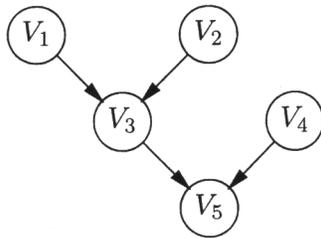


Probabilistic Reasoning (INFOPROB)

6 January 2006

Question 1

Consider a probabilistic network $B = (G, \Gamma)$, where $G = (V(G), A(G))$ is the following acyclic digraph and $\Gamma = \{\gamma_{V_i} \mid V_i \in V(G)\}$ is given by:



$$\begin{aligned} \gamma(v_1) &= 0.25 & \gamma(v_2) &= 0.6 \\ \gamma(v_3 \mid v_1 \wedge v_2) &= 0.2 & \gamma(v_4) &= 0.3 \\ \gamma(v_3 \mid \neg v_1 \wedge v_2) &= 0.1 & \gamma(v_5 \mid v_3 \wedge \neg v_4) &= 0.5 \\ \gamma(v_3 \mid v_1 \wedge \neg v_2) &= 0.7 & \gamma(v_5 \mid \neg v_3 \wedge v_4) &= 0.8 \\ \gamma(v_3 \mid \neg v_1 \wedge \neg v_2) &= 0.35 & & \end{aligned}$$

- a) Assume that the variables V_3 and V_4 exhibit a disjunctive interaction with respect to variable V_5 and that for specifying the assessment function for node V_5 , the noisy-or gate is used. Complete the assessment function γ_{V_5} for node V_5 . *(10 points)*

Let \Pr be the probability distribution defined by probabilistic network B . Consider the five computation rules of *Pearl's* data fusion algorithm.

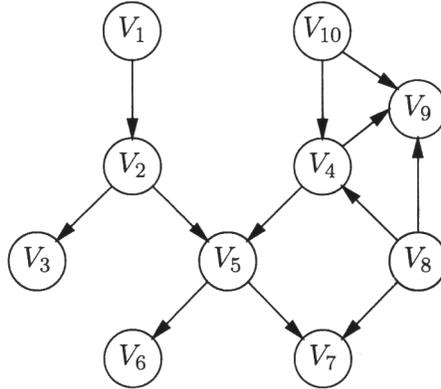
- b) Assume that the observation $V_3 = \text{true}$ is entered into probabilistic network B . Illustrate *Pearl's* algorithm by computing the probability $\Pr^{v_3}(v_1)$. Clearly indicate which messages are passed and how they are computed; explicitly mention all assumptions you make. *(15 points)*

Now use *Pearl's* algorithm as a “black box”: $\text{Pearl}(B, \tilde{c}_E) \rightarrow \Pr^{\tilde{c}_E}(V_i)$ taking as input network B and a partial configuration \tilde{c}_E of observations and returning for each node the probabilities of all its values, given the observations.

- c) Explain how the probability $\Pr^{v_3}(v_1 \vee \neg v_4)$ can be efficiently computed from the network. *(10 points)*

Question 2

Consider a probabilistic network $B = (G, \Gamma)$, where $G = (V(G), A(G))$ is the following acyclic digraph with > 6 loops:



- a) Give a loop cutset for graph G that can be found by applying the heuristic *Suermondt & Cooper* algorithm. (10 points)
- b) Consider a set $C \subseteq V(G)$. Let G' be the graph that results from G by removing all *outgoing* arcs of all nodes in C , that is, $G' = (V(G), A'(G))$ with $A'(G) = A(G) \setminus \{(V_i, V_j) | V_i \in C\}$. Similarly, let G'' be a graph that results from G by removing all nodes in C together with their incident arcs, that is, $G'' = (V(G) \setminus C, A''(G))$ with $A''(G) = A(G) \setminus \{(V_i, V_j) | V_i \in C \text{ or } V_j \in C\} = A'(G) \setminus \{(V_i, V_j) | V_j \in C\}$.

A well-known property of G' is that if G' is a singly connected graph then set C is a loop cutset for G . Prove, or give a counter-example for, the following similar statement:

if G'' is a singly connected graph then set C is a loop cutset for G .

(15 points)

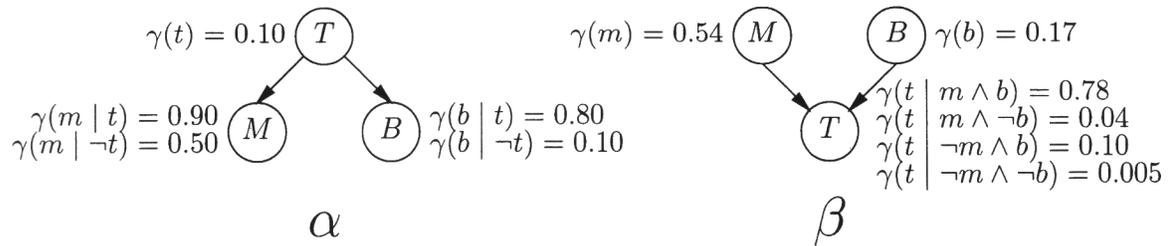
- c) Suppose we subject network B to a one-way sensitivity analysis, where we are interested in the probability distribution $\Pr(V_5)$. More specifically, we are interested in the *most likely value* of variable V_5 and how this changes upon parameter variation. Given that there are no observations in the network and assuming that all variables in the network can only adopt two values, a one-way sensitivity analysis results in 116 sensitivity functions describing the effect of varying each of the 58 parameter probabilities on each of the two output values of V_5 .

It is possible to use these sensitivity functions to establish whether or not an *observation* for one of the variables in the network could change the most likely value of variable V_5 ? Clearly motivate your answer. (10 points)

Question 3

Consider modelling the disease tuberculosis, together with its symptoms and the different tests that are used to diagnose the disease, in a probabilistic network. We focus on the variable T , with possible values t and $\neg t$ that describe whether or not a patient has tuberculosis, and two test variables M and B with values m and $\neg m$, and b and $\neg b$, respectively. Variable M models the outcome of a Mantoux test and variable B models the outcome of a bloodtest.

Consider the following two possible probabilistic networks:



a) Clearly explain which of the following statements is or are correct:

- A. Network α cannot diagnose the patient, but can only predict the outcome of the two tests M and B for patients for whom we know whether or not they have tuberculosis (T).
- B. Network β needs the outcome of both tests B and M to diagnose whether or not a patient has tuberculosis (T).
- C. Networks α and β can both predict whether or not a patient has tuberculosis based upon ≤ 0 test-results; both networks can in addition predict the outcome of the two tests M and B for patients for whom we know for sure whether or not they have tuberculosis (T).

Observation: the two networks do not capture the same independence relation: in network α we have for example that $I(\{M\}, \{T\}, \{B\})$ and $\neg I(\{M\}, \emptyset, \{B\})$, whereas in network β we find that $\neg I(\{M\}, \{T\}, \{B\})$ and $I(\{M\}, \emptyset, \{B\})$.

Argument: if the test results are somewhat reliable, then, for example, a positive outcome for test M would make a positive outcome for test B more likely.

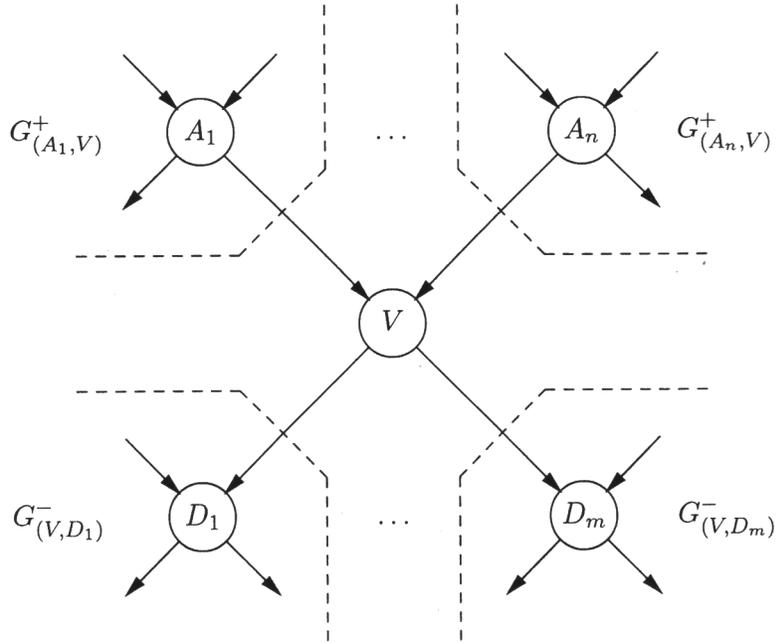
b) Assume that you agree with *Argument*, would you then prefer network α or network β ? Explain your answer.

Suppose that we change the network by summarising the two testresults in an additional intermediate variable I in such a way that $\Pr(TMB)$ remains unchanged. This intermediate variable is placed between the disease variable and the two test variables, resulting in two new networks α' and β' , with $A(G_{\alpha'}) = \{T \rightarrow I, I \rightarrow M, I \rightarrow B\}$ and $A(G_{\beta'}) = \{M \rightarrow I, B \rightarrow I, I \rightarrow T\}$.

c) For your network of preference from part b), are the (in)dependences among the variables T , M and B stated in *Observation* above still valid in the extended network? If not, would you prefer the simple network, or the extended one? Clearly motivate your answer(s).

Formulas

Pearl in a singly connected graph



Consider a node V in a probabilistic network $B = (G, \Gamma)$. Let $\rho(V) = \{A_1, \dots, A_n\}$ be the set of direct ancestors (parents) of V in G , and let $\sigma(V) = \{D_1, \dots, D_m\}$ be the set of its direct descendants (children). With Pearl's algorithm, node V computes the following parameters:

$$\begin{aligned} \pi(V) &= \sum_{c_{\rho(V)}} \left(\gamma(V | c_{\rho(V)}) \cdot \prod_{i=1, \dots, n} \pi_V^{A_i}(c_{A_i}) \right) \\ \lambda(V) &= \prod_{j=1, \dots, m} \lambda_{D_j}^V(V) \\ \pi_{D_j}^V(V) &= \alpha \cdot \pi(V) \cdot \prod_{\substack{k=1, \dots, m \\ k \neq j}} \lambda_{D_k}^V(V) \\ \lambda_V^{A_i}(A_i) &= \alpha \cdot \sum_{c_V} \lambda(c_V) \cdot \sum_{c_{\rho(V) \setminus \{A_i\}}} \left(\gamma(c_V | c_{\rho(V) \setminus \{A_i\}} \wedge A_i) \cdot \prod_{\substack{k=1, \dots, n \\ k \neq i}} \pi_V^{A_k}(c_{A_k}) \right) \end{aligned}$$