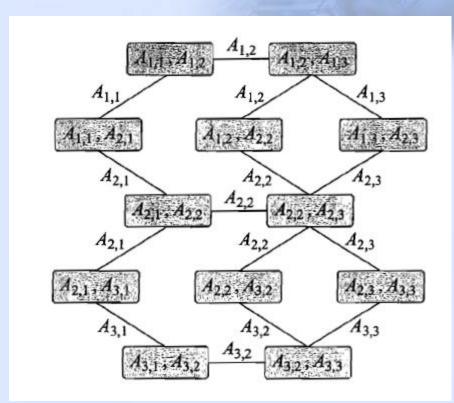
Cluster graph

Ising Model

Cluster Graph







 What's the result if run Sum-Product belief propagation on the cluster graph?

Loops?

Convergence?

Correctness?

This is the "Loopy Belief Propagation" algorithm!

Machine Learning and Graphical Models

(Lecture III)

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- What is inference in GM
- The hardness of inference in GM
- Exact inference algorithms
- Approximate inference algorithms
- Future research directions

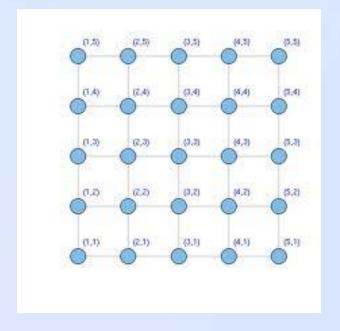
Approximate inference methods:

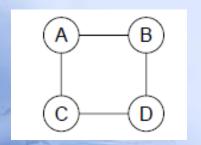
- Variational Methods.
 - (Loopy) belief propagation.
 - Mean field method.

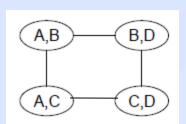
- Sampling based methods.
 - Metropolis-Hastings
 - Gibbs Sampling
 - Importance Sampling

Variational Methods

• (Loopy) belief propagation:







Cluster graph

- Loopy belief propagation: just let the belief propagates.
- On clique trees, belief propagation converges after propagating on all edges (two directions).
- For general cluster graphs, it is not guaranteed to converge. Even if it converges, it can converge to a wrong answer.

- Variational explanation of the (loopy) belief propagation:
 - Exact inference

Proposition: The following optimization problem gives the same result as exact inference:

Find a set of beliefs to minimize the Helmholtz free energy (or equivalently the relative entropy) to the factored distribution on clique trees under the constraints of calibration (local agreement).

 The fixed point characterization of the solution of the optimization problem:

$$\delta_{i \to j} = \sum_{C_i \setminus S_{ij}} \psi_i \prod_{k \in Nb(i) \setminus \{j\}} \delta_{k \to i}$$

Approximate inference

Find a set of beliefs to minimize the factored free energy to the factored distribution on cluster graphs under the constraints of calibration.

 The fixed point characterization of the solution of this optimization problem is the (loopy) belief propagation formula.

- Mean field method:
 - Using first order approximation of the target distribution:

$$P(X_1, \dots, X_n) = \prod_i P(X_i)$$

 The fixed point equation and updating rule have simple forms.

Sampling Based Methods

- Sampling-based (Monte Carlo) algorithms:
 - Idea: Using sample based frequency to approximate the probability.
 - Probability is Expectation:

$$P(X = x) = E[I(X = x)]$$

 Expectation (of a function) can be approximated by sample average:

$$E[f(X)] \approx \frac{1}{n} \sum_{i} f(X_i)$$

$$X, X_1, \dots, X_n$$
 i.i.d. ~ P

- Monte Carlo is useful in many problems:
 - High dimensional numerical integration.
- How to generate the sample when the target probability distribution is difficult to compute?
- Markov Chain Monte Carlo (MCMC)
 - Key idea: Generate data by Markov chain, whose stationary distribution is the target pdf.

- A brief review of finite state Markov chains:
 - Reducible vs. Irreducible
 - Periodic vs. Aperiodic
 - Ergodic: no matter which initial state is, the process will converge to a unique stationary distribution.

• Regular MC: 1) for every pair of states x, x', the prob. that x will reach x' in k steps for some finite k is positive; 2) for every state x, there is positive prob. that x will stay in x in the next step.

Easy to understand via the graph view

Theorem (sufficient condition for ergodic MC):
 If a finite state MC is regular, then it is ergodic.

- Regular Markov chains are good for our purpose.
- We run the Markov chain, and wait for it converges to the stationary distribution, then the data can be used for approximate calculation.
- But, how long will it take for the MC to converge?



Gibbs Sampling

- One of the simplest sampling algorithm.
- Assume for $X = (X^1, \dots, X^d)$, $P(X^j | X^{-j})$ is easy to sample.
- Easy to implement for Graphical Models.
- Proposed by Geman & Geman (TPAMI, 1984).

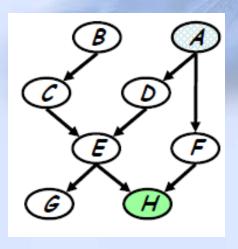
Gibbs sampling algorithm

- Goal: draw a sequence of examples x_1, x_2, \dots, x_n , when $n \to \infty$, $x_n \sim P$, where P is the target distribution; $x_i = (x_i^1, \dots, x_i^d) \in R^d$
- Algorithm:
 - Draw from some initial distribution $x_0 \sim P_0$
 - For $t = 1, \dots, n$
 - 1. $X_t \leftarrow X_{t-1}$
 - 2. For each $j \in [d]$
 - Sample X_t^j according to $P(X^j | X^{-j})$
 - Return x_1, x_2, \dots, x_n

Why Gibbs sampling is easy for PGM?

• Sample P(F | a,...e, g, h)

$$P(F | a,...e, g, h) = \frac{P(F, a,...e, g, h)}{\sum_{f} P(f, a,...e, g, h)}$$



$$= \frac{P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(F|a)P(g|e)P(h|e,F)}{\sum_{f} P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(g|e)P(h|e,f)}$$

$$= \frac{P(F|a)P(h|e,F)}{\sum_{f} P(f|a)P(h|e,f)}$$

Only the factors that involve F left!

- Generally, for both Bayesian networks and Markov networks, the conditional probability for Gibbs sampling involves only factors that the query random variable lives in.
- Trivially generalize to the case where there is evidence
 - Draw a sequence of examples where the target distribution is $P(\cdot | E = e)$

Theorem:

The Gibbs sampling process has a unique stationary distribution P (or $P(\cdot | E = e)$)

- Disadvantages of Gibbs sampling for PGMs:
 - Slow convergence to stationary distribution.



- For Gibbs sampling, we assume that it is easy to generate a sample from $P(X^{j}|X^{-j})$. But sometimes, this is difficult.
- More generally, for a target distribution P, it may be very difficult to generate sample directly according to P, does MCMC help?
- The idea of Metropolis-Hastings:
 - Using a *proposal* distribution $\widetilde{T}(x'|x)$: a transition model.

- An important result for Markov chain:
 - Detailed Balance (Reversible Markov chain):

Definition: A finite state MC with transition probability matrix T is said to be *reversible* if there exists a unique distribution P such that for all x, x'

$$P(x)T(x'|x) = P(x')T(x|x').$$

The above equation is called detailed balance.

• Reversible: for any sequence x_1, x_2, \dots, x_n , the probability that it occurs in the process is the same as the probability that x_n, x_{n-1}, \dots, x_1 occurs.

Theorem:

If the transition matrix T defines a regular Markov chain, and T satisfies the detailed balance w.r.p. to P, then P is the unique stationary distribution of the Markov chain T.

- Metropolis-Hastings algorithm:
 - Goal: draw a sequence of examples x_1, x_2, \dots, x_n , when $n \to \infty$, $x_n \sim P$, where P is the target distribution.
 - Algorithm:
 - Let T(x'|x) be a proposal transition model.
 - Define the transition matrix of a Markov chain as:

$$T(x'|x) = \min \left[1, \frac{P(x')\widetilde{T}(x|x')}{P(x)\widetilde{T}(x'|x)} \right]$$

Generate $x_1, x_2 \cdots, x_n$ according to the MC of T.

Proposition:

For any target distribution P, and any proposal transition model $\widetilde{T}(x'|x)$, the Markov chain defined by T in the Metropolis-Hastings algorithm satisfies the detailed balance w.r.p. P.

Thus if the Markov chain defined by T is regular, P is the unique stationary distribution.



Mixing time for MCMC:

- What we need is the stationary distribution of the Markov chain, but how long does it take to converge --- mixing time (burn in).
- Gibbs sampling sometimes has very slow convergence.
- Metropolis-Hastings's convergence depends on the proposal distribution.

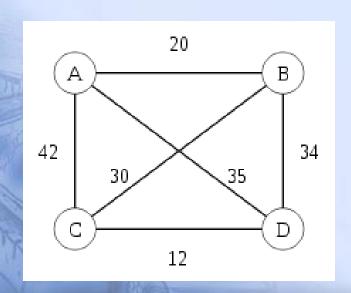
- Theory for the convergence for MCMC:
 - For a Markov chain, the largest eigenvalue of the transition matrix T is 1; the **gap** between the largest and the second largest (in absolute value) eigenvalue determines the mixing time.
- A main challenge for PGM:
 - Design MCMC algorithms that: 1) efficiently implementable for PGMs; 2) mixing time is not too long.

Some Thoughts and Open Problems



- Inference is NP-hard, what shall we do?
 - Develop practical algorithms. #P complete is a worst case result.

 To solve the inference problem, we are in a situation very similar to solving TSP (NP-hard):



TSP: Find the shortest path such that each vertex is visited exactly once.

- TSP (decision) is NP-complete.
- Euclidean TSP is NP-hard.
- Approximate TSP is NP-hard.

Arora proved (Gödel prize, 2010):

Euclidean TSP + approximation

polynomial approximation scheme



Can we find a reasonable class of graphical models such that (approximate) inference has polynomial time algorithm?



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Thanks!



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