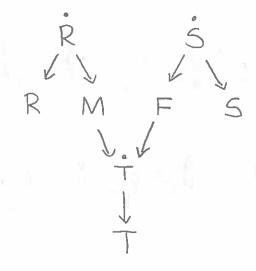
EXACT INFERENCE IN BAYESIAN NETWORKS

D Consider the following Bayesian network for blood types:



where

- R, S, T are the blood genotypes for Rhonda, Sam, and Tim (E { AA, AB, AO, BB, BO, OO}).
 - R, S, T are the blood types for Rhonda, Sam, and Tim (& 2 A, B, AB, O3)
 - M, F are the genes passed to Tim by Rhonda (his Mother) and Sam (his Father), M, F & 2 A, B, 03.
- 2) What if we want to compute the probability of Rhonda having blood type AB, given that we know Tim has blood type A?

EXACT NFERENCE

3) We could try reasoning about it intuitively...

well, if Tim has type A,
then Rhanda has genetype AA, AB, AO, or BO...
and I guess AA is more likely than AB,
but AB is more Weely than BO...



But it's a bit much. Plus what about harder questions like what is the probability of Rhonda having blood type AB, given we know Tim has blood type A, AND. Tim's cousin on his faller's side has blood type O?

4) We need a method to autemate complicated probability questions.

This problem is called probabilistic inference.

Note that P(R=AB|T=A) = P(R=AB,T=A), so if we P(T=A)

Can answer joint queries like P(R=AB, T=A) and P(T=A)then we can immediately also answer carditional queries 5) On the surface, it doesn't look that hard to solve:

$$P(R=AB,T=A) = \sum \sum \sum \sum \sum \sum P(R=AB,T=A,r,s,t,m,f,s)$$

But this expression has 6.6.6.3.3.4 = 7776 terms. While a computer could handle that, I certainly can't. Plus, as the number of variables in the Bayesian network increases, the number of terms grows exponentially.

6) We can begin by replacing the joint distribution by the factored representation assumed by the Bayesian network.

= ZZZZZP(i)P(s)P(r/i)P(m/i)P(f/s)P(s/s)P(t/m,f)P(t/t)

this is just a this is just a function $g_3(r,r) = P(r|r)$ g(t,m,f) = P(t|m,f)

F(rjt) is just a sum of products:

= $\mathbb{Z} \mathbb{Z} \mathbb{Z} \mathbb{Z} \mathbb{Z} g_{3}(\dot{r}) g_{2}(\dot{s}) g_{3}(r,\dot{r}) g_{4}(m,\dot{r}) g_{5}(f,\dot{s}) g_{6}(s,\dot{s}) g_{7}(\dot{t},m,f) g_{8}(t,\dot{t})$

How can we compute this efficiently?

(8) The first thing to notice is we can arrange the summations in whatever order we want, e.g. $P(r,t) = \sum \sum \sum \sum g_{s}(i)g_{2}(i)g_{3}(r,i)g_{4}(m,i)g_{5}(f,i)g_{6}(s,i)g_{7}(i,m,f)g_{8}(t,t)$ in first thing to notice is we can arrange the summations $P(r,t) = \sum \sum \sum \sum g_{s}(i)g_{2}(i)g_{3}(r,i)g_{4}(m,i)g_{5}(f,i)g_{6}(s,i)g_{7}(i,m,f)g_{8}(t,t)$

9 Naw, we can move the innermost summation inward:

 $= \sum_{n} \sum_{i} \sum_{j} g_{i}(r) g_{2}(s) g_{3}(r,r) g_{4}(m,r) g_{5}(f,s) g_{7}(t,m,f) g_{8}(t,t) \sum_{j} g_{6}(s,s)$

Note that $\sum_{s} g_{6}(s, s)$ is a function of s, so we can define a new function $h_{1}(s) = \sum_{s} g_{6}(s, s)$ and rewrite:

 $P(r,t) = \sum \sum \sum \sum g_1(i)g_2(i)g_3(r,i)g_4(m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_4(i,m,i)g_5(f,i)g_5(i,m$

We have "eliminated" the summation over variable 5 (at least, we're hiding the fact that it exists.)

1 We can continue to do this for each summartion:

$$P(r,t) = \sum_{m} \sum_{f} \sum_{g} (r) g_{3}(r,r) g_{4}(m,r) g_{7}(t,m,f) g_{8}(t,t) \sum_{g} g_{2}(s) g_{5}(f,s) h_{1}(s)$$

=
$$\sum \sum \sum g_1(r)g_3(r,r)g_4(m,r)g_7(t,m,f)g_8(t,t)h_2(f)$$

[letting
$$h_2(f) = \sum_{s} g_2(s)g_s(f,s)h_s(s)$$

=
$$\sum \sum g_{3}(\dot{t}, m, f)g_{8}(t, \dot{t})h_{2}(f) \sum_{\dot{r}} g_{1}(\dot{r})g_{3}(r, \dot{r})g_{4}(m, \dot{r})$$

$$= \sum_{m} \sum_{f} \sum_{f} g_{7}(\dot{t}, m, f) g_{8}(\dot{t}, \dot{t}) h_{2}(f) h_{3}(r, m)$$
[letting $h_{3}(r, m) = \sum_{f} g_{7}(\dot{r}) g_{3}(r, \dot{r}) g_{4}(m, \dot{r})$]
$$= \sum_{f} \sum_{f} \sum_{f} g_{7}(\dot{t}, m, f) g_{8}(\dot{t}, \dot{t}) h_{2}(f) h_{3}(r, m) = \sum_{f} g_{7}(\dot{r}) g_{3}(r, \dot{r}) g_{4}(m, \dot{r})$$

$$= \sum_{m} \sum_{f} h_{2}(f) h_{3}(r,m) \sum_{\dot{t}} g_{7}(\dot{t},m,f) g_{8}(t,\dot{t})$$

=
$$\sum_{m \neq 1} h_2(f) h_3(r,m) h_4(t,m,f)$$

[letting
$$h_4(t,m,f) = \sum_{\dot{t}} g_{\pm}(\dot{t},m,f) g_{8}(\dot{t},\dot{t})$$
]

$$= \sum_{m} h_{3}(r,m) \sum_{f} h_{2}(f) h_{4}(t,m,f)$$

$$= \sum_{m} h_3(r,m) h_5(t,m)$$

[letting
$$h_{6}(r,t) = \sum_{m} h_{3}(r,m)h_{5}(t,m)$$
]

1) What have we don? Well, we have laid out a strategy for computing P(r,t).

num operations

Num operations -745, compute $h_1(\dot{s}) = \sum_{s} g_{s}(s,\dot{s})$ $|\dot{s}| \cdot |\dot{s}|$

- $\forall f$, compute $h_2(f) = \sum_{\hat{s}} g_2(\hat{s})g_5(f,\hat{s})h_1(\hat{s})$ |5|. |F|

- Vr,m, compute h3 (r,m) = \(\frac{1}{2} \, \text{g}(\vec{r}) \, \text{g}(\vec{r}) \, \text{g}(\vec{r}) \, \text{g}(\vec{m}, \vec{r}) \)

- $\forall t, m, F, compute h_{4}(t, m, f) = \sum_{t} g_{+}(t, m, f) g_{8}(t, t)$ |T|.|T|.|M|.|F|

- Yt, m, compute h5(t,m) = \(\Sigma h_2(f) h_4(t,m,f) \) |F|. |T|. |M|

- $\forall r, t$, compute $h_c(r, t) = \sum_m h_3(r, m) h_5(t, m)$ $|M| \cdot |R| \cdot |T|$

Then return P(r,t) = ho (r,t).

Notice that:

ISI. ISI + ISI. IFI+ IRI. IRI. IMI + ITI. ITI. IMI. IFI+ IFI. ITI. IMI + IMI. IKI. ITI
is much better than directly computing the sun of products;
which is:

|R|.|R|.|S|.|S|.|T|.|T|.|M|.|F|

(414 operations, vesus 124416 operations)

12) If we execute this strategy, then we can get the probability that Rhonda has blood type AB and her son has type A. O

	1	
h,:	5	h. (s)
	Α̈́Α	
	A'A AB	1
	Ao	
	BB	1
\$ 11	BO	
	00	

$$h_2$$
: $f | h_2(f)$
 $A | \frac{1}{3}$
 $B | \frac{1}{3}$
 $O | \frac{1}{3}$

$$h_3$$
: $r m | h_3(r,m)$

AB A $\frac{1}{12}$

AB B $\frac{1}{12}$

AB O O

$$h_5$$
: $t m h_5(t,m)$
 $A A \frac{2}{3}$
 $A B O \frac{1}{3}$

$$h_{\omega}$$
: $r + h_{\omega}(r, t)$
AB A $\frac{1}{18}$

$$P(R=AB_{3}T=A) = \frac{1}{18}$$

We can use the same technique to compute $P(T=A) = \frac{1}{3}$. This gives us the probability that Rhanda's blood type is AB given Tim's blood type is A:

$$P(R=AB|T=A) = \frac{P(R=AB,T=A)}{P(T=A)} = \frac{1}{6}$$

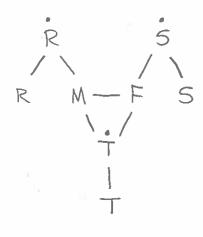
(13) To analyze the running time of this "variable elimination" algorithm, let d be the maximum size of any variable domain.

Compute all values o	E	num ops	
_ h, (s)		15/· s/ \le d2	< d+
$h_2(f)$		5 . F \ d2	< d4
$h_3(r,m)$		1R - 1R - M = d3	< d4
$h_4(t,m,f)$		T - T - M - F ≤ d'	t < 24
hs (t, m)		F . T . M = d3	< d+
$h_{G}(r,t)$		$ M \cdot R \cdot T \leq d^3$	5d4
		in.	$\leq n \cdot d^4$
			1
			number of variables
			ot variables

⁽H) If we let w be the maximum number of variables in our h-functions; then the running time is $O(n \cdot d^{w+1})$

This quantity w is called the width of the elimination order.

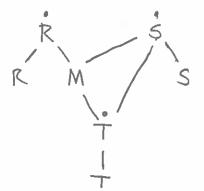
(15) It can be convenient to view this graphically. Create an undirected graph whose vertices are the variables. There is an edge between two vertices iff they appear in the same g-function:



 $P(r|t) = \sum \sum \sum \sum \sum g_{s}(r)g_{s}(s)g_{s}(r,r)g_{4}(m,r)g_{5}(f,s)g_{6}(s,s)g_{7}(t,m,f)g_{8}(t,t)$

(i) Suppose we choose to first eliminate F: $P(r|E) = \sum \sum \sum \sum g_{s}(i)g_{z}(i)g_{z}(i)g_{3}(r,i)g_{4}(m,i)g_{6}(s,i)g_{8}(t,i)h_{s}(m,i,i)$ $m \neq i \neq s \neq s$ where $h_{s}(m,i,i,i) = \sum g_{s}(F,i,i)g_{4}(E,i,i)$

We create an h-function over m, s, t. Graphically, these are all the nodes that F was connected to. The new graph becomes:



17) Note that "eliminating" F corresponds (graphically) to removing F and connecting all the nodes it was Connected to. The size of this new clique corresponds to the number of variables in the new h-function:

R M S 1-1-1 h, (m, s,t)

(19) Therefore, we can compute the width of an elimination order as a graph algorithm:

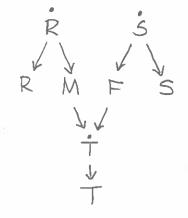
> " let G = (V, E), where V carresponds to the variables and two vertices are connected if they co-occur in a g-function.

-> Herate through Vising Vn:

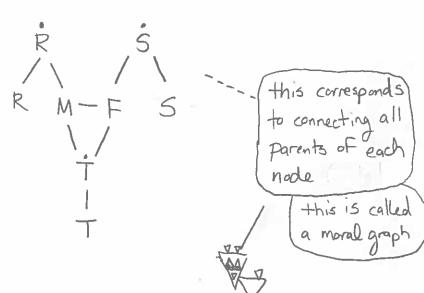
→ w = max (w, num vertices Vi is connected to)

→ remove Vi from G and connect all its neighbors return w

- 19) So given a Bayesian network, we can predict the runtime of probabilistic interence based solely on its graph structure, and the domain size of its variables.
 - (i) Take the network:



(ii) Create an undirected
graph where two vars
are connected if they
co-appear in a
conditional probability
function



- (iii) Compute the width w of your selected elimination order (as described in (B))
- (iv) The runtime of variable elimination is O(n.dwi), where n is the number of variables, and d is the maximum domain size.

20) This means that certain Bayesian networks enjoy good performance, like the following:

$$H_1 \rightarrow H_2 \rightarrow H_3 \rightarrow \cdots \rightarrow H_K$$

$$\downarrow \qquad \qquad \downarrow \qquad \qquad \downarrow$$

$$O_1 \qquad O_2 \qquad O_3 \qquad O_K$$

which is Sometimes called a Hidden Markov Model and has an elimination order of width 1 (i.e. O, H, O2, H2...)

Thus the runtime of variable elimination is $O(k \cdot d^2)$.

If the variables are tall binary (or constant-size domain), then this is O(k).

21) Unfortunately, others have guaranteed poor performance, like this grid network:

whose best width is O(m), thus the runtime of VE is O(m2.dm+1)