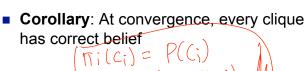


Convergence of Lauritzen-Spiegelhalter Algorithm

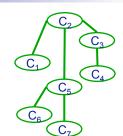


- Complexity: Linear in # cliques
 - for the "right" schedule over edges (leaves to root, then root to leaves)



$$\pi_i(c_i) = P(c_i)$$

$$\mu_{ij}(s_{ij}) = P(s_{ij})$$



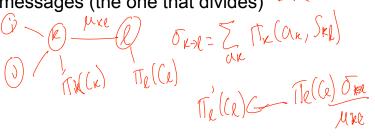
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VE versus BP in clique trees

■ VE messages (the one that multiplies) (x=Ax 0)

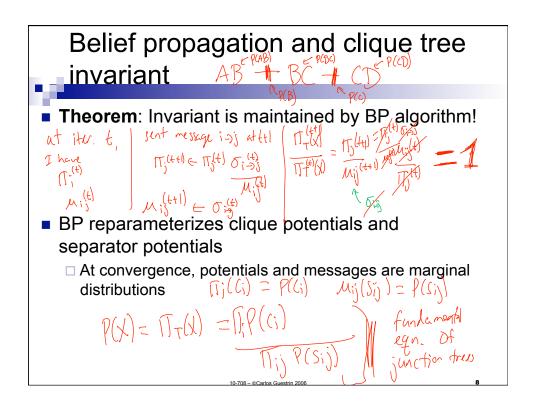


■ BP messages (the one that divides) とうし



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Clique tree invariant Clique tree potential: Product of clique potentials divided by separators potentials $T_{T}(X) = \prod_{i} \pi_{i}(C_{i}) \qquad T_{0}(C_{i}) = \text{product of } C \text{ PTs assigned to } C \text{ PTs$



Subtree correctness

- Informed message from i to j, if all messages into i (other than from j) are informed
 - □ Recursive definition (leaves always send informed messages)
- Informed subtree:
 - ☐ All incoming messages informed
- Theorem:
 - □ Potential of connected informed subtree T' is marginal over scope[T']
- Corollary:
 - ☐ At convergence, clique tree is calibrated
 - $\pi_i = P(scope[\pi_i])$
 - μ_{ii} = P(scope[μ_{ii}])

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Clique trees versus VE for inference

- Clique tree advantages
 - □ Multi-query settings
 - □ Incremental updates
 - □ Pre-computation makes complexity explicit
- Clique tree disadvantages
 - □ Space requirements no factors are "deleted"
 - ☐ Slower for single query
 - □ Local structure in factors may be lost when they are multiplied together into initial clique potential

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Clique tree summary



- Solve marginal queries for all variables in only twice the cost of query for one variable
- Cliques correspond to maximal cliques in induced graph
- Two message passing approaches
 - □ VE (the one that multiplies messages)
 - □ BP (the one that divides by old message)
- Clique tree invariant
 - ☐ Clique tree potential is always the same
 - □ We are only reparameterizing clique potentials
- Constructing clique tree for a BN
 - □ from elimination order
 - ☐ from triangulated (chordal) graph
- Running time (only) exponential in size of largest clique
 - □ Solve **exactly** problems with thousands (or millions, or more) of variables, and cliques with tens of nodes (or less)

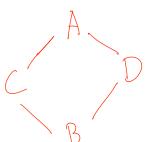
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Swinging Couples revisited



- This is no perfect map in BNs
- But, an undirected model will be a perfect map



CID LAE

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