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Graphical Models with R 1st talk: Graphs and Markov properties with R

Presentation · July 2018

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Graphical Models with R

1st talk: Graphs and Markov properties with R

Dhafer Malouche

essai.academia.edu/DhaferMalouche

- Terminology of graphs
 Undirected graphs (UG)
 Directed Acyclic Graphs (DAG)
- 2. Markov properties
- 3. Visualizing graphs with **R**

Terminology of graphs

What's a graph?

- A graph is $\mathcal{G} = (V, E)$ where V is a finite set and $E \subseteq V \times V$.
 - V the of vertices
 - *E* is the set edges (its elements are denoted by $\alpha\beta$)
- undirected when $\alpha \beta$:

$$\alpha\beta\in E \iff \beta\alpha\in E$$

- directed when $\alpha \rightarrow \beta$ or $\alpha \leftarrow \beta$

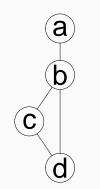
If
$$\alpha\beta \in E \Rightarrow \beta\alpha \notin E$$

• bi-directed when $\alpha \leftrightarrow \beta$

Undirected Graphs

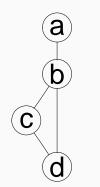
Undirected Graph (UG)

- $V = \{a, b, c, d, e\}$
- $E = \{ab, bc, cd, bd\}$
- cliques: ab, bcd, e



- $V = \{a, b, c, d, e\}$
- $E = \{ab, bc, cd, bd\}$
- cliques: ab, bcd, e

A *clique* in *G* is a maximal complete subset of *V*



UG with R, gRbase

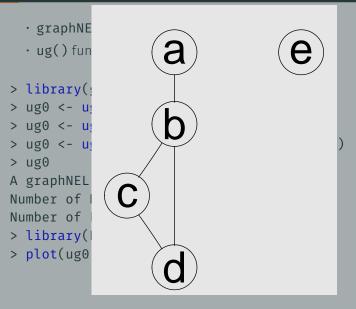
- graphNEL objects
- \cdot ug() function

UG with R, gRbase

- graphNEL objects
- \cdot ug() function

```
> library(gRbase)
> ug0 <- ug(~a:b,~b:c:d,~e)
> ug0 <- ug(~a:b+b:c:d+e)
> ug0 <- ug(c("a","b"),c("b","c","d"),"e")
> ug0
A graphNEL graph with undirected edges
Number of Nodes = 5
Number of Edges = 4
> library(Rgraphviz)
> plot(ug0)
```

UG with R, gRbase



Adjacency matrix with R, gRbase

$$\mathcal{G} = (V, E) \longmapsto A = [a(\alpha\beta)] \in \{0, 1\}^{|V| \times |V|} \text{ such that}$$
$$a(\alpha\beta) = 1 \iff \alpha\beta \in E$$

Adjacency matrix with R, gRbase

$$\mathcal{G} = (V, E) \longmapsto A = [a(\alpha\beta)] \in \{0, 1\}^{|V| \times |V|} \text{ such that}$$
$$a(\alpha\beta) = 1 \iff \alpha\beta \in E$$

```
> ug01 <- ug(~a:b+b:c:d+e,result="matrix")
> ug01
    a b c d e
a 0 1 0 0 0
b 1 0 1 1 0
c 0 1 0 1 0
d 0 1 1 0 0
e 0 0 0 0
```

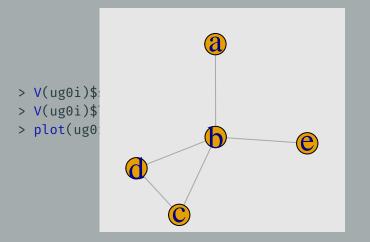
```
> nodes(ug0)
[1] "a" "b" "c" "d" "e"
> edges(ug0)
$a
[1] "b"
$b
[1] "c" "d" "a"
$c
[1] "d" "b"
$d
[1] "b" "c"
$e
character(0)
```

UG with R, igraph

```
> ug0i <- ug(c("a","b"),c("b","c","d"),c("e","b"),</pre>
                result="igraph")
+
> ug0i
TGRAPH UNW- 5 5 --
+ attr: name (v/c), label (v/c), weight (e/n)
+ edges (vertex names):
[1] a--b b--c b--d b--e c--d
> library(igraph)
> ## vertices
> V(ug0i)
+ 5/5 vertices, named:
[1] a b c d e
> ## edges
> E(ug0i)
+ 5/5 edges (vertex names):
[1] a--b b--c b--d b--e c--d
```

- > V(ug0i)\$size <- 25
- > V(ug0i)\$label.cex <- 2</pre>
- > plot(ug0i, layout=layout.spring)

UG with R, igraph



Cliques in an UG with R,

```
> library(RBGL)
> is.complete(ug0, c("b","c","d"))
[1] TRUE
> maxClique(ug0)
$maxCliques
$maxCliques[[1]]
[1] "b" "c" "d"
```

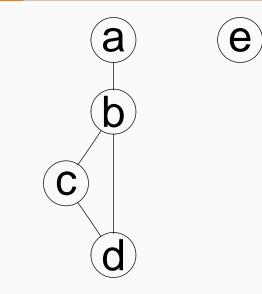
\$maxCliques[[2]] [1] "b" "a"

```
$maxCliques[[3]]
[1] "e"
```

Paths and separators in an UG

- A path (of length *n*) between α and β in an undirected graph is a set of vertices $\alpha = \alpha_0, \alpha_1, \dots, \alpha_n = \beta$ where $\alpha_{i-1} \sim \alpha_i$ for $i = 1, \dots, n$.
- If $\alpha = \beta$ then the path is said to be a cycle of length *n*.
- A subset $S \subset V$ in an undirected graph is said to separate $A \subseteq V$ from $B \subseteq V$ if every path between a vertex in A and a vertex in B intersects S.

Paths and separators in an UG



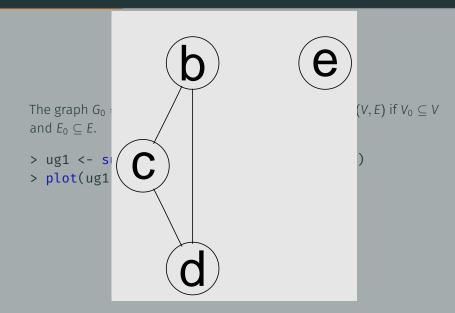
> separates(a = "a", b = "d",S1 = c("b", "c"), ug0)
[1] TRUE
> separates(a = "a", b = "b",S1 = c("d", "c"), ug0)
[1] FALSE

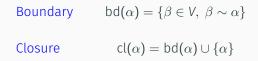
The graph $G_0 = (V_0, E_0)$ is said to be a subgraph of G = (V, E) if $V_0 \subseteq V$ and $E_0 \subseteq E$.

The graph $G_0 = (V_0, E_0)$ is said to be a subgraph of G = (V, E) if $V_0 \subseteq V$ and $E_0 \subseteq E$.

> ug1 <- subGraph(c("b","c","d","e"), ug0)
> plot(ug1)

Subgraphs





```
Boundary bd(\alpha) = \{\beta \in V, \beta \sim \alpha\}

Closure cl(\alpha) = bd(\alpha) \cup \{\alpha\}

> adj(object = ug0, "c")

$c

[1] "d" "b"

> closure(object = ug0,set = "c")

[1] "c" "d" "b"
```

A node in an undirected graph is simplicial if its boundary is complete.

A node in an undirected graph is simplicial if its boundary is complete.

```
> is.simplicial(set = "b", object = ug0)
[1] FALSE
> simplicialNodes(object = ug0)
[1] "a" "c" "d" "e"
> connectedComp(g = ug0)
$`1`
[1] "a" "b" "c" "d"
$`2`
[1] "e"
```

- An edge $\alpha_i \sim \alpha_j$ is a chord if the nodes of this edge belong to a cycle $\alpha = \alpha_0 \sim \alpha_1 \sim \ldots \alpha_n = \alpha$ and where $j \notin \{i 1, i + 1\}$
- A graph where all the cycle of length \geq 4 are *chorldless* is called triangulated graph

- An edge $\alpha_i \sim \alpha_j$ is a chord if the nodes of this edge belong to a cycle $\alpha = \alpha_0 \sim \alpha_1 \sim \ldots \alpha_n = \alpha$ and where $j \notin \{i 1, i + 1\}$
- A graph where all the cycle of length \geq 4 are *chorldless* is called triangulated graph
- > is.triangulated(ug0)
 [1] TRUE

Let (A, B, S) be a triplet of subsets of V. (A, B, S) is a decomposition of \mathcal{G} if

- i. (A, B, S) are disjoints and $V = A \cup B \cup S$
- ii. S is complete
- iii. S separates A and B in ${\cal G}$

Let (A, B, S) be a triplet of subsets of V. (A, B, S) is a decomposition of \mathcal{G} if

- i. (A, B, S) are disjoints and $V = A \cup B \cup S$
- ii. S is complete
- iii. S separates A and B in $\mathcal G$

```
> is.decomposition(set = "a", set2 = "d", set3 = c("b","c"), ug0)
[1] FALSE
> ug1<-subGraph(c("b","c","d","a"), ug0)
> is.decomposition(set = "a", set2 = "d", set3 = c("b","c"), ug1)
[1] TRUE
> is.decomposition(set = "a", set2 = c("d","b"), set3 = "c", ug1)
[1] FALSE
```

 $\mathcal{G} = (V, E)$ is called a decomposable if

i. G is complete, i.e; $E = V \times V$.

ii. or it can be decomposed into a decomposable subgraphs.

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i. \mathcal{G} is complete, i.e; $E = V \times V$.

ii. or it can be decomposed into a decomposable subgraphs.

Theorem

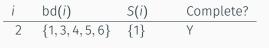
 ${\mathcal G}$ is decomposable if and only if ${\mathcal G}$ is triangulated

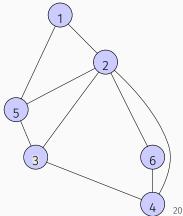
Perfect ordering

Assume $V = \{1, ..., |V|\}$. This is order called **perfect** if $\forall i = 2, ..., |V|$, $S(i) = bd(i) \cap \{1, ..., i - 1\}$ is complete

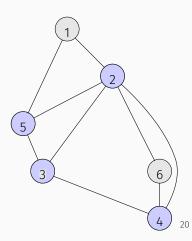
Perfect ordering

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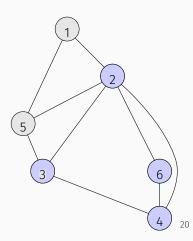




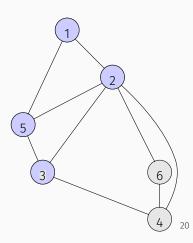
i	bd(i)	S(i)	Complete?
2	$\{1, 3, 4, 5, 6\}$	{1}	Y
3	$\{2, 3, 4, 5\}$	{2}	Y



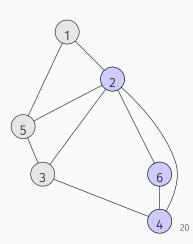
i	bd(i)	S(i)	Complete?
2	$\{1, 3, 4, 5, 6\}$	{1}	Y
3	$\{2, 3, 4, 5\}$	{2}	Υ
4	{2,3,6}	{2,3}	Y

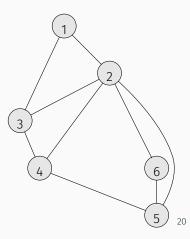


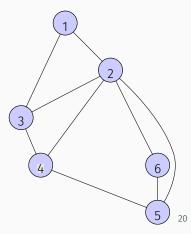
i	bd(i)	S(i)	Complete?
2	$\{1, 3, 4, 5, 6\}$	{1}	Y
3	$\{2, 3, 4, 5\}$	{2}	Υ
4	{2,3,6}	{2,3}	Υ
5	{1,2,3}	{1,2,3}	Ν

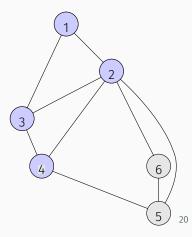


i	bd(i)	S(i)	Complete?
2	$\{1, 3, 4, 5, 6\}$	{1}	Y
3	$\{2, 3, 4, 5\}$	{2}	Y
4	{2,3,6}	{2,3}	Y
5	{1,2,3}	{1,2,3}	Ν
6	{2,4}	{2,4}	Υ

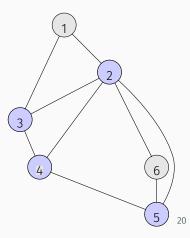




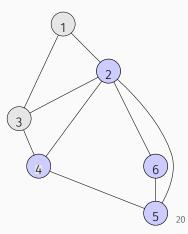




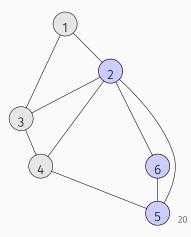
i	bd(i)	S(i)	Complete?
2	$\{1, 3, 4, 5, 6\}$	{1}	Y
3	$\{1, 2, 4\}$	{1,2}	Υ
4	{2,3,5}	{2,3}	Y



i	bd(i)	S(i)	Complete?
2	{1,3,4,5,6}	{1}	Y
3	$\{1, 2, 4\}$	{1,2}	Υ
4	{2,3,5}	{2,3}	Υ
5	{2, 4, 6}	{2}	Y



i	bd(i)	S(i)	Complete?
2	$\{1, 3, 4, 5, 6\}$	{1}	Y
3	$\{1, 2, 4\}$	{1,2}	Υ
4	$\{2, 3, 5\}$	{2,3}	Υ
5	$\{2, 4, 6\}$	{2}	Υ
6	{2,5}	{2,5}	Υ



Perfect ordering and decomposability

• If *G* is decomposable, then the perfect ordering can be obtained using the *maximum cardinality search* algorithm

• If *G* is decomposable, then the perfect ordering can be obtained using the *maximum cardinality search* algorithm

Theorem

 \mathcal{G} is decomposable if and only if \mathcal{G} is triangulated if and only if the vertices of \mathcal{G} admit a perfect ordering.

• If *G* is decomposable, then the perfect ordering can be obtained using the *maximum cardinality search* algorithm

Theorem

 \mathcal{G} is decomposable if and only if \mathcal{G} is triangulated if and only if the vertices of \mathcal{G} admit a perfect ordering.

```
> g2<-ug(~1*2*5,~2*5*3,~2*6*4,~2*4*3)
> mcs(g2)
[1] "1" "2" "5" "3" "4" "6"
```

RIP ordering for the cliques

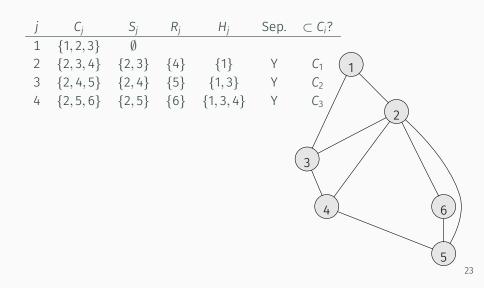
- Let $\{C_1, \ldots, C_p\}$ be the set of cliques for \mathcal{G}
- A Running Intersection Property of $\{C_1, \ldots, C_p\}$ means that for all $j = 2, \ldots, p, \exists i < j$ such that

 $C_j \cap (C_1 \cup \ldots \cup C_{j-1}) \subset C_i$

•
$$S_1 = \emptyset$$
, $S_2 = C_2 \cap C_1$, $S_3 = C_3 \cap (C_1 \cup C_2)$, ...,
 $S_p = C_p \cap (C_1 \cup \ldots \cup C_{p-1})$

- $R_1 = C_1, R_2 = C_2 \setminus S_2, \ldots, R_j = C_j \setminus S_j, \ldots, R_p = C_p \setminus S_p.$
- $S_2 = C_2 \cap C_1$ separates R_2 from $H_2 = C_1 \setminus S_2$
- $\forall j \geq 2$, S_j separates R_j from $H_j = (C_1 \cup \ldots \cup C_{j-1}) \setminus S_j$.

RIP ordering for the cliques



RIP ordering for the cliques

- + If ${\mathcal{G}}$ is triangulated RIP ordering exists (iff)
- $\exists i < j \text{ such } S_j \subset C_i, C_i \text{ is called the parent and the } S_j \text{ are called separators.}$

```
> g1<-ug(~1*2*3,~2*3*4,~2*4*5,~2*6*5)
> rip(g1)
cliques
 1:231
 2:234
 3:254
 4:256
separators
 1 :
 2:23
 3:24
 4:25
parents
 1:0
 2:1
 3:2
 4:3
```

.

.

- $\begin{array}{ccc} \mathcal{G} & \Rightarrow & \text{Cliques} \\ & & & \text{Separators} \end{array}$
 - ${\cal G}$ decomposable = Triangulated Only chordless cycles
 - ${\cal G}$ decomposable = Perfect ordering for vertices RIP ordering for cliques

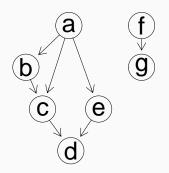
Directed Acyclic Graphs

A Directed Acyclic Graph

- $\vec{\mathcal{G}} = (V, E),$ if $\alpha\beta \in E$ then $\beta\alpha \notin E.$
- edges= arrows
- there's no cycles acyclics: arrows pointing in the same direction all the way around.

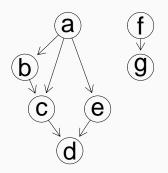
A Directed Acyclic Graph

- $\vec{\mathcal{G}} = (V, E),$ if $\alpha\beta \in E$ then $\beta\alpha \notin E.$
- edges= arrows
- there's no cycles acyclics: arrows pointing in the same direction all the way around.
- $V = \{a, b, c, d, e, f, g\}$
- $E = \{ab, ac, ae, bc, cd, ed, fg\}$



• If $\alpha\beta \in E$ then α is the parent of β

- If $\alpha\beta\in E$ then α is the parent of β
- \cdot *a* is the parent of *b*
- in **R**:
 - ~b*a means b is the child of a
 - ~d*c*e means d is the child of c and e



```
> dag0 <- dag(~a, ~b*a, ~c*a*b, ~d*c*e, ~e*a, ~g*f)
> dag0 <- dag(~a + b*a + c*a*b + d*c*e + e*a + g*f)
> dag0 <- dag(~a + b|a + c|a*b + d|c*e + e|a + g|f)
> dag0 <- dag("a", c("b","a"), c("c","a","b"), c("d","c","e"),
+ c("e","a"),c("g","f"))
> dag0
A graphNEL graph with directed edges
Number of Nodes = 7
Number of Edges = 7
```

```
> dag0a=dag(~a, ~b*a, ~c*a*b, ~d*c*e, ~e*a, ~g*f,
          result="matrix")
+
> dag0a
 abcdegf
a 0 1 1 0 1 0 0
b 0 0 1 0 0 0 0
c 0 0 0 1 0 0 0
d 0 0 0 0 0 0 0
e 0 0 0 1 0 0 0
g 0 0 0 0 0 0 0
f 0 0 0 0 0 1 0
```

- A path (of length *n*) from α to β is a sequence of vertices $\alpha = \alpha_0, \ldots, \alpha_n = \beta_n$ such that $\alpha_{i-1} \to \alpha_i$ is an edge in the graph. If there is a path from α to β we write $\alpha \mapsto \beta$.
- If $\alpha \rightarrow \beta \alpha$ is a parent of β and β is a children of α .

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- If $\alpha \rightarrow \beta \alpha$ is a parent of β and β is a children of α .

```
> parents("d",dag0)
[1] "c" "e"
> children("c",dag0)
[1] "d"
```

Ancestrals

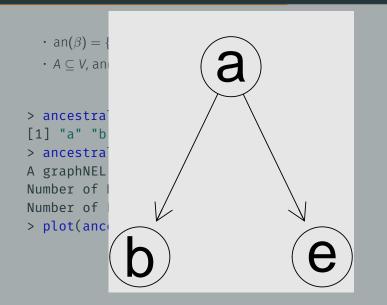
- $\operatorname{an}(\beta) = \{ \alpha \in \mathsf{V} \text{ such } \alpha \mapsto \beta \}$ ancestrors of β
- $A \subseteq V$, $an(A) = \bigcup_{\beta \in A} an(\beta)$ ancestral set of A.

Ancestrals

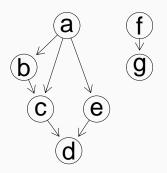
- $\operatorname{an}(\beta) = \{ \alpha \in \mathsf{V} \text{ such } \alpha \mapsto \beta \}$ ancestrors of β
- $A \subseteq V$, $\operatorname{an}(A) = \bigcup_{\beta \in A} \operatorname{an}(\beta)$ ancestral set of A.

```
> ancestralSet(c("b","e"),dag0)
[1] "a" "b" "e"
> ancestralGraph(c("b","e"),dag0)
A graphNEL graph with directed edges
Number of Nodes = 3
Number of Edges = 2
> plot(ancestralGraph(c("b","e"),dag0))
```

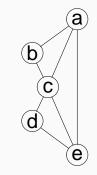
Ancestrals



Moralizing a DAG = Transforming it into an UG (arrows become non-directed) and adding an edge to all parents Moralizing a DAG = Transforming it into an UG (arrows become non-directed) and adding an edge to all parents



Moralizing a DAG = Transforming it into an UG (arrows become non-directed) and adding an edge to all parents





 $\vec{\mathcal{G}} \longleftrightarrow \mathcal{G}_m$

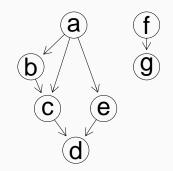
- Let $\alpha \mapsto \beta$ in the DAG $\mathcal{G} = (V, E)$ and $S \subset V$.
- + $\alpha \mapsto \beta$ is active according to S if two following conditions hold:
 - i. every node with converging edges $(\rightarrow \alpha_i)$ is either in S or has a descendant in S,
 - ii. every other node is not in S.
- + $\alpha \mapsto \beta$ is **blocked** by S if it is not active according to \mathcal{G}
- (*A*, *B*, *S*) three disjoint subsets of *V*, *S d*—separate *A* from *B* if for any path from *A* to *B* is blocked by *S*.

d-Separation, Examples

$$A = \{a\}, B = \{d\}, S = \{c, e\}$$

а	\rightarrow	С	\rightarrow	d	blocked by S
а	\rightarrow	е	\rightarrow	d	blocked by S

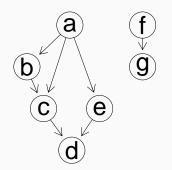
Then S d-separates A and B.



d-Separation, Examples

$$A = \{a\}, B = \{d\}, S = \{c, e\}$$

Then S d-separates A and B.

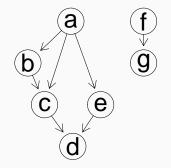


d-Separation, Examples

$$A = \{b\}, B = \{e\}, S = \{c, d\}$$

No paths btw A and B can be blocked by S

Then S doesn't *d*-separate A and B.

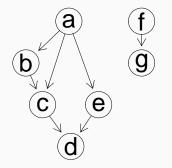


d-Separation, Examples

$$A = \{b\}, B = \{e\}, S = \{c, d\}$$

No paths btw A and B can be blocked by S

Then S doesn't *d*-separate A and B.



> dSep(amat = as(dag0, "matrix"), + first = "b", second = "e",cond = c("c","d"))
[1] FALSE

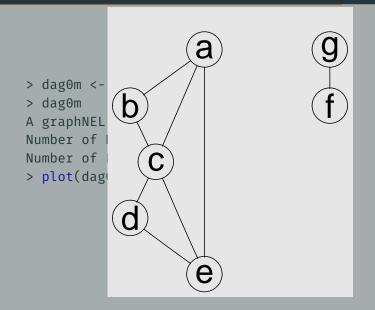
Theorom

Let $\vec{\mathcal{G}}$ be a DAG and \mathcal{G}_m its moral UG associated to $\vec{\mathcal{G}}$.

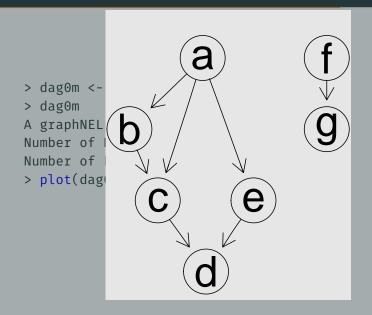
S d-separates A and B if and only if S separates A and B in the sub-graph deduced from \mathcal{G}_m .

```
> dag0m <- moralize(dag0)
> dag0m
A graphNEL graph with undirected edges
Number of Nodes = 7
Number of Edges = 8
> plot(dag0m)
```

Moralization with **R**



Moralization with **R**



Markov properties

Conditional Independence

• $\mathbf{X}_{V} = (X_{v}, v \in V) \sim P$ a random vector ($\in \mathbb{R}^{|V|}$)

• For
$$A \subseteq V$$
, $X_A = (X_v, v \in A)$

- for all A, B, S \subseteq V, A $\perp\!\!\!\perp$ B | S means that $X_A \perp\!\!\!\perp X_B \mid X_S$.
- If f(.) is the generic density

$$A \perp\!\!\!\perp B \mid S \iff f(x_A, x_B \mid x_c) = f(x_A \mid x_S) f(x_B \mid x_S)$$
$$\iff f(x_A, x_B, x_S) = h(x_A, x_S) g(x_B, x_S)$$

Markov properties for UG

 $\mathcal{G} = (V, E)$ is an undirected graph.

(P) We say that P is pairwise Markov w.r.t \mathcal{G} , if

 $\alpha \not\sim_{\mathcal{G}} \beta \Rightarrow \alpha \perp \!\!\!\perp \beta \mid \mathsf{V} \setminus \{\alpha, \beta\}$

(G) We say that P is global Markov w.r.t. $\mathcal{G} = (V, E)$,

S separates A and B in $\mathcal{G} \Rightarrow X_A \perp\!\!\!\perp X_B \mid X_S$

(F) If P has a density f, C is the set of cliques of G, we say that P factorized Markov w.r.t G, then

$$f(x_V) = \prod_{c \in \mathcal{C}} g_c(x_c)$$

Markov properties for UG

 $\mathcal{G} = (V, E)$ is an undirected graph.

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(G) We say th Theorem

if P has a density f, then

(F) If P has a

$$(F) \iff (G) \iff (P)$$

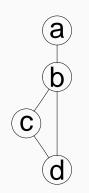
say that P

Xς

factorized markov w.i.e.y, men

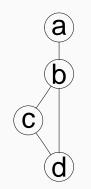
$$f(X_V) = \prod_{c \in \mathcal{C}} g_c(X_c)$$

• a ⊥⊥ c | b



е

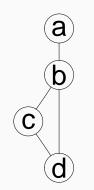
- a ⊥⊥ c | b
- а ⊥⊥ е





е

- a ⊥⊥ c | b
- ∙аше
- a ⊥⊥ d | b, c.





Markov properties for DAGs

 $\vec{\mathcal{G}} = (V, E)$ is a directed acyclic graph.

(Fd) We say that P admits a recusive factorisation according to $\vec{\mathcal{G}}$ if

$$f(x_V) = \prod_{c \in \mathcal{C}} g_c(x_c \mid x_{\mathsf{pa}(c)})$$

(Gd) P obeys to the directed global Markov property w.r.t $\vec{\mathcal{G}}$

$$S d$$
 – separates A and B in $\mathcal{G} \Rightarrow X_A \perp \!\!\!\perp X_B \mid X_S$

(Pd) P obeys to the directed pairwise Markov property w.r.t $\vec{\mathcal{G}}$ if

$$\alpha \not\sim_{\mathcal{G}} \beta \Rightarrow \alpha \perp \beta \mid \mathsf{nd}(\alpha) \setminus \{\beta\}$$

 $nd(\alpha) = V \setminus desc(\alpha)$ where

$$\mathsf{desc}(\alpha) = \{\beta \in \mathsf{V}, \ \alpha \mapsto \beta\}$$

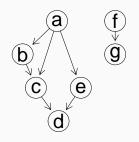
Markov properties for DAGs

 $\vec{\mathcal{G}} = (V, E)$ is a directed acyclic graph.

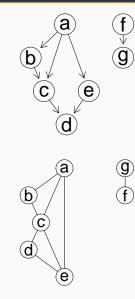
(Fd) We say that P admits a recusive factorisation according to $\vec{\mathcal{G}}$ if

$$f(x_V) = \prod_{c \in C} g_c(x_c \mid x_{\mathsf{pa}(c)})$$

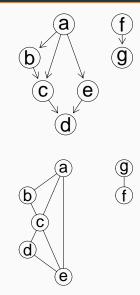
(Gd) P obeys t **Theorem**t $\vec{\mathcal{G}}$ if P has a density f, then $(Pd) \text{ P obeys t} \quad (Fd) \iff (Gd) \iff (Pd)$ $\mu.r.t \vec{\mathcal{G}} \text{ if}$ $\alpha \not\sim_{\mathcal{G}} \beta \Rightarrow \alpha \perp \beta \mid nd(\alpha) \setminus \{\beta\}$ $nd(\alpha) = V \setminus desc(\alpha) \text{ where}$ $desc(\alpha) = \{\beta \in V, \ \alpha \mapsto \beta\}$



Examples

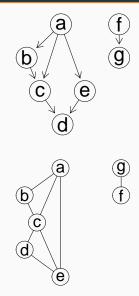


Examples



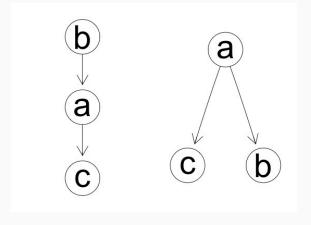
• a ⊥⊥ d | {b, c, e}

Examples



- a ⊥⊥ d | {b, c, e}
- $\cdot b \perp d \mid \{a, c, e\}$

Markov equivalence



b ⊥⊥ c | a

•
$$\mathbf{X}_V = (X_V, v \in V) \sim P$$
 a random vector ($\in \mathbb{R}^{|V|}$):

 $\mathcal{I}(P) = \{ (A, B, S) \subset V \text{ such that } A \perp\!\!\!\perp B \mid S \}$

• $\mathbf{X}_{V} = (X_{v}, v \in V) \sim P$ a random vector $(\in \mathbb{R}^{|V|})$:

 $\mathcal{I}(P) = \{ (A, B, S) \subset V \text{ such that } A \perp\!\!\!\perp B \mid S \}$

• If UG,
$$\mathcal{G} = (V, E)$$
:

 $\mathcal{S}(G) = \{(A, B, S) \subset V \text{ such that } S \text{ separates } A \text{ and } B \text{ in } \mathcal{G}\}$

• (P, \mathcal{G}) is a Graphical Model then $\mathcal{S}(\mathcal{G}) \subseteq \mathcal{I}(P)$

• $X_V = (X_V, v \in V) \sim P$ a random vector ($\in \mathbb{R}^{|V|}$):

 $\mathcal{I}(P) = \{(A, B, S) \subset V \text{ such that } A \perp\!\!\!\perp B \mid S\}$

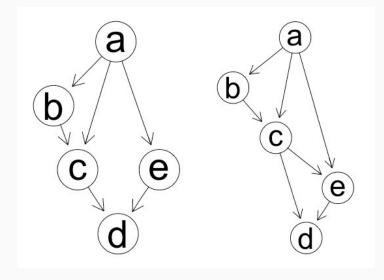
• If DAG, $\vec{\mathcal{G}} = (V, E)$:

 $S(\vec{\mathcal{G}}) = \{(A, B, S) \subset V \text{ such that } Sd - \text{separates } A \text{ and } B \text{ in } \mathcal{G}\}$

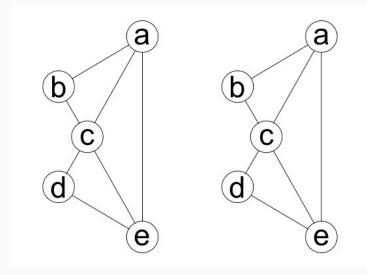
- $(P, \vec{\mathcal{G}})$ is a Graphical Model then $\mathcal{S}(\vec{\mathcal{G}}) \subseteq \mathcal{I}(P)$
- \cdot Two DAG \mathcal{G}_1 and \mathcal{G}_2 are Markov Equivalence if

$$\mathcal{S}(\vec{\mathcal{G}_1}) = \mathcal{S}(\vec{\mathcal{G}_2})$$

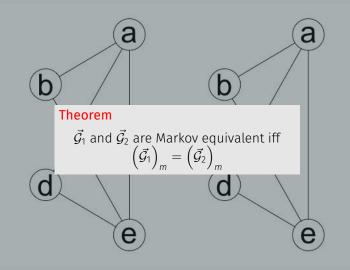
Example, Markov equivalence



Example, Markov equivalence



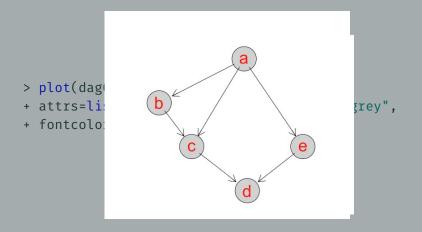
Example, Markov equivalence



Visualizing graphs with R

- > plot(dag0,
- + attrs=list(node = list(fillcolor="lightgrey",
- + fontcolor="red")))

Example 1, Customizing the graph



Example 1, Transforming the graph en LTEX

```
> library(tikzDevice)
> tikz("g.tex",standAlone = T)
> plot(dag0,
+ attrs=list(node = list(fillcolor="lightgrey",
+ fontcolor="red")))
> dev.off()
```

Example 1, Transforming the graph en LTEX

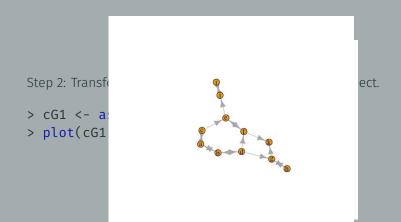
```
> library(tikzDevice)
> tikz("g.tex",standAlone = T)
> plot(dag0,
+ attrs=list(node = list(fillcolor="lightgrey",
+ fontcolor="red")))
> dev.off()
% Created by tikzDevice version 0.10.1 on 2017-03-31 14:50:06
% !TEX encoding = UTF-8 Unicode
\documentclass[10pt]{article}
\usepackage{tikz}
\usepackage[active.tightpage.psfixbb]{preview}
\PreviewEnvironment{pgfpicture}
\setlength\PreviewBorder{0pt}
\begin{document}
\begin{tikzpicture}[x=1pt.v=1pt]
\definecolor{fillColor}{RGB}{255,255,255}
\path[use as bounding box,fill=fillColor,fill opacity=0.00] (0,0) rectangle (505.89,505.89);
\begin{scope}
```

Step 1: Construct the adjacency matrix

```
> d1 <- matrix(0,11,11)</pre>
> d1[1,2] <- d1[2,1] <- d1[1,3] <- d1[3,1] <- d1[2,4] <- d1[4,2] <-
+ d1[5,6] <- d1[6,5] <- 1
> d1[9,10] <- d1[10,9] <- d1[7,8] <- d1[8,7] <- d1[3,5] <-
+ d1[5,10] <- d1[4,6] <- d1[4,7] <- 1
> d1[6,11] <- d1[7,11] <- 1
> rownames(d1) <- colnames(d1) <- letters[1:11]</pre>
> d1
  abcdefghijk
a 0 1 1 0 0 0 0 0 0 0 0
b 1 0 0 1 0 0 0 0 0 0 0
c 1 0 0 0 1 0 0 0 0 0 0
d 0 1 0 0 0 1 1 0 0 0 0
e 0 0 0 0 0 1 0 0 0 1 0
f 0 0 0 0 1 0 0 0 0 1
g 0 0 0 0 0 0 0 1 0 0 1
h 0 0 0 0 0 0 1 0 0 0 0
i 0 0 0 0 0 0 0 0 0 1 0
j00000000100
k 0 0 0 0 0 0 0 0 0 0 0 0
```

Step 2: Transform the adjacency matrix into igraph object.

- > cG1 <- as(d1, "igraph")</pre>
- > plot(cG1)



Step 3: Changing the type of the edges (only directed and undirected edges

```
> E(cG1)
+ 18/18 edges (vertex names):
[1] a->b a->c b->a b->d c->a c->e d->b d->f d->g e->f e->j f->e f->k g->h
[15] g->k h->g i->j j->i
> is.mutual(cG1) ## checks the reciproc pair of the supplied edges
[1] TRUE TRUE TRUE TRUE TRUE FALSE TRUE FALSE TRUE FALSE
[12] TRUE FALSE TRUE FALSE TRUE TRUE TRUE
> ## Change the bidirected edges to undirected edges
> E(cG1)$arrow.mode <- c(2,0)[1+is.mutual(cG1)]
> plot(cG1, layout=layout.spring)
```

```
Step 3: Chang
edges
> E(cG1)
+ 18/18 edges
[1] a->b a->c
[15] g->k h->g
> is.mutual(cG
[1] TRUE TFA
[12] TRUE FAI
> ## Change tf
> E(cG1)$arrow
> plot(cG1, la
```



nd undirected

e->j f->e f->k g->h

oplied edges _<mark>SE TRUE FALSE</mark>

Step 4: Renaming and reshaping nodes, Recoloring edges according to the type

```
> cG1a <- as(cG1, "graphNEL")</pre>
> nodes(cG1a)
 [1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "i" "k"
> nodes(cG1a) <- c("alpha","theta","tau","beta","pi","upsilon","gamma",</pre>
                     "iota", "phi", "delta", "kappa")
+
> edges <- buildEdgeList(cG1a)</pre>
> for (i in 1:length(edges))
   if (edges[[i]]@attrs$dir=="both") {
     edges[[i]]@attrs$dir <- "none"</pre>
+
     edges[[i]]@attrs$color <- "blue"
   }
+
    if (edges[[i]]@attrs$dir=="forward") {
+
     edges[[i]]@attrs$color <- "red"
+
÷
÷
 nodes <- buildNodeList(cG1a)</pre>
>
   for (i in 1:length(nodes)) {
>
   nodes[[i]]@attrs$fontcolor <- "red"</pre>
+
   nodes[[i]]@attrs$shape <- "ellipse"</pre>
   nodes[[i]]@attrs$fillcolor <- "lightgrey"</pre>
   if (i <= 4) {
   nodes[[i]]@attrs$fillcolor <- "lightblue"</pre>
   nodes[[i]]@attrs$shape <- "box"</pre>
+
+
> cG1al <- agopen(cG1a, edges=edges, nodes=nodes, name="cG1a",</pre>
+ layoutType="neato")
> plot(cG1al)
```

Step 4: Renaming and reshaping nodes, Recoloring edges according to the type > cG1a <- as(cG1. > nodes(cG1a) [1] "a" "b" "c" > nodes(cG1a) <-</pre> iota edges <- buildE for (i in 1:len theta if (edges[[i]] beta edges[[i]]@a edges[[i]]@a alpha kappa if (edges[[i] edges[[i]]@a nodes <- build > for (i in 1:le > nodes[[i]]@att nodes[[i]]@att nodes[[i]]@att if (i <= 4) { nodes[[i]]@att

+ nodes[[i]]@attrs\$shape <- "box"</p>

```
> }
> cG1al <- agopen(cG1a, edges=edges, nodes=nodes, name="cG1a",
+ lavoutType="neato")
```

```
> plot(cG1al)
```

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