

Bayesian Networks

III. Exact Inference (section 3)

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I. Inference in Probabilistic Networks

II. Variable elimination

3. Clustering

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

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

the **(connection) component of v in G** .

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

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

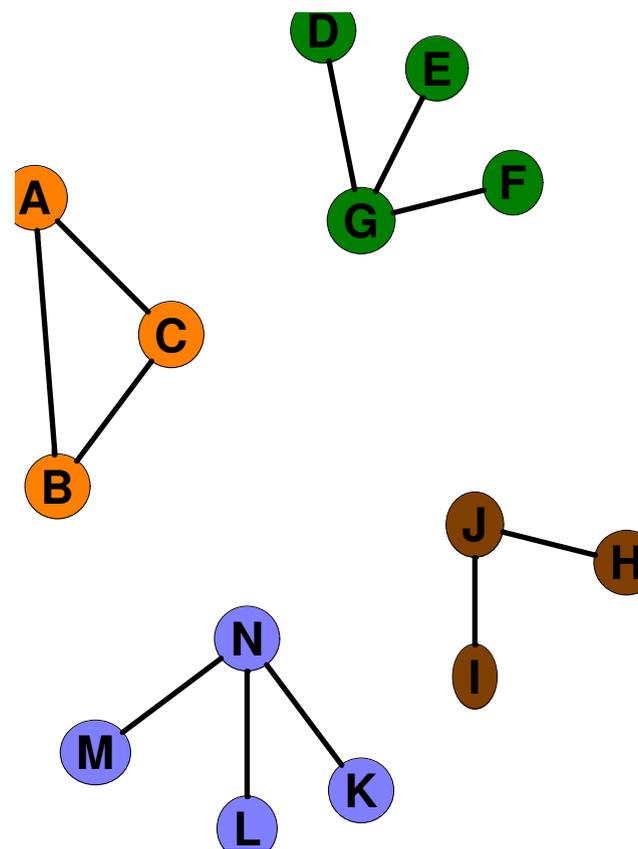


Figure 1: Graph with four components (colored).

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 path between any two vertices:

$$|G^*(v, w)| = 1, \quad \forall v, w \in V$$

The unique path between v and w is denoted by $\text{path}_G(v, w)$.

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:

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

Rooted trees are special DAGs.

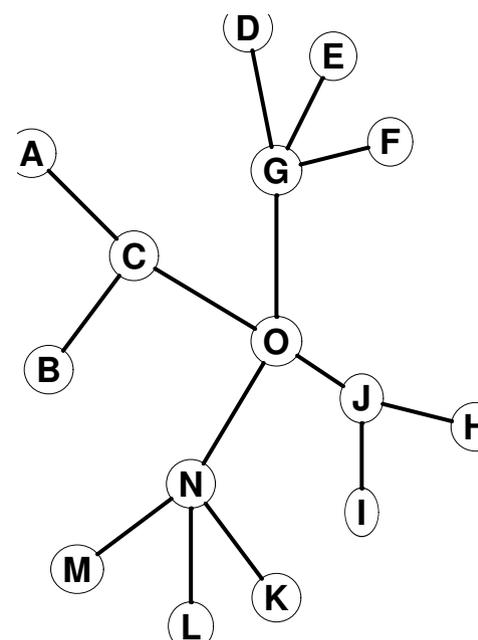


Figure 2: An unrooted tree.

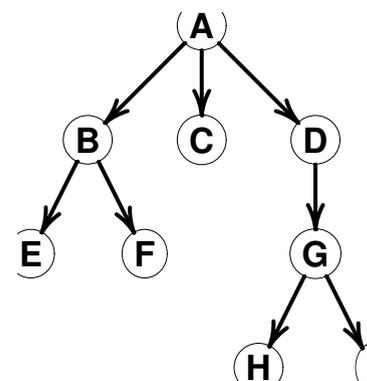


Figure 3: A (rooted) tree.

Trees / leaves

Definition 3. Let $G = (V, E)$ be an unrooted tree and $r \in V$ any vertex. Then the directed graph $\text{tree}(G, r) := (V, E')$ with

$$E' := \{(v, w) \mid \{v, w\} \in E, |\text{path}(r, v)| < |\text{path}(r, w)|\}$$

is called **tree rooted at r of G** . Obviously the tree rooted at r is a rooted tree with root r .

For an unrooted tree the vertices with only one neighbor are called **leaves**.

For a rooted tree vertices other than the root with only one neighbor are called **leaves**; the root is called a **leaf** if it is the only vertex of the tree.

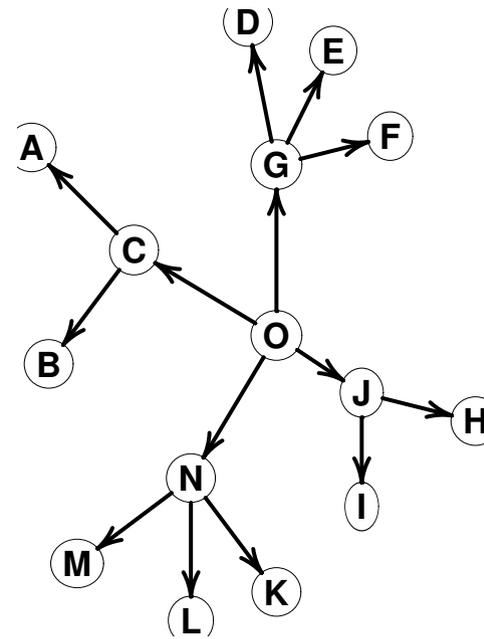


Figure 4: Unrooted tree from figure 2 rootet at O .

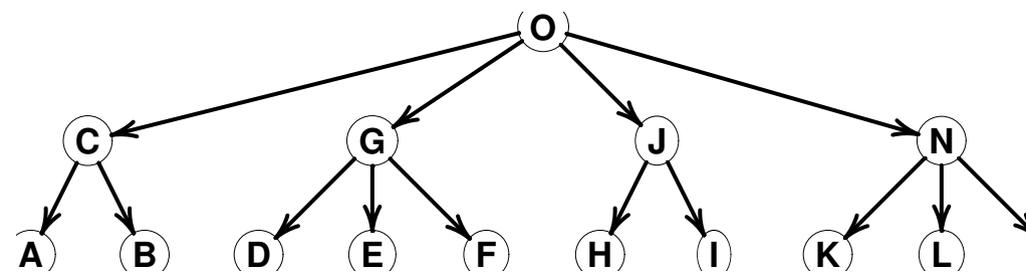


Figure 5: The same tree as in figure 4

Trees / level maps

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 $\text{depth}(G)$.

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

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

is called **level map of G** if

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

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

$$\text{depth}(v) := |\text{path}(r, v)|$$

and

$$\text{height}(v) := \text{depth}(G) - \max\{|p| \mid w \in V \text{ leaf}, p \in G^*(v, w), r \notin p\} + 1$$

are examples for level maps.

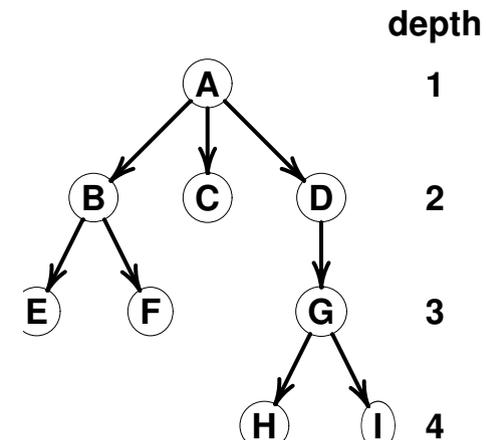


Figure 6: The depth level map for a tree.

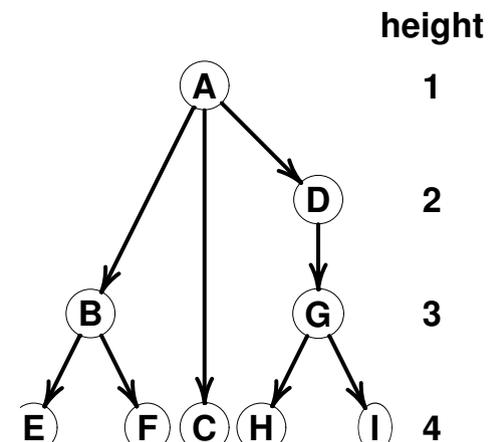


Figure 7: The height level map for a tree.

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 $\text{desc } r \cup \{r\}$ form a tree.

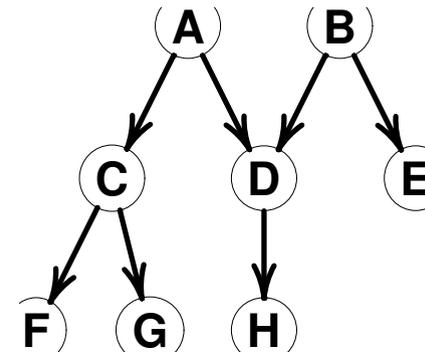


Figure 8: A polytree with roots A and B .

Cluster trees

Definition 7. Let V be a set (of variables).

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)

$$U \cap W = U \cap \bigcup_{\text{comp}_{G \setminus \{U\}}(W)}$$

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**.

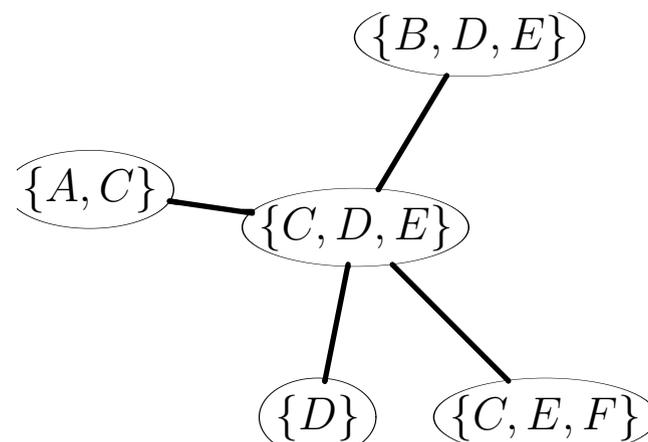


Figure 9: A cluster tree on $V := \{A, B, C, D, E, F\}$.

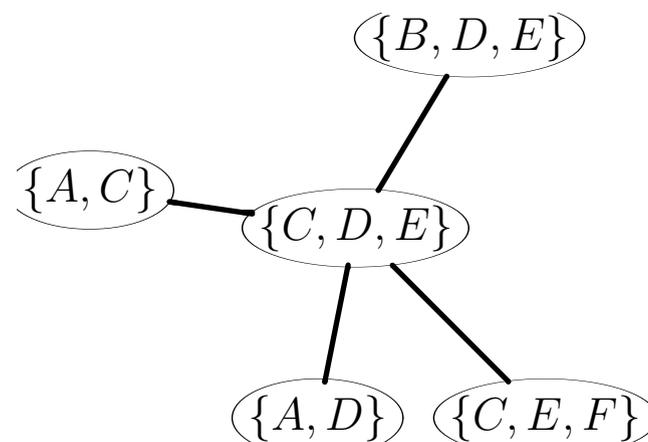


Figure 10: Not a cluster tree.

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

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

s.t.

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

(ii) $\text{Im}(Q_G)$ covers Q , i.e.,

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

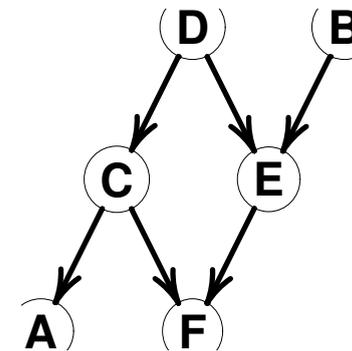
and

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

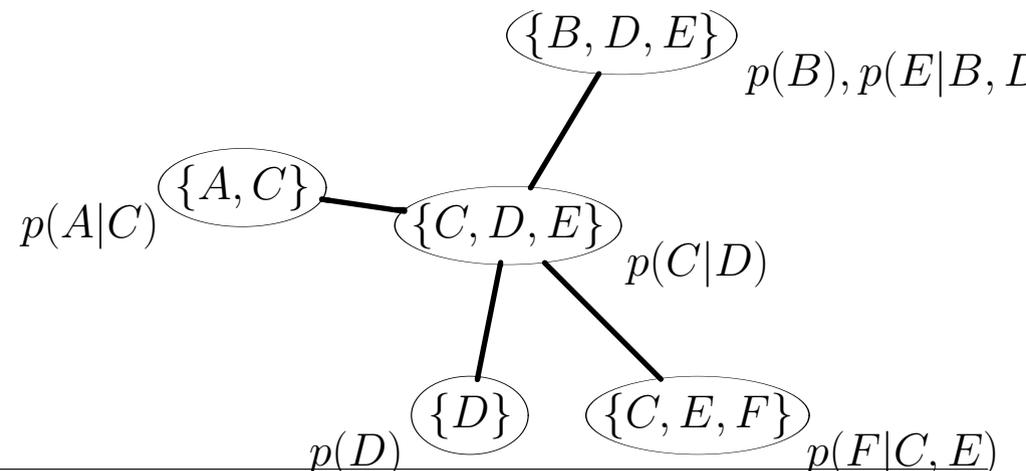
$$Q_G(W) \cap Q_G(U) \neq \emptyset \Rightarrow W = U, \quad \forall W, U \in \mathcal{V}$$

is called a **cluster tree for Q** .

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



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



A simple cluster tree for polytree Bayesian networks

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

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

is called the **family of** v .

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

$$\mathcal{V} := \{\text{fam}(v) \mid v \in V\}$$

and

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

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

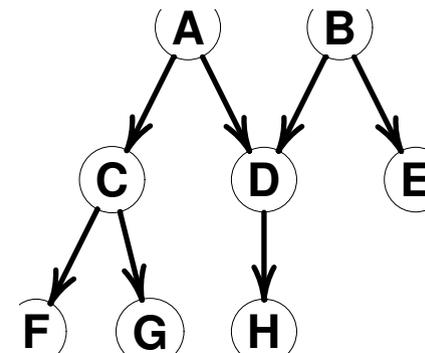


Figure 11: Polytree Bayesian network.

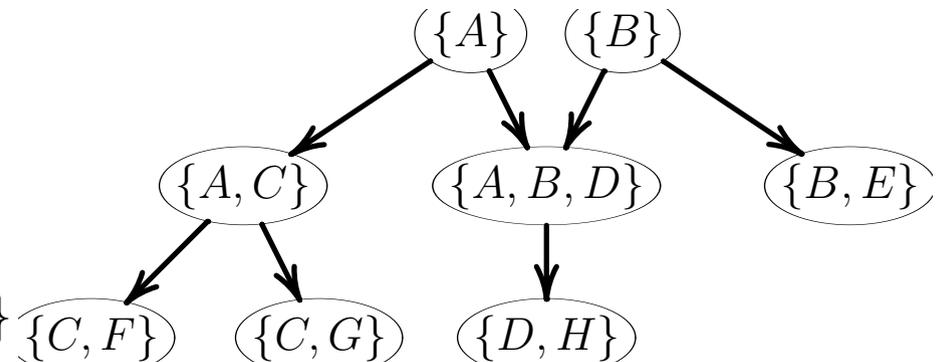


Figure 12: Cluster tree of polytree Bayesian network above.

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, \dots, 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 Markov networks.

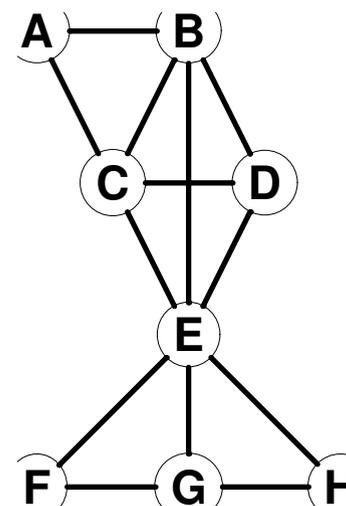


Figure 13: Markov network.

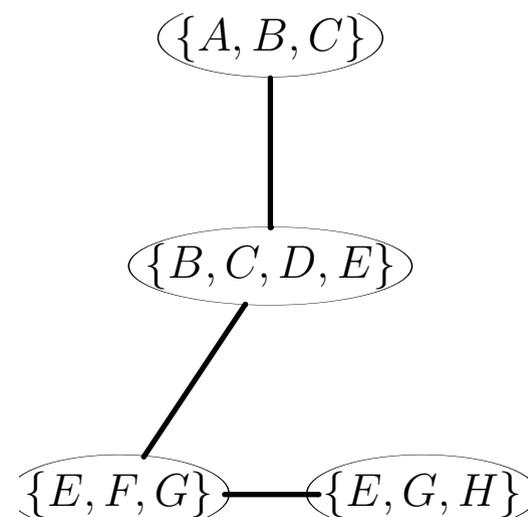


Figure 14: Clique cluster tree of Markov network above.

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

$$\text{moral}(G)$$

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

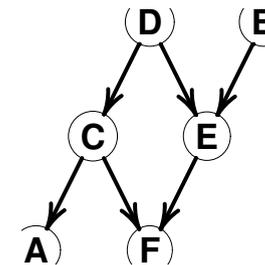


Figure 15: Bayesian network.

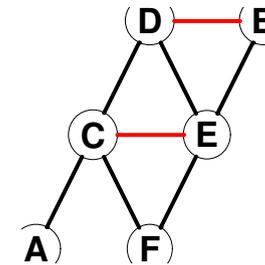


Figure 16: Markov network for Bayesian network above.

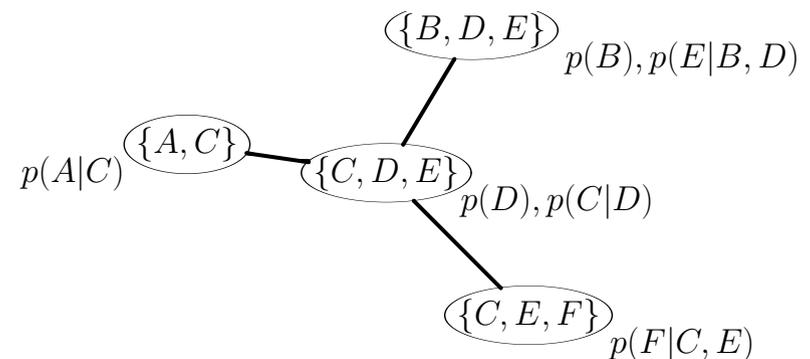


Figure 17: Clique cluster tree for Markov network above.

Vertex marginals and link potentials

Let Q be a set of potentials and G be a cluster tree for Q .

Inference for all variables separately can be accomplished by

- (i) adding the evidence potentials to Q (and to Q_G),
- (ii) computing the **vertex marginals**

$$q_V := \left(\prod_{q \in Q} q \right)^{\downarrow V}$$

- (iii) computing the single variable marginals

$$q_v := (q_V)^{\downarrow v}, \quad \text{for } V \in \mathcal{V} \text{ with } v \in V$$

This can be done by a recursive computation of the **link potentials**:

$$q_{U,W} := \left(\prod_{q \in Q_G(\text{comp}_{G \setminus \{W\}}(U))} q \right)^{\downarrow U \cap W}$$

traditionally called **messages**

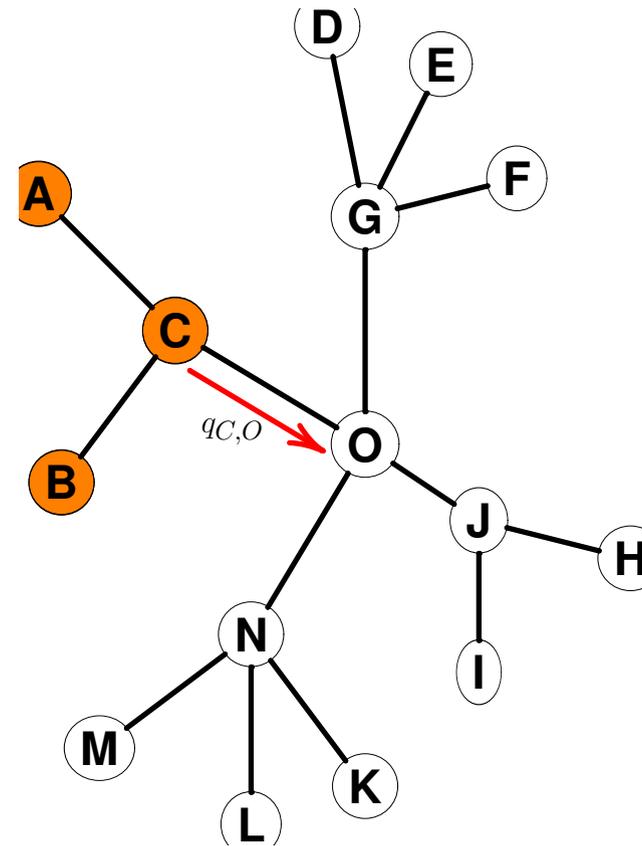


Figure 18: The link potential $q_{C,O}$ describes the potentials in the component $\text{comp}_{G \setminus \{O\}}(C)$ (orange).

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)

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

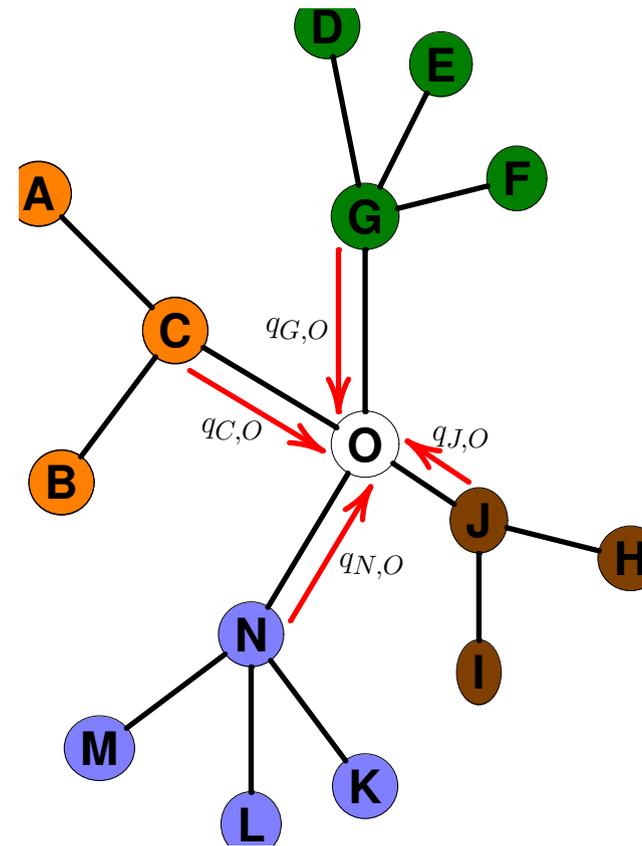


Figure 19: Expressing the vertex potential q_O by the linkpotentials $q_{.,O}$.

Link potentials

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

(i)

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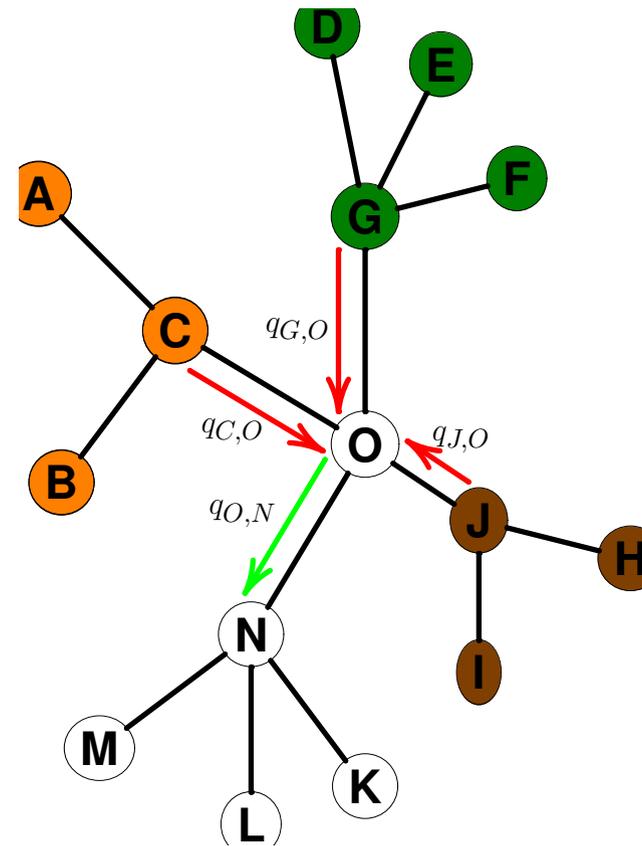


Figure 20: Expressing the link potential $q_{O,N}$ by the linkpotentials $q_{.,O}$.

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 $\lambda_{\min}, \lambda_{\max}$ its minimal and maximal values.

I. up links (**collect evidence**): induction on $n := \lambda(U)$ for link potentials $q_{U, \text{pa}(U)}$.

$n = \lambda_{\max}$: 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

$q_{U, \text{pa}(U)}$ can be computed (G is a tree, thus U has at most one parent).

□

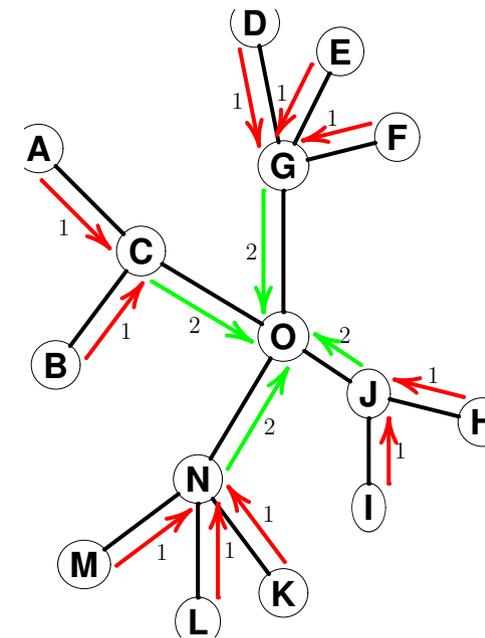


Figure 21: Collect evidence.

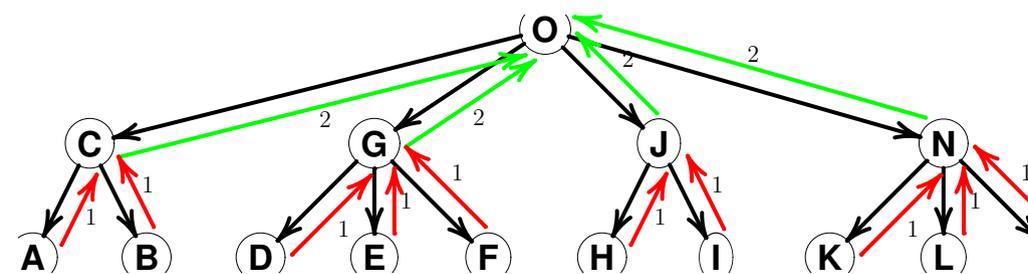


Figure 22: Collect evidence.

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

$n = \lambda_{\min}$: $\text{pa}(U)$ is the root. All of its neighboring link potentials have been computed by step I. $\Rightarrow q_{\text{pa}(U),U}$ can be computed.

$n \rightarrow n + 1$: the link potentials from childs into $\text{pa}(U)$ have already been computed by step I, the link potential $q_{\text{pa}(\text{pa}(U)),\text{pa}(U)}$ has already been computed by induction hypothesis. $\Rightarrow q_{\text{pa}(U),U}$ can be computed.

□

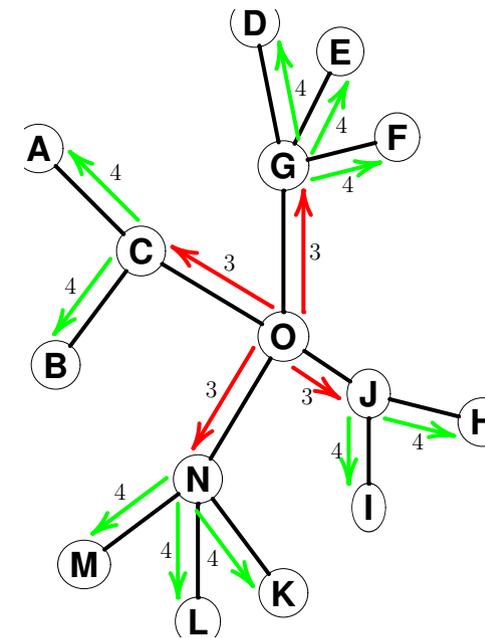


Figure 23: Distribute evidence.

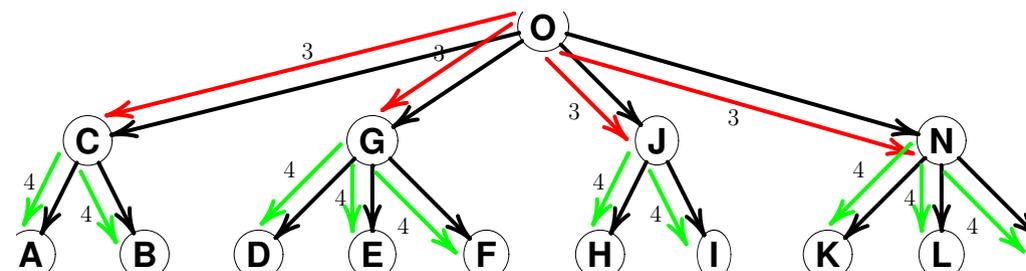


Figure 24: Distribute evidence.

Shafer-Shenoy propagation

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

(i) collect evidence:

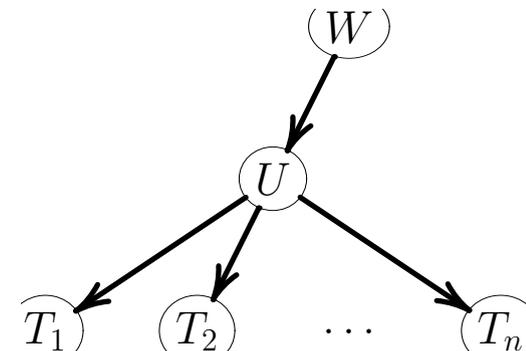
$$q_{U,W} = \left(\prod_{q \in Q_G(U)} q \prod_{\substack{T \in \text{fan}(U), \\ T \neq W}} q_{T,U} \right)^{\downarrow U \cap W} = \left(\left(\prod_{q \in Q_G(U)} q \right) \cdot q_{T_1,U} \cdots q_{T_n,U} \right)^{\downarrow U \cap W}$$

(ii) distribute evidence:

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

(iii) marginalize:

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



Hugin propagation

The following computation scheme is called **Hugin propagation** []:

(i) collect evidence:

$$q'_U = \prod_{q \in Q_G(U)} q \prod_{\substack{T \in \text{fan}(U) \\ T \neq W}} q_{T,U} = \left(\prod_{q \in Q_G(U)} q \right) \cdot q_{T_1,U} \cdots q_{T_n,U}$$

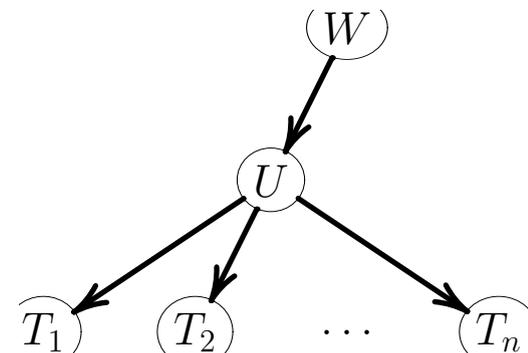
$$q_{U,W} = q'_U \downarrow^{U \cap W}$$

(ii) marginalize and distribute evidence:

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

$$q_{U,T_i} = \left(\frac{q_U}{q_{T_i,U}} \right) \downarrow^{U \cap T_i}$$

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



Shafer-Shenoy vs. Hugin propagation

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)

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

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

$$\text{fam}_{\sigma(\{1, \dots, i\})}(\sigma(i))$$

is complete.

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

```

1 perfect-ordering-MCS( $G = (V, E)$ ) :
2 for  $i = 1, \dots, |V|$  do
3    $\sigma(i) := v \in V \setminus \sigma(\{1, \dots, i-1\})$  with maximal  $|\text{fan}_G(v) \cap \sigma(\{1, \dots, i-1\})|$ 
4   breaking ties arbitrarily
5 od
6 return  $\sigma$ 

```

Figure 26: MCS algorithm to compute a perfect ordering [TY84].

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

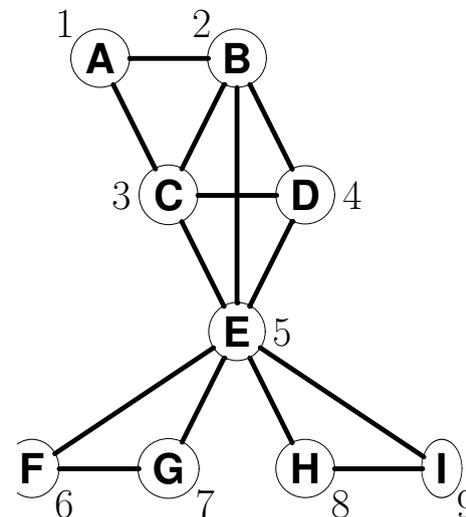


Figure 25: Perfect ordering of a triangulated graph obtained by MCS.

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., $\text{fam}_{\sigma(\{1, \dots, i\})}(\sigma(i))$ is complete.
2. we get all cliques this way, as for each clique C let $i := \max \sigma^{-1}(C)$, then $C = \text{fam}_{\sigma(\{1, \dots, i\})}(\sigma(i))$.

1 **enumerate-cliques-MCS**($G = (V, E)$) :

2 $\mathcal{C} := \emptyset$

3 **for** $i = 1, \dots, |V|$ **do**

4 $\sigma(i) := v \in V \setminus \sigma(\{1, \dots, i-1\})$ with maximal $|\text{fan}_G(v) \cap \sigma(\{1, \dots, i-1\})|$

5 breaking ties arbitrarily

6 $\mathcal{C} := \mathcal{C} \cup \{\text{fam}_{\sigma(\{1, \dots, i\})}(\sigma(i))\}$

7 **od**

8 $\mathcal{C} := \{C \in \mathcal{C} \mid \nexists D \in \mathcal{C} : D \supseteq C\}$

9 **return** \mathcal{C}

Let $C_i := \text{fam}_{\sigma(\{1, \dots, i\})}(\sigma(i))$ and

$$C_i = \{\sigma(j_1), \dots, \sigma(j_n), \sigma(i)\}$$

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

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

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

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

Clique trees for triangulated graphs (3/3)

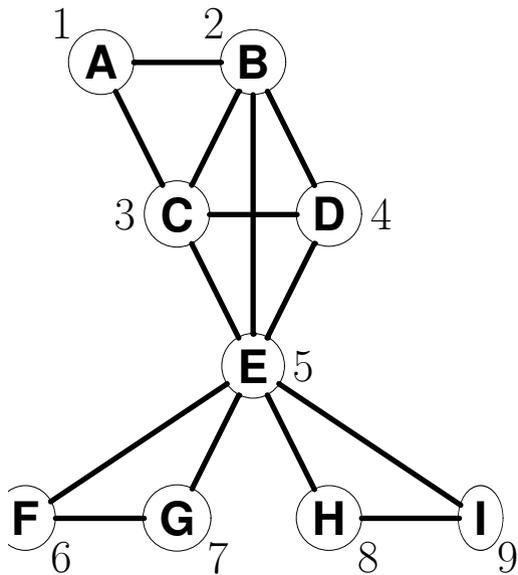


Figure 28: Perfect ordering of a triangulated graph obtained by MCS.

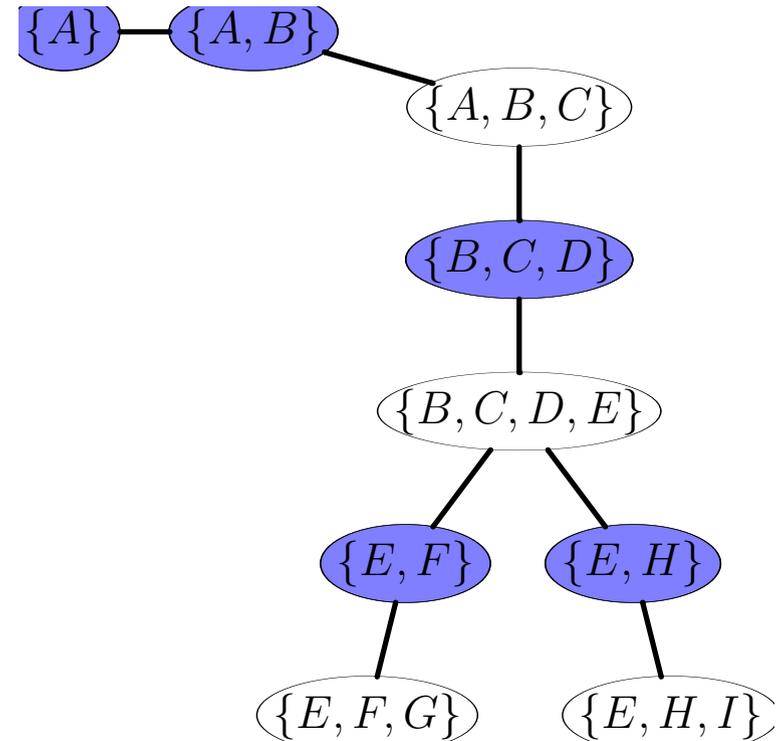


Figure 29: Clique cluster tree for triangulated graph at the left (blue nodes are temporary and pruned).

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.

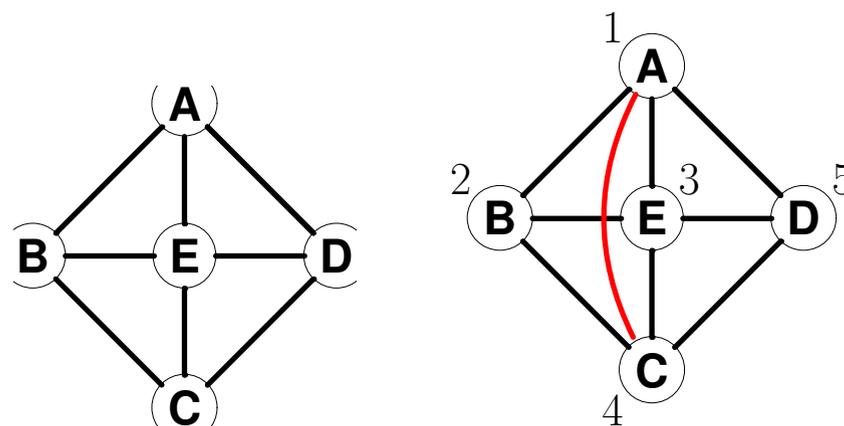


Figure 30: Non-triangulated graph and its triangulation obtained by MCS.

```

1 triangulate-MCS( $G = (V, E)$ ) :
2  $\sigma := \text{perfect-ordering-MCS}(G)$ 
3  $\text{fillin} := \emptyset$ 
4 for  $i = |V|, \dots, 1$  do
5    $\text{fillin} := \text{fillin} \cup \{(u, w) \mid u, w \in \text{fan}_{(V, E \cup \text{fillin})}(\sigma(i)) \cap \sigma(\{1, \dots, i-1\}), \{u, w\} \notin E\}$ 
6 od
7 return  $G' := (V, E \cup \text{fillin})$ 

```

Figure 31: Maximum cardinality search algorithm for triangulating a graph [TY84].

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).

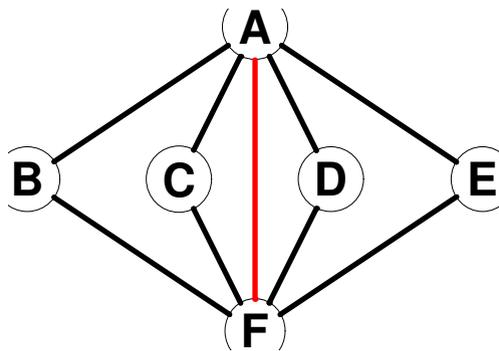


Figure 32: Optimal triangulation.

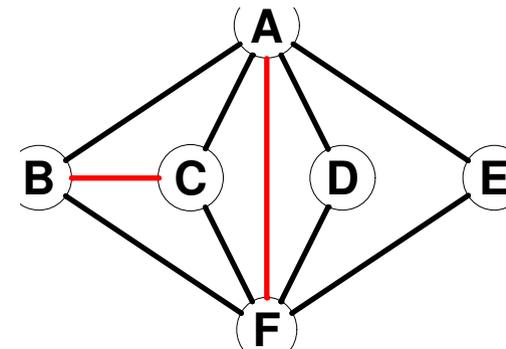


Figure 33: Non-optimal triangulation obtained by MCS (with smallest index rule).

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].

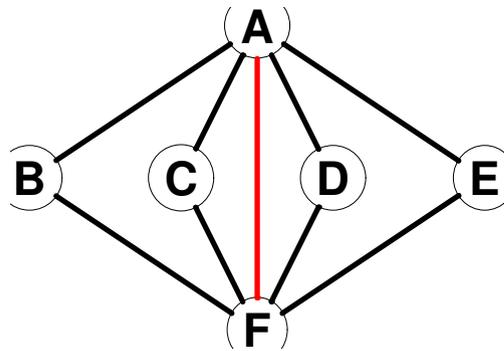


Figure 34: A minimum triangulation (here: unique).

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].

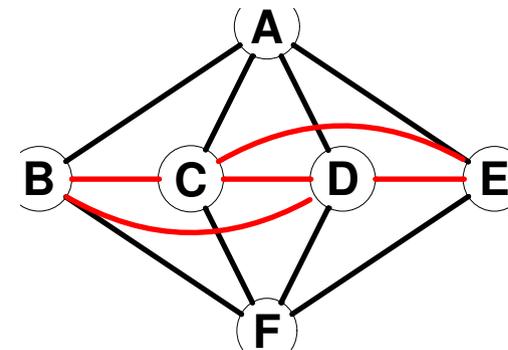


Figure 35: A minimal triangulation.

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