

Bayesian Networks

5. Exact Inference / Clustering

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Bayesian Networks



1. Trees

- 2. Cluster Trees
- **3. Recursive Computation of Link Potentials**
- 4. Clique (Cluster) Trees
- 5. Triangulation

Components and cycles



Definition 1. Let G be an undirected graph. G is called **connected**, if there is a path from any vertex to any other vertex:

 $G^*(v,w)\neq \emptyset, \quad \forall v,w \in V$

For a vertex $v \in V$ we call

 $\operatorname{comp}_G(v) := \{ w \, | \, G^*(v,w) \neq \emptyset \}$

the (connection) component of v in G.

A proper path $p = (v_1, \ldots, v_n)$ is called **cyclic**, if $v_1 = v_n$ and v_i are pairwise different otherwise:

 $v_i = v_j \Leftrightarrow i = 1 \text{ and } j = n$

A proper path $p = (v_1, \ldots, v_n)$ is called **simple**, if v_i are pairwise different.

An undirected graph G is called **acyclic**, if it does not contain a cyclic path.

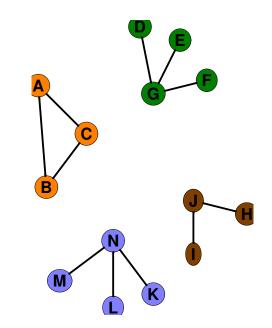


Figure 1: Graph with four components (colored).

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Bayesian Networks / 1. Trees

Trees

Definition 2. An undirected graph *G* is called **unrooted/undirected tree**, if

- (i) it is connected and acyclic or equivalently
- (ii) there is exactly one simple path between any two vertices:

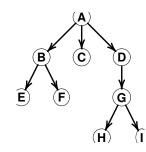
$$|G^*_{\mathsf{simple}}(v,w)| = 1, \quad \forall v, w \in V$$

The unique simple path between v and w is denoted by $path_G(v, w)$.

A directed graph *G* is called **(rooted/directed) tree**, if every vertex but one (called **root**) has exactly one parent and the root has no parents:

$$\exists r \in V : \operatorname{pa}(r) = \emptyset \text{ and } \forall v \in V, v \neq r : |\operatorname{pa}(v)| = 1$$

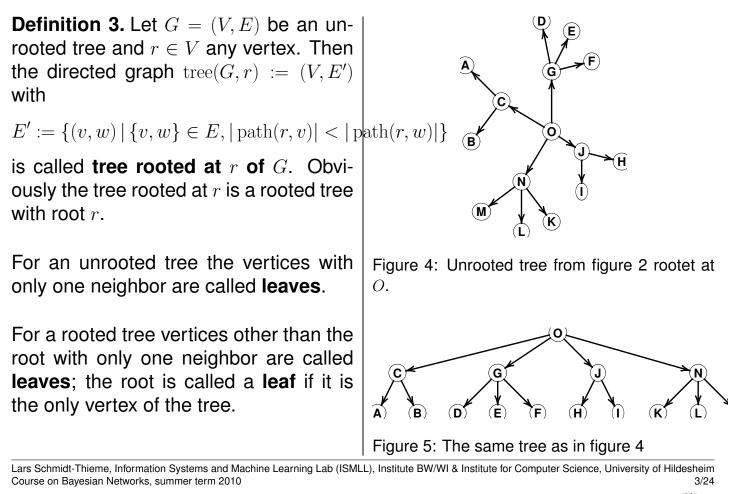
Figure 2: An unrooted tree.



Rooted trees are special DAGs. Figure 3: A (rooted) tree.

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Bayesian Networks / 1. Trees



Definition 4. Let G be a DAG (e.g., a rooted tree). The length of the longest path is called the **depth of** G and denoted by depth(G).

Let G := (V, E) be a DAG (e.g., a rooted tree). A map

 $\lambda:V\to\mathbb{N}$

is called level map of G if

 $\lambda(v) > \lambda(\mathrm{pa}(v)), \quad \forall v \in V$

For a rooted tree G := (V, E) with root r,

depth(v) := |path(r, v)|

and

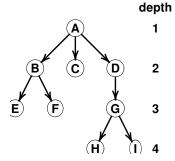


Figure 6: The depth level map for a tree.

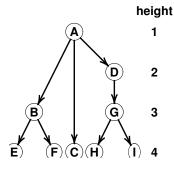


Figure 7: The height level map for a tree.

 $\operatorname{height}(v) := \operatorname{depth}(G) - \max\{|p| \mid w \in V \text{ leaf}, p \in G^*(v, w), r \not\in p\} + 1$

are examples for level maps.

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Links, polytrees



Definition 5. Let G := (V, E) be an undirected graph. The set

$$L_G := \{ (v, w) \mid \{ v, w \} \in E \}$$

is called its set of links.

Definition 6. A directed graph G is called **polytree**, if for each vertex r without parents (called a root) its descendants desc $r \cup \{r\}$ form a tree.

or equivalently

if every vertex has at most one parent that is not a root (i.e., has parents itself). Figure 8: A polytree with roots A and B.

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Bayesian Networks



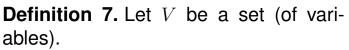
1. Trees

2. Cluster Trees

- 3. Recursive Computation of Link Potentials
- 4. Clique (Cluster) Trees
- 5. Triangulation

Cluster trees





An unrooted tree $G := (\mathcal{V}, E)$ on $\mathcal{V} \subseteq \mathcal{P}(V)$ is called a **cluster tree on** V, if

(i) the induced subgraph on all vertices containing a given variable *v*, i.e.,

$$\{W \in \mathcal{V} \mid v \in W\}$$

is connected for all variables $v \in V$. or equivalently

(ii) for any $U, W \in \mathcal{V}$:

$$U \cap W = U \cap \bigcup \operatorname{comp}_{G \setminus \{U\}}(W)$$

For two vertices U, W of a cluster tree $U \cap W$ is called their **separator**.

Cluster trees are also called **join trees** and **junction trees**.

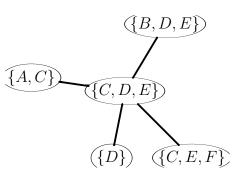


Figure 9: A cluster tree on $V := \{A, B, C, D, E, F\}.$

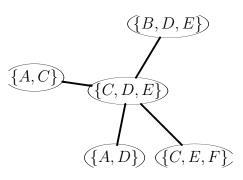


Figure 10: Not a cluster tree.

 $Q := \{ p(D), p(B), p(C|D), \}$

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Bayesian Networks / 2. Cluster Trees

Cluster trees



Definition 8. Let V be a set of variables and Q be a set of potentials on V.

A cluster tree $G := (\mathcal{V}, E)$ on V with a map

 $Q_G: \mathcal{V} \to \mathcal{P}(Q)$

(i)
$$\operatorname{dom}(q) \subseteq C$$
 for all $q \in Q_G(C)$, $C \in \mathcal{V}$,

(ii) $Im(Q_G)$ covers Q, i.e.,

$$\bigcup_{W\in\mathcal{V}}Q_G(W)=Q$$

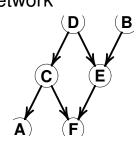
and

(iii) $Q_G(W)$ and $Q_G(U)$ are pairwise disjunct, i.e.,

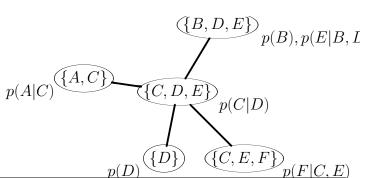
$$Q_G(W) \cap Q_G(U) \neq \emptyset \Rightarrow W = U, \forall W, U \in \mathcal{V}$$

is called a cluster tree for Q.

 $p(E|D,B), p(A|C), p(F|C,E)\}$ are the conditional probabilities of the bayesian network



A cluster tree for Q is, e.g.,



Lars Schmidt-Thieme, Information Systems and Machine Learning Lab (ISMLL), Institute BW/WI & Institute for Computer Science, University of Hildesheim Course on Bayesian Networks, summer term 2010 7/24 A simple cluster tree for polytree Bayesian networks

Let G be a directed graph. For $v \in V$

$$fam(v) := \{v\} \cup pa(v)$$

is called the familiy of v.

Let $(G = (V, E), (p_v)_{v \in V})$ be a polytree Bayesian network. Let

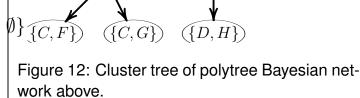
$$\mathcal{V} := \{ \operatorname{fam}(v) \, | \, v \in V \}$$

and

$$F := \{ (\operatorname{fam}(\operatorname{pa}(v)), \operatorname{fam}(v)) \mid v \in V, \operatorname{pa}(v) \neq 0 \}$$

Then $H := (\mathcal{V}, F)$ is a cluster tree for $Q := \{p_v | v \in V\}$ called family tree.





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Bayesian Networks / 2. Cluster Trees

Clique cluster tree for Markov networks

Markov networks $(G, (q_C)_{C \in \mathcal{C}(G)})$ use potentials on cliques to specify the JPD. If G is triangulated, it allows a chain of cliques, i.e., an ordering C_1, \ldots, C_n of the cliques that satisfies the running intersection property:

$$C_i \cap \bigcup_{j < i} C_j \subseteq C_{k(i)}, \quad \forall i \exists k(i) < i$$

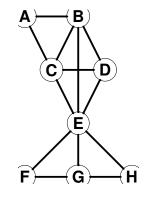
We can construct the clique (cluster) tree $H := (\mathcal{V}, F)$ from

$$\mathcal{V} := \mathcal{C}(G) = \{C_1, \dots, C_n\}$$

and

$$F := \{ (C_{k(i)}, C_i) \mid i = 2, \dots, n \}$$

We will later address the problem of cluster trees for non-triangulated





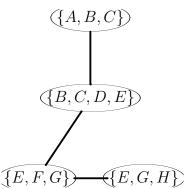
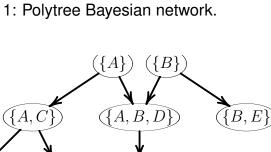


Figure 14: Clique cluster tree of Markov network

Markov networks. Lars Schmidt-Thieme, Information Systems and Machine Learning Lab (ISMLL), Institute BW/WI & Institute for Computer Science, University of Hildesheim Course on Bayesian Networks, summer term 2010 9/24





Clique cluster tree for Bayesian networks



Cluster trees for Bayesian networks can be constructed by a two phase approach:

- (i) construct an equivalent Markov network representation of the Bayesian network,
- (ii) construct the clique cluster tree for the Markov network.

An equivalent Markov network for a Bayesian network $(G = (V, E), (p_v)_{v \in V})$ can be constructed by

moral(G)

and assigning the conditional probabilities to cliques that contain their domain.

Figure 15: Bayesian network.

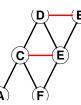


Figure 16: Markov network for Bayesian network above.

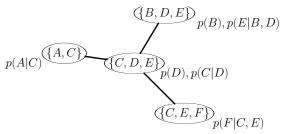


Figure 17: Clique cluster tree for Markov net-

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- 1. Trees
- 2. Cluster Trees

3. Recursive Computation of Link Potentials

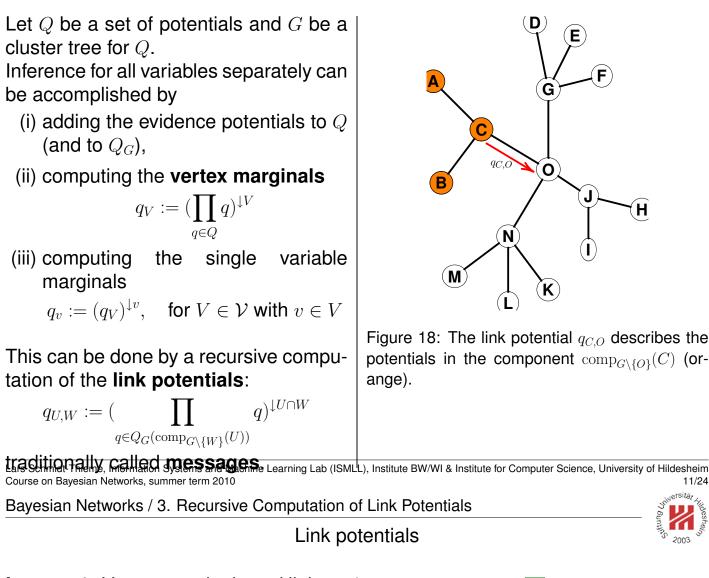
- 4. Clique (Cluster) Trees
- 5. Triangulation

Bayesian Networks / 3. Recursive Computation of Link Potentials

Vertex marginals and link potentials



11/24



Lemma 1. Vertex marginals and link potentials can be expressed by link potentials:

(i)

$$q_U = \prod_{q \in Q_G(U)} q \prod_{T \in \text{fan}(U)} q_{T,U}$$

(ii)

$$q_{U,W} = \left(\prod_{q \in Q_G(U)} q \prod_{\substack{T \in \text{fan}(U), \\ T \neq W}} q_{T,U}\right)^{\downarrow U \cap W}$$

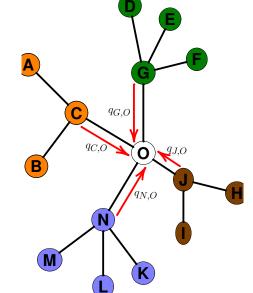


Figure 19: Expressing the vertex potential q_O by the linkpotentials $q_{.O}$.

Link potentials



Lemma 1. Vertex marginals and link potentials can be expressed by link potentials:

(i)

$$q_U = \prod_{q \in Q_G(U)} q \prod_{T \in \text{fan}(U)} q_{T,U}$$

(ii)

$$q_{U,W} = \left(\prod_{q \in Q_G(U)} q \prod_{\substack{T \in \text{fan}(U), \\ T \neq W}} q_{T,U}\right)^{\downarrow U \cap W}$$

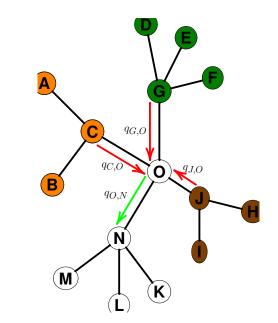


Figure 20: Expressing the link potential $q_{O,N}$ by the linkpotentials $q_{.,O}$.

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Bayesian Networks / 3. Recursive Computation of Link Potentials



Recursive computation of link potentials

Lemma 2. The formula of the previous lemma allows the recursive computation of link potentials in a cluster tree *G*.

Proof. Choose an arbitrary vertex as root and replace *G* by its rooted tree. Let λ be a level map of *G* and λ_{\min} , λ_{\max}

its minimal and maximal values. I. up links (**collect evidence**): induction

on $n := \lambda(U)$ for link potentials $q_{U, pa(U)}$.

- $n = \lambda_{\max}$: U is a leaf and has no other neighbors other than its parent.
- $n \rightarrow n-1$: the link potentials from childs into U have already been computed by induction hypothesis. \Rightarrow $q_{U,\mathrm{pa}(U)}$ can be computed (G is a tree, thus U has at most one parent).

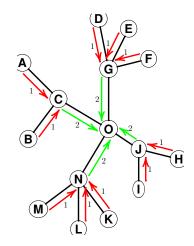


Figure 21: Collect evidence.

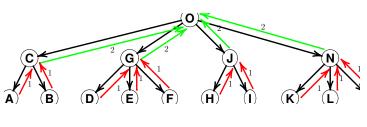


Figure 22: Collect evidence.

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Recursive computation of link potentials



Lemma 2. The formula of the previous lemma allows the recursive computation of link potentials in a cluster tree *G*.

Proof (cont.).

II. down links (**distribute evidence**): induction on $n := \lambda(pa(U))$ for link potentials $q_{pa(U),U}$.

- $n = \lambda_{\min}$: pa(U) is the root. All of its neighboring link potentials have been computed by step I. $\Rightarrow q_{pa(U),U}$ can be computed.
- $n \rightarrow n+1$: the link potentials from childs into pa(U) have already been computed by step I, the link potential $q_{pa(pa(U)),pa(U)}$ has already been computed by induction hypothesis. \Rightarrow $q_{pa(U),U}$ can be computed.

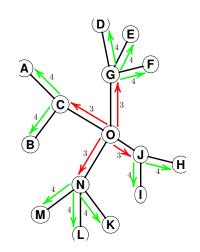


Figure 23: Distribute evidence.

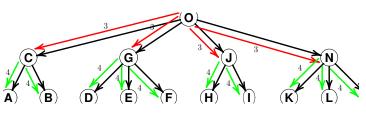


Figure 24: Distribute evidence.

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Bayesian Networks / 3. Recursive Computation of Link Potentials



Shafer-Shenoy propagation

The following computation scheme is called Shafer-Shenoy propagation []:

(i) collect evidence:

$$q_{U,W} = \left(\prod_{q \in Q_G(U)} q \prod_{\substack{T \in \text{fan}(U), \\ T \neq W}} q_{T,U}\right)^{\downarrow U \cap W} = \left(\left(\prod_{q \in Q_G(U)} q\right) \cdot q_{T_1,U} \cdots q_{T_n,U}\right)^{\downarrow U \cap W}$$

(ii) distribute evidence:

$$q_{U,T_i} = (\prod_{q \in Q_G(U)} q \prod_{\substack{T \in \text{fan}(U), \\ T \neq T_i}} q_{T,U})^{\downarrow U \cap T_i} = ((\prod_{q \in Q_G(U)} q) \cdot q_{W,U} \cdot q_{T_1,U} \cdots \widehat{q_{T_i,U}} \cdots q_{T_n,U})^{\downarrow U \cap T_i}$$

(iii) marginalize:

$$q_U = \prod_{q \in Q_G(U)} q \prod_{T \in \text{fan}(U)} q_{T,U} \qquad = (\prod_{q \in Q_G(U)} q) \cdot q_{W,U} \cdot q_{T_1,U} \cdots q_{T_n,U}$$

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Hugin propagation

The following computation scheme is called Hugin propagation []:

(i) collect evidence:

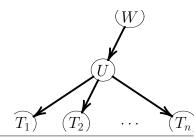
$$q'_{U} = \prod_{q \in Q_{G}(U)} q \prod_{\substack{T \in \text{fan}(U) \\ T \neq W}} q_{T,U} = (\prod_{q \in Q_{G}(U)} q) \cdot q_{T_{1},U} \cdots q_{T_{n},U}$$
$$q_{U,W} = q'_{U}^{\downarrow U \cap W}$$

(ii) marginalize and distribute evidence:

$$q_U = q'_U \cdot q_{W,U}$$

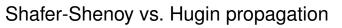
 $q_{U,T_i} = (rac{q_U}{q_{T_i,U}})^{\downarrow U \cap T_i}$

but store separator marginal $(q_U)^{\downarrow U \cap T_i}$



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Bayesian Networks / 3. Recursive Computation of Link Potentials



Hugin propagation compared to Shafer-Shenoy propagation:

- (i) Hugin propagation allows the reuse of the storage space of the link potentials $q_{U,W}$ for $q_{W,U}$ (one "postbox" instead of two),
- (ii) Hugin propagation affords extra storage space for the vertex potentials q_U and thus its overall space requirements are higher,
- (iii) Hugin propagation requires a smaller number of total operations (additions, multiplications, divisions) than Shafer-Shenoy propagation at vertices with degree > 3 (that can be avoided by the use of binary cluster trees),

- (iv) Hugin propagation allows the marginalization of the smaller separator marginals,
- (v) Some of the operations required by Hugin propagation are more costly (divisions) than those required by Shafer-Shenoy.



Lazy propagation



The idea of **lazy propagation** [MJ98] is to keep the link potentials in factored form, i.e., to replace the link potential $q_{U,W}$ with a set of potentials $Q_{U,W}$ with

$$q_{U,W} = \prod_{q \in Q_{U,W}} q$$

The formulas of lemma 1 then read as:

(i)

$$q_U = \prod_{q \in Q_G(U)} q \prod_{\substack{T \in \text{fan}(U)\\q \in Q(T,U)}} q$$

(ii)

$$q_{U,W} = \operatorname{elim}(Q_G(U) \cup \bigcup_{\substack{T \in \operatorname{fan}(U), \\ T \neq W}} Q_{T,U}, c(U \cap W))$$

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Bayesian Networks



- 1. Trees
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- **3. Recursive Computation of Link Potentials**
- 4. Clique (Cluster) Trees
- 5. Triangulation

Clique trees for triangulated graphs (1/3)

Clique cluster trees can easily be computed of triangulated graphs.

(i) Triangulated graphs admit a perfect ordering of G, i.e., an ordering σ with

$$fam_{\sigma(\{1,\dots,i\})}(\sigma(i))$$

is complete.

(ii) A perfect ordering can be computed by the maximum cardinality search algorithm (MCS).

```
i perfect-ordering-MCS(G = (V, E)):
```

- $2 \, \underline{\mathbf{for}} \, i = 1, \dots, |V| \, \underline{\mathbf{do}}$
 - $\sigma(i) := v \in V \setminus \sigma(\{1, \dots, i-1\})$ with maximal $|\operatorname{fan}_G(v) \cap \sigma(\{1, \dots, i-1\})|$
- breaking ties arbitrarily
- 5 <u>od</u>
- 6 <u>return</u> σ

Figure 26: MCS algorithm to compute a perfect ordering [TY84]. Lars Schmidt-Thieme, Information Systems and Machine Learning Lab (ISMLL), Institute BW/WI & Institute for Computer Science, University of Hildesheim Course on Bayesian Networks, summer term 2010 18/24

Bayesian Networks / 4. Clique (Cluster) Trees

Clique trees for triangulated graphs (2/3)

All cliques can be enumerated by a variant of the MCS algorithm:

- 1. if *G* is triangulated, MCS computes a perfect ordering of *G*, i.e., $fam_{\sigma(\{1,...,i\})}(\sigma(i))$ is complete.
- 2. we get all cliques this way, as for each clique C let $i := \max \sigma^{-1}(C)$, then $C = \operatorname{fam}_{\sigma(\{1,\ldots,i\})}(\sigma(i))$.
- *i* enumerate-cliques-MCS(G = (V, E)): 2 $C := \emptyset$
- $s \operatorname{\underline{\mathbf{for}}} i = 1, \dots, |V| \operatorname{\underline{\mathbf{do}}}$

$$\sigma(i) := v \in V \setminus \sigma(\{1, \dots, i-1\}) \text{ with maximal } |\operatorname{fan}_G(v) \cap \sigma(\{1, \dots, i-1\})|$$

breaking ties arbitrarily

$$\delta \qquad \mathcal{C} := \mathcal{C} \cup \{ \operatorname{fam}_{\sigma(\{1,\dots,i\})}(\sigma(i)) \}$$

7 <u>od</u>

$$\mathcal{C} := \{ C \in \mathcal{C} \mid \exists D \in \mathcal{C} : D \supseteq C \}$$

9 return
$$C$$

Figure 27: MCS algorithm to compute cliques of a triangulated graph [TY84].

Proving the correctness of MCS affords some work (e.g., [Sha94, p. 43–46]).

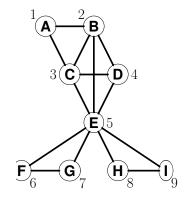
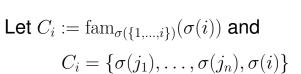


Figure 25: Perfect ordering of a triangulated graph obtained by MCS.



with $j_1 < j_2 < \ldots < j_n$. Due to the completeness of C_i then $\sigma(j_n)$ is a neighbor of all $\sigma(j_l)$, $l = 1, \ldots, n-1$, and thus

$$C_i \cap \bigcup_{k < i} C_k \subseteq C_{j_n}$$

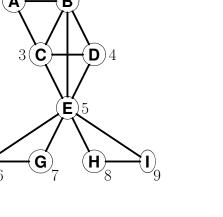
i.e., the sequence $(C_i)_{i=1,...,|V|}$ has the running intersection property (that can be telescoped if a C_i gets pruned).





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Clique trees for triangulated graphs (3/3)



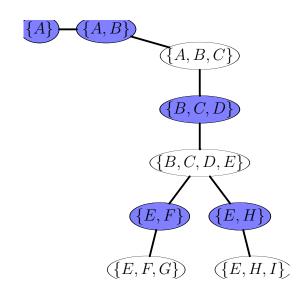


Figure 28: Perfect ordering of a triangulated graph obtained by MCS.

Figure 29: Clique cluster tree for triangulated graph at the left (blue nodes are temporary and pruned).

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Bayesian Networks



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Triangulation of graphs (1/3)

As clique cluster trees can easily be computed of triangulated graphs, we triangulate non-triangulated graphs by filling-in additional edges.

However, additional edges mean, that the graph represents a smaller portion of the independency statements, and thus, inference becomes harder.

The fewer edges have to be filled-in, the better.

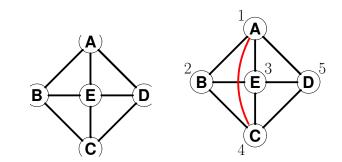


Figure 30: Non-triangulated graph and its triangulation obtained by MCS.

 $\begin{array}{l} \text{triangulate-MCS}(G = (V, E)): \\ \text{2 } \sigma := \textit{perfect-ordering-MCS}(G) \\ \text{3 } \textit{fillin} := \emptyset \\ \text{4 } \underbrace{\textbf{for}}_{5} i = |V|, \dots, 1 \underbrace{\textbf{do}}_{(V,E \cup \textit{fillin})}(\sigma(i)) \cap \sigma(\{1, \dots, i-1\}), \{u, w\} \notin E \} \end{array}$

7 **return**
$$G' := (V, E \cup fillin)$$

Figure 31: Maximum cardinality search algorithm for triangulating a graph [TY84]. Lars Schmidt-Thieme, Information Systems and Machine Learning Lab (ISMLL), Institute BW/WI & Institute for Computer Science, University of Hildesheim Course on Bayesian Networks, summer term 2010 21/24

Bayesian Networks / 5. Triangulation

Triangulation of graphs (2/3)



MCS does not guarantee to give best results (i.e., minimal fill-ins). It is just a heuristics that gives useable results (in most cases).

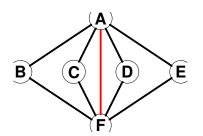


Figure 32: Optimal triangulation.

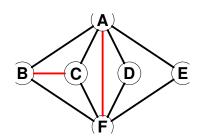


Figure 33: Non-optimal triangulation obtained by MCS (with smallest index rule).

Triangulation of graphs (3/3)

Beneath the heuristic triangulation algorithms one distinguishes between:

minimum triangulations: no other triangulation has a smaller number of filled-in edges (global minimum).

This task is known to be NP-complete [Yan81].

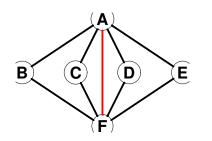


Figure 34: A minimum triangulation (here: unique).

minimal triangulations: no subset of the filled-in edges results in a triangulation (local minimum).

There are several algorithms for the minimal triangulation task, e.g., Lex-M [RTL76], MCS-M [BBH02], and LB-triang [BBH⁺03].

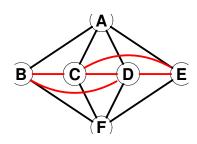


Figure 35: A minimal triangulation.

Lars Schmidt-Thieme, Information Systems and Machine Learning Lab (ISMLL), Institute BW/WI & Institute for Computer Science, University of Hildesheim Course on Bayesian Networks, summer term 2010 23/24

Bayesian Networks / 5. Triangulation

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