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Efficient Calculation of Optimal Decisions in Graphical Models

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Preface

This project report is my Master's thesis, and represents 20 weeks of work. Writing this thesis is the last part of the Master of Science in Industrial Mathematics program at the Norwegian University of Science and Technology. The first years of studies, I tried not to think about the fact that I would have to write a Master's thesis during my tenth and last semester. I have loved being a student, and I have very much enjoyed each and every one of the last five years. Especially, I have loved this Spring semester. Working with this project and this thesis, it turned out to be maybe the best semester of all. I have learned a lot, both of statistical knowledge, but also a lot about writing a report.

First and foremost, I would like to thank my adviser Professor Håkon Tjelme-land for the invaluable cooperation, for each of our many discussions, and for lots of inspiration and constructive feedback. I am very thankful for all the time he has spent on this project.

I would also like to thank "Matteland", and with that, all of my classmates, for creating a great place to work, but also space and time for much needed breaks. Then, I would like to thank the Norwegian Computing Center for hopes and dreams for the future. My mother very much deserves a special thanks for her continuous support, as does my grandmother for being the most original person I know. Also, a big thanks to my dad, for telling his 16 year old daughter that girls are not able to do Mathematics.

At last, I would like to thank Black Sabbath, longboarding, Thin Lizzy, cross country skiing, coffee, unicycling, Led Zeppelin, Rubik's cube and the wonderful spring in Trondheim for making each of my days special.

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Abstract

We present a method for finding the optimal decision on Random Variables in a graphical model. Upper and lower bounds on the exact value for each decision are used to reduce the complexity of the algorithm, while we still ensure that the decision chosen actually represents the exact optimal choice. Since the highest lower bound value is also a lower bound on the value of the optimal decision, we rule out any candidate with an upper bound of lower value than the highest lower bound. By this strategy, we try to reduce the number of candidates to a number we can afford to do exact calculations on.

We generate five Bayesian Networks with corresponding value functions, and apply our strategy to these. The bounds on the values are obtained by use of an available computer program, where the complexity is controlled by an input constant. We study the number of decisions accepted for different values of this input constant. From the first Network, we learn that the bounds does not work well unless we split the calculations into parts for different groups of the nodes. We observe that this splitting works well on the next three Networks, while the last Network illustrates how the method fails when we add more edges to the graph. We realize that our improved strategy is successful on sparse graphs, while the method is unsuccessful when we increase the density of edges among the nodes.

Sammendrag

Vi presenterer en metode for å finne optimal beslutning på tilfeldige variable i grafiske modeller. Øvre og nedre skranke for eksakt verdi blir brukt for å redusere metodens kompleksitet, samtidig som det sørger for at beslutningen som velges til slutt faktisk er den optimale. Da den høyeste nedre skranken også er en nedre skranke for verdien av den optimale beslutningen, kan vi forkaste enhver kandidat med lavere øvre skranke enn den høyeste nedre skranken. Slik prøver vi å redusere antallet kandidater til et antall vi klarer å regne eksakt på.

Vi genererer fem Bayesianske nettverk med tilhørende verdifunksjoner, og tester metoden på disse. Skrankene blir funnet ved hjelp av et tilgjengelig dataprogram, hvor kompleksiteten styres av brukeren. Vi har variert denne, og sett på hvor mange kandidater vi klarer å forkaste. Av det første nettverket ser vi at skrankene fungerer dårlig dersom vi ikke splitter utregningene i deler for ulike grupper av nodene. Fra neste tre nettverkene, ser vi at oppsplittingen fungerer godt, samtidig som det siste nettverket viser hvordan metoden slutter å fungere når vi legger til flere kanter på grafen. Vi ser derfor at resultatet blir bra for grafer med relativt få koblinger mellom nodene, men når tettheten av disse øker, ser vi at metoden slutter å fungere.

Contents

1	Introduction	1
2	Bayesian Networks	2
2.1	Mathematical Results	5
2.2	Conditional Independence	9
2.3	Markov Random Fields	12
3	Pseudo-Boolean functions	16
3.1	A graph structure for coefficients	19
3.2	Converting between exponential and linear form	21
3.3	Probability Distribution for a Bayesian Network	21
3.3.1	An example Network	23
4	Approximate Forward-Backward Algorithm	25
5	Mathematical Formulation of the Problem	29
6	Mathematical Formulation of the Solution	32
7	Generating Bayesian Networks	37
7.1	The Directed Acyclic Graph	38
7.2	The Joint Probability Distribution	40
7.3	The Cost and Income functions	41
8	First Set of Tests: BN1	43
9	Second Set of Tests	47
9.1	Sparse graphs: BN2-BN4	51
9.1.1	BN2	52
9.1.2	BN3	59
9.1.3	BN4	65
9.2	A less sparse graph: BN5	72
10	Some Analysis on Complexity	77
11	Closing Remarks	80
A	Implementation of Graph Structure	84

B	A short User Oriented Introduction to "The AMRF Programs"	85
B.1	A pseudo-Boolean function as input	86
B.2	Calculating an approximate normalizing constant	86
B.3	Calculating bounds for the normalizing constant	87
B.4	An approximate Viterbi algorithm	88
C	Probability Distribution, Cost and Income Functions for Section 8 and Section 9	89
C.1	Distribution, Cost and Income functions BN1	89
C.2	Distribution, Cost and Income functions BN2	92
C.3	Distribution, Cost and Income functions BN3	101
C.4	Distribution, Cost and Income functions BN4	110
C.5	Distribution, Cost and Income functions BN5	123

1 Introduction

Continuously, we try our best to find the best decision for something. Usually, we want to "the right thing", but even with the best intentions, our decisions can lead to outcomes we don't like. This is because we usually don't control all surroundings that will influence the result of our decision, when set into action. That means that we don't know for sure what will be the end result of acting according to some decision. Then, we want to reason about how good a decision is on average. To handle this problem, we set up a probabilistic model for the surroundings, and associate Random Variables to the sources of influence to our end result. On these Random Variables, we construct a Joint Probability Distribution.

The Joint Probability Distribution associates a probability to each event concerning the surroundings, that is, some number telling how often we would expect this collection of criteria on the surroundings to be true. But the Joint Probability Distribution also contains more information about how the Random Variables depend on each other in a probabilistic setting. So, when setting up such a probabilistic model, we have to be able to encode all such information. In general, a causal model is both the easiest to set up, and the easiest to work with after. That means setting up a marginal probability distribution for each Random Variable to depend on the other Random Variables having a direct influence on the first one, and merging all these marginals to the Joint Probability Distribution.

To visualize these dependencies, it is usually convenient to set it all up in a graphical model. That means drawing a circle to represent each Random Variable, and put arrows between them to encode dependencies. In a directed graphical model, the arrows have a direction, and correspondingly, in an undirected graphical model, arrows are undirected. For the directed causal model, we would set an arrow to point from a Random Variable to another if the first one has direct influence on the other. After specifying the graphical model, that is, the circles and arrows, the graphical model encodes some structure on the Joint Probability Distribution. This structure usually also makes it easier to specify the numbers we need to have the full Joint Probability Distribution.

Given the graphical probabilistic model, we can start reasoning about the effect of our decision; what could happen, and how often it would. To figure out what is actually the best decision, we have to be able to associate a value to each realization. That is, a combination of our decision and an assignment to the unknown surroundings should correspond to some number representing the value. These values are then a function depending both on our deterministic

decision and several unknown Random Variables. And then, with our probabilistic graphical model in hand, we can find the expected value of a decision. Comparing the expected values for different decisions, we choose the decision with the highest expected value, and call this the optimal decision.

As an example of application, we have a search for hydrocarbons, as in Martinelli et al. (2011). The Random Variables represent different places where hydrocarbons might be present. The structure of the graph is specified by where hydrocarbons can be generated, and where it can flow. That is, the probabilistic model and the Joint Probability Distribution is set up by looking at some geological data. We would like to expect as high a gross income as possible, and our decision would be which areas to check. The value function would represent the gross income of a realization, that is, the sum over incomes for findings, minus both development costs for the findings and costs for checking all of the areas we decided to.

The general problem described above is exactly what we try to solve in this report. First, in Section 2, we introduce theory about such probabilistic models, where we focus on Bayesian Networks, which is a type of directed graphical models. In Section 3, we present the representation of functions that will be used throughout the report, and also, in Section 4, we present an algorithm that will be used when we calculate the expected value. Then, we give a mathematical formulation of our problem in Section 5, and a mathematical formulation of the solution is given in Section 6. Sections 7, 8 and 9 present information and the actual results from tests of our solution method, and Section 10 discusses the complexity of the algorithms. Finally, Section 11 provides some closing remarks.

2 Bayesian Networks

From Cormen et al. (2009), we have the following definition of a directed graph.

Definition 2.1. *A directed graph G is a pair (V, E) , where V is a finite set and E is a binary relation on V . The set V is called the vertex set of G , and its elements are called vertices or nodes. The set E is called the edge set of G , and its elements are called edges.*

We will refer to the elements in V as the nodes in G , and represent them as circles in our figures. As described in Definition 2.1, the elements in E are ordered pairs of nodes. That is, if $e = (X_i, X_j) \in E$, there is an edge e from node X_i to node X_j . This edge e will be represented by an arrow from node X_i to node X_j in our figures. Correspondingly, an undirected graph has

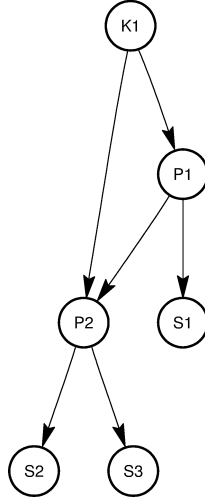


Figure 1: An example graph $G = (V, E)$. All graphs in this report are drawn by use of the MATLAB Biograph tool.

undirected edges. That is, the elements in E are unordered pairs of nodes, and can therefore be viewed as if there was a directed edge in both directions. From now on, when the term graph is used, we assume that the graph is a directed one, unless otherwise specified.

A visualization of an example graph can be seen in Figure 1. Observe that in this graph,

$$V = \{ K_1, P_1, P_2, S_1, S_2, S_3 \}$$

and

$$E = \{ (K_1, P_1), (K_1, P_2), (P_1, P_2), (P_1, S_1), (P_2, S_2), (P_2, S_3) \}.$$

A Bayesian Network is a way to express conditional independence assumptions among a set of Random Variables by use of a graph. Each Random Variable is represented as a node X_i , and arrows between the nodes encode a possible conditional dependence relationship. In this discussion, we will not distinguish between the Random Variable and its corresponding node. In graph theory the terminology for nodes at the different end points of an edge is as follows.

Definition 2.2. *If there is an edge $e = (X_j, X_k)$ from node X_j to node X_k , node X_j is a parent of node X_k , and node X_k is a child of node X_j .*

If there are no edges e from any node to node X_j , we say that X_j is a root node. If there are no edges e to any node from node X_j , we say that X_j is a leaf node. Also, a path is a list of edges, such that the end node at each edge is the start node of the next. Then, the path describes a way to traverse the graph from the first start node to the last end node by just following the directed edges in the graph. If there is a path along the directed edges $E = \{e_i\}_i$ from node X_j to node X_k , we say that X_j is an ancestor of X_k , and X_k is a descendant of X_j . Letting 2^V denote the power set of V , this introduces the following functions.

$$\begin{aligned} \text{Pa} : V &\rightarrow 2^V & \text{such that} & & \text{Pa}(X_i) &= \{X_k \in V \mid (X_k, X_i) \in E\} \\ \text{Ch} : V &\rightarrow 2^V & \text{such that} & & \text{Ch}(X_i) &= \{X_k \in V \mid (X_i, X_k) \in E\} \end{aligned}$$

That is, $\text{Pa}(X_i)$ is the set of nodes X_k such that X_k is a parent of X_i and $\text{Ch}(X_i)$ is the set of nodes X_k such that X_k is a child of X_i . Correspondingly, we also let $\text{Anc}(X_i)$ denote the set of ancestors of X_i and $\text{Des}(X_i)$ denote the set of descendants of X_i .

With some graph terminology in hand, we present the following definition of a Bayesian Network from Russell and Norvig (2003).

Definition 2.3. *A Bayesian Network is a graph, consisting of a set of nodes $V = \{X_i\}_{i=1}^n$ and a set of directed edges $E = \{e_i\}_{i=1}^{n_e}$ between pairs of the nodes. It is required that the graph has no directed cycles, i.e. it is a Directed Acyclic Graph. In addition, each node X_i has a set of Local Probability Distributions $P(X_i | \text{Pa}(X_i))$ associated with it.*

That is, for each assignment of $\text{Pa}(X_i)$, $P(X_i | \text{Pa}(X_i))$ is a probability distribution for the Random Variable X_i .

The nodes and the edges specify what is called the topology of the network. Since the topology of the Bayesian Network constitutes a Directed Acyclic Graph, there is a topological ordering of the nodes. That is, there exists a bijective numbering of the nodes

$$\ell : \{X_i\}_{i=1}^n \rightarrow \{1, \dots, n\}$$

such that for any edge $e = (X_j, X_k)$ in the network, we have $\ell(X_j) < \ell(X_k)$. Also, this means that for any nodes X_j, X_k with $\ell(X_j) < \ell(X_k)$, there is no directed path from X_k to X_j .

Given a Bayesian Network with its topology and the sets of Local Probability Distributions, one uniquely determines the full Joint Probability Distribution over all the Random Variables represented in the network by the formula

$$\mathbb{P}(X_1, \dots, X_n) = \prod_{i=1}^n P(X_i | \text{Pa}(X_i)). \quad (2.1)$$

2.1 Mathematical Results

Notice that, given a subset A_j of the Random Variables, we can divide the probability distribution into

$$\mathbb{P}(X_1, \dots, X_n) = \prod_{X_k \in A_j} P(X_k | \text{Pa}(X_k)) \prod_{X_k \notin A_j} P(X_k | \text{Pa}(X_k)).$$

Note that we have introduced the short hand $P(X_j | \text{Pa}(X_j))$ for the more correct expression $P(X_j = x_j | X_k = x_k \forall X_k \in \text{Pa}(X_j))$. Given a Random Variable X_j , we let $C_j \subseteq \text{Ch}(X_j)$, and extend the definition of the function Anc to sets C_j of nodes such that

$$\text{Anc}(C_j) = \bigcup_{X_k \in C_j} \text{Anc}(X_k).$$

Now, let

$$A_j = C_j \cup \text{Anc}(C_j) \cup \text{Anc}(X_j) \cup \{X_j\}, \quad (2.2)$$

and also

$$A_j^c = \{X_1, \dots, X_n\} \setminus A_j.$$

Notice that if $C_j \neq \emptyset$,

$$\text{Anc}(X_j) \cup \{X_j\} \subseteq \text{Anc}(C_j).$$

We are going to prove that for any assignment of the Random Variables in A_j ,

$$\sum_{X_i \in A_j^c} \prod_{X_k \in A_j^c} P(X_k | \text{Pa}(X_k)) = 1. \quad (2.3)$$

Without loss of generality, assume that the numbering i of the nodes X_i is such that

$$A_j = \{X_1, X_2, \dots, X_{m-1}\}, \quad A_j^c = \{X_m, X_{m+1}, \dots, X_n\},$$

and that X_1, X_2, \dots, X_n is actually a topological ordering of all the Random Variables $\{X_1, \dots, X_n\}$ in the Bayesian Network. Then, given an assignment of the Random Variables in A_j ,

$$\begin{aligned}
\sum_{X_i \in A_j^c} \prod_{X_k \in A_j^c} P(X_k | \text{Pa}(X_k)) &= \sum_{X_m} \cdots \sum_{X_{n-1}} \sum_{X_n} \prod_{X_k \in A_j^c} P(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_m} \cdots \sum_{X_{n-1}} \sum_{X_n} \prod_{k=m}^n P(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_m} \cdots \sum_{X_{n-1}} \prod_{k=m}^{n-1} P(X_k | \text{Pa}(X_k)) \sum_{X_n} P(X_n | \text{Pa}(X_n)) \\
&= \sum_{X_m} \cdots \sum_{X_{n-1}} \prod_{k=m}^{n-1} P(X_k | \text{Pa}(X_k)) \\
&\quad \cdots \\
&= \sum_{X_m} P(X_m | \text{Pa}(X_m)) \\
&= 1,
\end{aligned}$$

since for each k , and any assignment of the Random Variables in $\text{Pa}(X_k)$, $P(X_k | \text{Pa}(X_k))$ is a probability distribution for the Random Variable X_k , and thus sums to one. The fact that $P(X_k | \text{Pa}(X_k))$ does not depend on any variable X_i for which X_i is succeeding X_k in a topological ordering, is used to move products outside the innermost sum. That is, in a topological ordering, a given node is succeeding all of its ancestors. The Local Probability Distribution $P(X_k | \text{Pa}(X_k))$, varies only with the different assignments to X_k 's parents, which is a subset of X_k 's ancestors. Therefore, this Local Probability Distribution cannot depend on a node X_i succeeding the node X_k in the topological ordering, in this setting denoted by $k < i$.

By Bayes Rule,

$$\mathbb{P}(X_j | \text{Pa}(X_j)) = \frac{\mathbb{P}(X_j, \text{Pa}(X_j))}{\mathbb{P}(\text{Pa}(X_j))}. \tag{2.4}$$

Let

$$C_j = \emptyset,$$

and note that this implies that

$$A_j = \text{Anc}(X_j) \cup \{X_j\}.$$

Define

$$\begin{aligned}
\tilde{A}_j &= A_j \setminus \text{Pa}(X_j) \setminus \{X_j\} \\
&= (\text{Anc}(X_j) \cup \{X_j\}) \setminus (\text{Pa}(X_j) \cup \{X_j\}) \\
&= \text{Anc}(X_j) \setminus \text{Pa}(X_j).
\end{aligned}$$

For any assignment of the Random Variables in $\text{Pa}(X_j)$, the denominator in (2.4) can be expressed as

$$\begin{aligned}
\mathbb{P}(\text{Pa}(X_j)) &= \sum_{X_i \in (\text{Pa}(X_j))^c} \mathbb{P}(X_1, \dots, X_n) \\
&= \sum_{X_{i_2} \in \tilde{A}_j \cup \{X_j\}} \sum_{X_{i_1} \in A_j^c} \prod_{X_k \in A_j} P(X_k | \text{Pa}(X_k)) \prod_{X_k \in A_j^c} P(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_{i_2} \in \tilde{A}_j \cup \{X_j\}} \prod_{X_k \in A_j} P(X_k | \text{Pa}(X_k)) \left(\sum_{X_{i_1} \in A_j^c} \prod_{X_k \in A_j^c} P(X_k | \text{Pa}(X_k)) \right) \\
&= \sum_{X_i \in \tilde{A}_j \cup \{X_j\}} \prod_{X_k \in A_j} P(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_i \in \tilde{A}_j} \sum_{X_j} P(X_j | \text{Pa}(X_j)) \prod_{X_k \in \text{Anc}(X_j)} P(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_i \in \tilde{A}_j} \prod_{X_k \in \text{Anc}(X_j)} P(X_k | \text{Pa}(X_k)) \left(\sum_{X_j} P(X_j | \text{Pa}(X_j)) \right) \\
&= \sum_{X_i \in \tilde{A}_j} \prod_{X_k \in \text{Anc}(X_j)} P(X_k | \text{Pa}(X_k)),
\end{aligned}$$

where we have used (2.3) for $C_j = \emptyset$. Correspondingly, for any assignment of the Random Variables in $\text{Pa}(X_j) \cup \{X_j\}$,

$$\begin{aligned}
\mathbb{P}(X_j, \text{Pa}(X_j)) &= \sum_{X_i \in (\{X_j\} \cup \text{Pa}(X_j))^c} \mathbb{P}(X_1, \dots, X_n) \\
&= \sum_{X_{i_2} \in \tilde{A}_j} \sum_{X_{i_1} \in A_j^c} \prod_{X_k \in A_j} P(X_k | \text{Pa}(X_k)) \prod_{X_k \in A_j^c} P(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_{i_2} \in \tilde{A}_j} \prod_{X_k \in A_j} P(X_k | \text{Pa}(X_k)) \left(\sum_{X_{i_1} \in A_j^c} \prod_{X_k \in A_j^c} P(X_k | \text{Pa}(X_k)) \right)
\end{aligned}$$

$$\begin{aligned}
&= \sum_{X_i \in \bar{A}_j} \prod_{X_k \in A_j} P(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_i \in \bar{A}_j} P(X_j | \text{Pa}(X_j)) \prod_{X_k \in \text{Anc}(X_j)} P(X_k | \text{Pa}(X_k)) \\
&= P(X_j | \text{Pa}(X_j)) \sum_{X_i \in \bar{A}_j} \prod_{X_k \in \text{Anc}(X_j)} P(X_k | \text{Pa}(X_k)) \\
&= P(X_j | \text{Pa}(X_j)) \cdot \mathbb{P}(\text{Pa}(X_j)).
\end{aligned}$$

That is, the Local Probability Distributions $P(X_k | \text{Pa}(X_k))$ equals the Conditional Probability Distributions $\mathbb{P}(X_k | \text{Pa}(X_k))$, since (2.4) reduces to

$$\mathbb{P}(X_j | \text{Pa}(X_j)) = \frac{\mathbb{P}(X_j, \text{Pa}(X_j))}{\mathbb{P}(\text{Pa}(X_j))} = P(X_k | \text{Pa}(X_k)).$$

Thus, we can write (2.1) as

$$\mathbb{P}(X_1, \dots, X_n) = \prod_{i=1}^n \mathbb{P}(X_i | \text{Pa}(X_i)). \quad (2.5)$$

Then again, let A_j be any set as in (2.2), and $h(X_1, \dots, X_n)$ be a function whose value only depends on the assignment of the variables in A_j . We know that

$$\begin{aligned}
\mathbb{E}h &= \sum_{\text{All RVs}} h(X_1, \dots, X_n) \mathbb{P}(X_1, \dots, X_n) \\
&= \sum_{X_{i_1} \in A_j} \sum_{X_{i_2} \in A_j^c} h(X_1, \dots, X_n) \mathbb{P}(X_1, \dots, X_n) \\
&= \sum_{X_{i_1} \in A_j} \sum_{X_{i_2} \in A_j^c} h(X_1, \dots, X_n) \prod_{X_k \in A_j} \mathbb{P}(X_k | \text{Pa}(X_k)) \prod_{X_k \in A_j^c} \mathbb{P}(X_k | \text{Pa}(X_k)).
\end{aligned}$$

Since h only depends on variables in A_j , we could write $h = h(X_{A_j})$, and get

$$\begin{aligned}
\mathbb{E}h &= \sum_{X_{i_1} \in A_j} \sum_{X_{i_2} \in A_j^c} h(X_{A_j}) \prod_{X_k \in A_j} \mathbb{P}(X_k | \text{Pa}(X_k)) \prod_{X_k \in A_j^c} \mathbb{P}(X_k | \text{Pa}(X_k)) \\
&= \sum_{X_{i_1} \in A_j} h(X_{A_j}) \prod_{X_k \in A_j} \mathbb{P}(X_k | \text{Pa}(X_k)) \left(\sum_{X_{i_2} \in A_j^c} \prod_{X_k \in A_j^c} \mathbb{P}(X_k | \text{Pa}(X_k)) \right) \\
&= \sum_{X_{i_1} \in A_j} h(X_{A_j}) \prod_{X_k \in A_j} \mathbb{P}(X_k | \text{Pa}(X_k)). \quad (2.6)
\end{aligned}$$

Actually,

$$\prod_{X_k \in A_j} \mathbb{P}(X_k | \text{Pa}(X_k)) = \mathbb{P}(X_{A_j}),$$

which, in fact, is easily proved by (2.6) by letting $h(X_1, \dots, X_n)$ be the indicator function for some assignment of the Random Variables in A_j . However, the main observation from (2.6), is that to find the expected value $\mathbb{E}h$, it is sufficient to use the probability distribution we get by ignoring all the factors in (2.5) which are Conditional Probability Distributions for the Random Variables in A_j^c . That is, letting $\mathbb{P}(X_{A_j})$ denote the product in the above equation, we get

$$\mathbb{E}h(X_{A_j}) = \sum_{A_j} h(X_{A_j}) \mathbb{P}(X_{A_j}). \quad (2.7)$$

2.2 Conditional Independence

As mentioned, each edge in the graph encodes a possible conditional dependence relationship between two nodes. That is, the edges in the graph determines the factors in the formula (2.5), which in turn allows for certain conditional dependence relationships. Whether a set of variables actually are conditionally dependent of each other, is determined by the conditional probability distributions $\{\mathbb{P}(X_i | \text{Pa}(X_i))\}$.

We let $X \perp Y$ denote that two Random Variables X, Y are independent, i.e.

$$\mathbb{P}(X = x) = \mathbb{P}(X = x | Y = y) \quad \forall x, y.$$

An equivalent statement is

$$\mathbb{P}(X = x, Y = y) = \mathbb{P}(X = x) \cdot \mathbb{P}(Y = y) \quad \forall x, y.$$

Also, the symbol $|$ indicates that we are discussing a conditional probability distribution, that is, $X \perp Y | Z$ means

$$\mathbb{P}(X = x, Y = y | Z = z) = \mathbb{P}(X = x | Z = z) \cdot \mathbb{P}(Y = y | Z = z) \quad \forall x, y, z.$$

Note that this means that $X \perp Y$ and $X \perp Y | \emptyset$ are equivalent statements.

Correspondingly, we let $X \not\perp Y$ denote that the graph does not encode that the Random Variables X, Y are independent. Also, $X \not\perp Y | Z$ means that the graph does not encode that X and Y are independent conditional on Z . That does not mean, however, that there are no set of conditional probability distributions that makes them independent. In fact, if we let X or Y be a constant,

they will always be conditionally independent, but their joint distribution can still be represented on a graph where they are connected by edges. The edges just allow for Random Variables not to be independent.

As an example, we study the graph in Figure 1. There are algorithms for finding all conditional independence relationships, see for example the Bouncing Ball Algorithm in Jordan (n.d.). However, this will not be covered in our scope. To introduce some intuition, we will present some of the possible conditional independence relationships from the graph in Figure 1, and give a brief, non rigorous discussion of what they mean for a given example.

We could look at the nodes in Figure 1 as persons, and let the edges indicate where messages could be sent. Say, K_1 suddenly wants to have a dinner party, and on his phone he only has the phone numbers to P_1 and P_2 . Correspondingly, P_1 can only communicate a message to P_2 and S_1 , and P_2 to S_2 and S_3 . Say, K_1 would want all to come, but he can only say this to P_1 and P_2 . However, through them all persons can be reached, but P_1 and P_2 would only forward the message if they are going themselves. The Random Variable for each node would here be a Boolean variable $\{\mathbf{True}, \mathbf{False}\}$ indicating whether the person is attending the dinner party or not. If K_1 is having the party for sure, he would be a constant Random Variable, having value \mathbf{True} with probability 1. Otherwise, his node could be associated with some probability distribution indicating a probability for him inviting to a dinner party or not. In that case, assuming all telephones are working perfectly, and all persons are going if they get the message, we know that all Random Variables are either \mathbf{True} or all Random Variables are \mathbf{False} . Thus, we cannot have

$$S_1 \perp S_2,$$

$$K_1 \perp S_1$$

nor

$$S_2 \perp S_3.$$

However, if we know the value of P_2 , then S_1 and S_2 are independent, written

$$S_2 \perp S_3 \mid P_2.$$

Then, allow for the telephones not to be perfect, that is, not all messages sent will be received. For each message sent, whether it is received or not is assumed to be independent of whether any other sent message is received. For example, the statement

$$S_1 \perp S_2 \mid P_2$$

is explained by the fact that knowing whether P_2 is going or not leaves only what we non rigorously can call independent randomness for the impact on the values of S_1 and S_2 , respectively.

Note that any $\not\perp$ statement can be proved by a counter example, while the explanations for the \perp statements above just serves as describing examples. The above examples can be summarized in the following list of conditional independence statements for the graph in Figure 1.

- $K_1 \not\perp S_1$
- $K_1 \perp S_1 \mid P_1$
- $S_2 \not\perp S_3$
- $S_2 \perp S_3 \mid P_2$
- $S_1 \not\perp S_2$
- $S_1 \perp S_2 \mid P_1$
- $S_1 \perp S_2 \mid P_2$
- $S_1 \perp S_2 \mid P_1, P_2$
- $S_1 \not\perp S_2 \mid K_1$

Note that this is just a small excerpt of all the possible conditional independence statements we could deduce from Figure 1. To find more of them, or for a more rigorous check, the reader is referred to either the Bouncing Ball Algorithm in Jordan (n.d.), or to do the more tedious work of checking by use of Bayes Rule on the general Joint Probability Distribution (2.5) for the given graph.

In general, there are two standard conditional independence relations that are characteristic for all Bayesian Networks. They are to be found in the following two Theorems from Russell and Norvig (2003), and assume a Joint Probability Distribution where each assignment of the Random Variables X_1, \dots, X_n has a positive probability,

$$\mathbb{P}(X_1, \dots, X_n) > 0.$$

Theorem 2.1. *A Random Variable in a Bayesian Network is conditionally independent of its non-descendants, given its parents.*

To state the last theorem, we have to make the following definition as in Russell and Norvig (2003).

Definition 2.4. *The Markov Blanket of a Node in a Bayesian Network is defined to be the set of its neighboring nodes, that is, its parents, its children and the parents of its children.*

Theorem 2.2. *A Random Variable in a Bayesian Network is conditionally independent of all other nodes in the network, given its Markov Blanket, as in Definition 2.4*

Theorems 2.1 and 2.2 can be proved by applying Bayes rule on the Joint Probability Distribution as in (2.5).

2.3 Markov Random Fields

In a Bayesian Network, a Directed Acyclic Graph is used to represent some dependence properties of the Joint Probability Distribution for the Random Variables. But the same Joint Probability Distribution (2.5) can also be represented on an undirected graph. A Markov Random Field is an undirected graphical model to represent probabilistic relationships. That is, as for a Bayesian Network, each node still represents a Random Variable, but in a Markov Random Field the edges are undirected. Thus, we cannot talk about a node's parents nor its children, descendants or ancestors. Instead we have the following definition from Bishop (2006).

Definition 2.5. *A clique is a subset of the nodes in the undirected graph such that there exists an (undirected) edge between all pairs of nodes in the subset. Furthermore, a maximal clique is a clique for which it is not possible to include any other nodes in the graph without it ceasing to be a clique.*

A potential function ψ on a maximal clique is a non negative function on the Random Variables in the clique. By letting Λ denote a maximal clique, and X_Λ denote the Random Variables in it, we can define a probability distribution on an undirected graph as the product of all potential functions $\psi_\Lambda(X_\Lambda)$, as in

$$\mathbb{P}(X_1, \dots, X_n) = \frac{1}{C} \prod_{\Lambda} \psi_\Lambda(X_\Lambda), \quad (2.8)$$

where C is the normalization constant that ensures it to be a probability distribution. The undirected graph and the corresponding probability distribution is then a Markov Random Field.

As for Bayesian Networks, there are also conditional independence statements that are true in general for Markov Random Fields. The following is

from Bishop (2006), and assumes a Joint Probability Distribution where

$$\mathbb{P}(X_1, \dots, X_n) > 0$$

for any assignment to the Random Variables X_1, \dots, X_n .

Theorem 2.3. *For three sets of nodes A, B, D , we have*

$$A \perp B \mid D$$

if all possible (undirected) paths between a node in A and a node in B pass through one or more nodes in D .

From this, we can deduce that a Random Variable is conditionally independent of any subset of the other Random Variables given its neighbors. Thus, the Markov Blanket of a Random Variable X_i in a Markov Random Field, is the set of nodes X_j which shares an edge $e = \{X_i, X_j\} \in E$ with X_i .

Given a Bayesian Network, we would like to find a corresponding Markov Random Field. For the same conditional independence properties to hold, we need the Markov Blanket for each Random Variable to be the same in both representations. Hence, when going from the Bayesian Network representation to the Markov Random Field representation, edges must be added between any two nodes that have common children. In fact, according to Bishop (2006), in addition to replacing all directed edges in the original Bayesian Network by undirected ones, this is all we need to do. The process of adding the extra edges between any pair of parents of the same child is called "moralization", and is also referred to as "marrying the parents". The resulting undirected graph is called the moral graph, and is also the corresponding undirected graphical representation of the probability distribution.

For the Joint Probability Distribution, note from (2.5) and (2.8), that we need to have

$$\frac{1}{C} \prod_{\Lambda} \psi_{\Lambda}(X_{\Lambda}) = \prod_{i=1}^n \mathbb{P}(X_i | \text{Pa}(X_i)). \quad (2.9)$$

As an example, we study the directed graph in Figure 1. Note that no moralization has to be done, since K_1 and P_1 are the only nodes with common children, and they already share an edge. Thus, the moral graph of Figure 1 has the same nodes and the same edges, except that now the edges are undirected ones. For this graph, the Bayesian Network tells us that

$$\mathbb{P}(K_1, \dots, S_3) = \mathbb{P}(K_1) \mathbb{P}(P_1 | K_1) \mathbb{P}(S_1 | P_1) \mathbb{P}(P_2 | K_1, P_1) \mathbb{P}(S_2 | P_2) \mathbb{P}(S_3 | P_2).$$

The corresponding moral graph tells us that

$$\mathbb{P}(K_1, \dots, S_3) = \frac{1}{C} \psi_1(P_1, S_1) \psi_2(P_2, S_2) \psi_3(P_2, S_3) \psi_4(K_1, P_1, P_2).$$

Comparing the two above equations, we observe from (2.9) that we can choose

$$\begin{aligned} \psi_1(P_1, S_1) &= C_1 \mathbb{P}(S_1|P_1) \\ \psi_2(P_2, S_2) &= C_2 \mathbb{P}(S_2|P_2) \\ \psi_3(P_2, S_3) &= C_3 \mathbb{P}(S_3|P_2) \\ \psi_4(K_1, P_1, P_2) &= C_4 \mathbb{P}(K_1) \mathbb{P}(P_1|K_1) \mathbb{P}(P_2|K_1, P_1) \end{aligned}$$

for some constants C_1, C_2, C_3, C_4 such that $C = \frac{1}{C_1 C_2 C_3 C_4}$. The general case is similar, and we can always choose all C_i s to be 1, and thus get $C = 1$. From now on, we will assume this choice.

In general, note that moralization ensures that for any node X_i , there exists a maximal clique Λ in the moral graph such that

$$\{X_i\} \cup \text{Pa}(X_i) \subseteq \Lambda.$$

Thus, we observe from (2.9), that for any maximal clique Λ , there is a non empty subset $\lambda \subseteq \Lambda$ such that

$$\psi_\Lambda = \prod_{X_i \in \lambda} \mathbb{P}(X_i | \text{Pa}(X_i)).$$

Now, let us study a subset of the Random Variables in Figure 1. Let $\tilde{G} = (\tilde{V}, \tilde{E})$, where

$$\tilde{V} = \{ P_2, S_2, S_3 \} \subset V \quad \text{and} \quad \tilde{E} = \{ (P_2, S_2), (P_2, S_3) \} \subset E$$

for $G = (V, E)$ in Figure 1. A visualization of \tilde{G} can be found in Figure 2, and the corresponding moral graph can be found in Figure 3. Observe that the maximal cliques are

$$\Lambda_1 = \{ S_2, P_2 \} \quad \text{and} \quad \Lambda_2 = \{ S_3, P_2 \}.$$

Correspondingly, the potential functions must be of the form

$$\psi_1 = \Phi_1(P_2) \mathbb{P}(S_2|P_2) \quad \text{and} \quad \psi_2 = \Phi_2(P_2) \mathbb{P}(S_3|P_2),$$

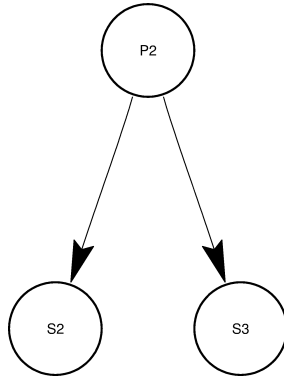


Figure 2: The graph \tilde{G} created by a subset of the nodes in Figure 1.

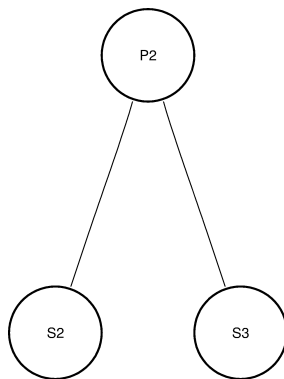


Figure 3: A visualization of the moral graph for \tilde{G} , where \tilde{G} is the graph visualized in Figure 2. Note that no extra edges had to be added to obtain the moral graph.

where $\Phi_1(P_2)\Phi_2(P_2) = \mathbb{P}(S_2)$, and we can choose either $\Phi_1(P_2) \equiv 1$ or $\Phi_2(P_2) \equiv 1$.

From this example, we note that the choice of λ might not be unique for each maximal clique Λ . In any case, the λ s define a partition of all the nodes X_1, \dots, X_n . That is, the λ s are pairwise disjoint and their union contains all nodes in the graph.

3 Pseudo-Boolean functions

In several applications, we are working with a set of variables that can take on two values. In this section, we will present some theory about functions on such a set. As in Hammer and Rudeanu (1968), we begin by the following definition.

Definition 3.1. *The set $B_2 = \{0, 1\}$ together with the operations of disjunction, conjunction and negation is called the Boolean algebra.*

That is, the operations mentioned is defined as follows.

Definition 3.2. *Negation, denoted by $\bar{}$, is associated with the function $1 - x$, and we get the following formulas*

$$\bar{0} = 1 \quad \text{and} \quad \bar{1} = 0.$$

Definition 3.3. *Conjunction, denoted by \cdot , works like usual multiplication. That is*

$$0 \cdot 0 = 0 \quad 0 \cdot 1 = 0 \quad 1 \cdot 0 = 0 \quad 1 \cdot 1 = 1.$$

Definition 3.4. *Boolean disjunction is denoted by \cup , and is defined by the formulas*

$$0 \cup 0 = 0 \quad 0 \cup 1 = 1 \quad 1 \cup 0 = 1 \quad 1 \cup 1 = 1.$$

Also, we let the term Boolean variable denote a variable that can take values in the set B_2 . A Boolean function f is then a function

$$f : B_2^n \rightarrow B_2,$$

that is, a function whose value and n arguments are all Boolean variables. This concept of a Boolean function is then expanded to the following.

Definition 3.5. A pseudo-Boolean function f is a function

$$f : B_2^n \rightarrow \mathbb{R},$$

that is, a function from an n -tuple of bivalent variables to the real numbers.

Note that this definition allows us to look at the space of Boolean functions as a subspace of all pseudo-Boolean functions.

The following result is from Hammer and Rudeanu (1968).

Theorem 3.1. Every pseudo-Boolean function may be written as a polynomial, which is linear in each variable, and which, after the reduction of the similar terms, is uniquely determined up to the order of the sums and products.

That is, let $X = [X_k]_{k=1}^n$ be a vector of Boolean variables and f be a pseudo-Boolean function. Then, there exists a collection T of sets $\lambda \subseteq \{1, \dots, n\}$ of indexes and a corresponding set of coefficients $\{\alpha^\lambda\}_{\lambda \in T}$ such that

$$f(X) = \sum_{\lambda \in T} \alpha^\lambda \prod_{k \in \lambda} X_k. \quad (3.1)$$

In fact, as mentioned in Theorem 3.1, there might be several choices of T . T can always be the power set of $\{1, \dots, n\}$, and in general

$$T \subseteq 2^{\{1, \dots, n\}}.$$

As an example, we look at a function h acting on the variables in Figure 1,

$$h(K_1, P_1, P_2, S_1, S_2, S_3) = \begin{cases} 1 & \text{if } S_2 = 0, S_3 = 0, \\ e & \text{if } S_2 = 1, S_3 = 0, \\ e^2 & \text{if } S_2 = 0, S_3 = 1, \\ e + e^2 - 1 & \text{if } S_2 = 1, S_3 = 1. \end{cases} \quad (3.2)$$

For this function, we can let

$$T = \{ \emptyset, \{S_1\}, \{S_2\} \},$$

and write h as

$$h(K_1, P_1, P_2, S_1, S_2, S_3) = 1 + (e - 1)S_2 + (e^2 - 1)S_3.$$

Theorem 3.1 is easily proved by a calculation of the coefficients $\{\alpha^\lambda\}_{\lambda \in T}$ for a general function, which can be done recursively as follows. First, for simplicity, assume $T = 2^{\{1, \dots, n\}}$, and let

$$v_\Lambda = [v_k]_{k=1}^n \quad \text{such that} \quad v_k = 1 \Leftrightarrow k \in \Lambda.$$

That is, we assume T to be the power set of all indexes appearing in S , and v_Λ to be a vector where all entries with indexes in Λ are on, and all other entries are off. This allows us to set

$$\alpha^\emptyset = f(v_\emptyset) = f(\vec{0}),$$

and for increasing size of Λ , set

$$\alpha^\Lambda = f(v_\Lambda) - \sum_{\lambda \subsetneq \Lambda} \alpha^\lambda.$$

After determining the full set of coefficients $\{\alpha_\lambda\}$, the choice of T can be changed by not including all λ s where $\alpha^\lambda = 0$.

In Hammer and Rudeanu (1968), several other representations of a pseudo-Boolean function, also using combinations of the operations in Definition 3.2 and Definition 3.4 are presented. In the following, we will assume the representation from (3.1).

From Theorem 3.1, we also deduce that any such positive valued function $f(X)$ can be represented as the exponential of a pseudo-Boolean function. That is,

$$f(X) = \exp \left(\sum_{\lambda \in S} \beta^\lambda \prod_{k \in \lambda} X_k \right). \quad (3.3)$$

As an example, the function h in (3.2) can be written as

$$h(K_1, P_1, P_2, S_1, S_2, S_3) = \exp (S_2 + 2S_3 + S_2S_3(\ln(e + e^2 - 1) - 3)).$$

In the following, we will continue to write α^λ for coefficients of the pseudo-Boolean representation (3.1), and β^λ for coefficients of the pseudo-Boolean representation of the logarithm of the given function as in (3.3). Correspondingly, T will denote the subsets we are summing over in the pseudo-Boolean representation (3.1), and S for the exponential version (3.3).

3.1 A graph structure for coefficients

When a computer program is given a pseudo-Boolean representation of a function, or when calculating the coefficients of that representation for a given function, a structure for storing coefficients of type α^λ is needed. Also, expressions of type

$$\sum_{\lambda \subseteq \Lambda} \alpha^\lambda$$

appear frequently in the formulas, so given Λ , its coefficient and the coefficients of subsets of Λ should be easily accessed. This is easily solved by a graph structure. The graph is a set of nodes, with one root node corresponding to α^\emptyset , and all other nodes divided into layers depending on the size of Λ for the coefficient α^Λ the node corresponds to. Let N denote the set of indexes for the variables the given function f depends on. For simplicity, assume $N = \{1, 2, \dots, n\}$, and also that we study the full representation

$$f(X) = \sum_{\lambda \subseteq N} \alpha^\lambda \prod_{k \in \lambda} X_k.$$

Then, for each $\Lambda \subseteq N$, in the coefficient graph, the corresponding node has all subsets of Λ as ancestors. Especially, the parents of that given node are the nodes corresponding to the sets in

$$\{ \lambda \subset \Lambda : |\lambda| = |\Lambda| - 1 \} = \{ \lambda : \lambda = \Lambda \setminus \{m\}, m \in \Lambda \}.$$

That is, to obtain a child of a given node λ , one augments λ with an $m \in N \setminus \lambda$. Correspondingly, each parent is found by removing one of the elements in λ . Such a coefficient graph with $n = 4$ and four layers, can be seen in Figure 4.

As an example, given $\Lambda = \{3, 4, 7\}$ and $n = 7$, we observe

$$\text{Pa}(\Lambda) = \{ \{4, 7\}, \{3, 7\}, \{3, 4\} \}$$

and correspondingly,

$$\text{Ch}(\Lambda) = \{ \{1, 3, 4, 7\}, \{2, 3, 4, 7\}, \{3, 4, 5, 7\}, \{3, 4, 6, 7\} \}.$$

An illustration of the node $\Lambda = \{3, 4, 7\}$ and its parents and children is to be found in Figure 5. We let layer r denote the set of nodes of distance r from the root node, so that all children of a given node are in the same layer, and similarly for all its parents.

For a further description of how we have implemented this graph structure, the reader is referred to Appendix A.

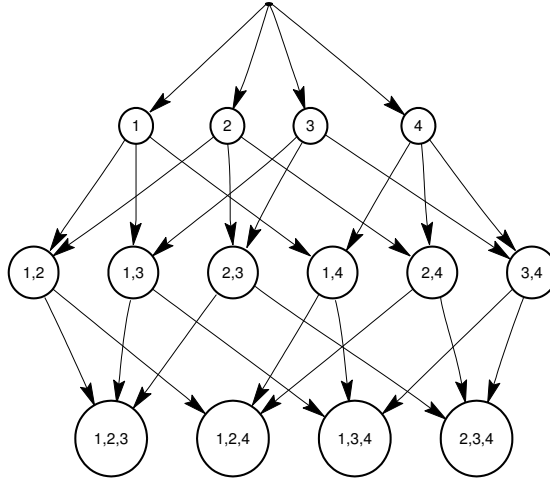


Figure 4: An example coefficient graph of four layers including the root node, with $n = 4$. That is, coefficient nodes corresponding to $|\lambda| \leq 3$ are represented. The arrows are from parents to children.

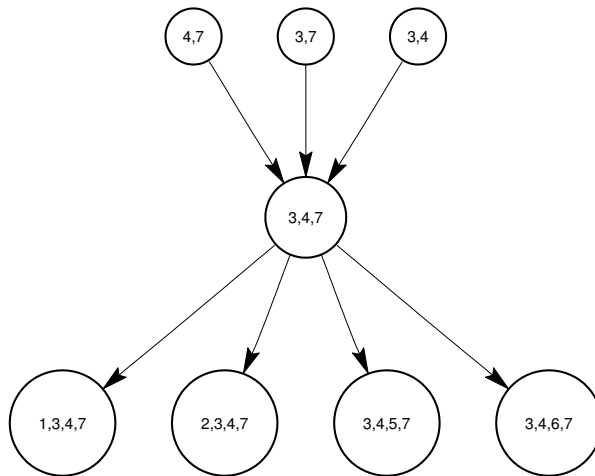


Figure 5: An illustration of the example showing parents and children for the node $\Lambda = \{3, 4, 7\}$ when $n = 7$.

3.2 Converting between exponential and linear form

Given a set T of index sets λ , and the set coefficients $\{\alpha^\lambda\}_{\lambda \in T}$ to describe a function $f(X)$ of the form

$$f(X) = \sum_{\lambda \in T} \alpha^\lambda \prod_{k \in \lambda} X_k,$$

we want to find a corresponding set S , and the set of coefficients $\{\beta^\lambda\}_{\lambda \in S}$ to write $f(X)$ as

$$f(X) = \exp \left(\sum_{\lambda \in S} \beta^\lambda \prod_{k \in \lambda} X_k \right).$$

First, note that in general $S \neq T$. As an example, note for $a, b \in \mathbb{R}$, we get the following pair of equivalent representations

$$1 + aX_1 + bX_2 = \exp \left(X_1 \ln(1 + a) + X_2 \ln(1 + b) + X_1 X_2 \ln \frac{1 + a + b}{(1 + a)(1 + b)} \right).$$

That is, in general, the exponential representation is full even though the linear representation is sparse. Correspondingly, the linear representation might be full for a function with sparse exponential representation.

For simplicity, assume S to be the power set of all indexes appearing in T , that is,

$$S = \mathcal{2}^{\{j : \exists \lambda \in T \text{ such that } j \in \lambda\}}.$$

We get, for $\Lambda \in S$,

$$\beta^\Lambda = \ln \left(\sum_{\lambda \in T : \lambda \subseteq \Lambda} \alpha^\lambda \right) - \sum_{\lambda \subsetneq \Lambda} \beta^\lambda.$$

This introduces an algorithm for calculating the β^λ s in order of increasing size of λ . Correspondingly, for the converse case, we have for $\Lambda \in T$

$$\alpha^\Lambda = \exp \left(\sum_{\lambda \in S : \lambda \subseteq \Lambda} \beta^\lambda \right) - \sum_{\lambda \subsetneq \Lambda} \alpha^\lambda.$$

3.3 Probability Distribution for a Bayesian Network

Given a Bayesian Network with nodes $X = [X_i]_{i=1}^n$, we have the full Joint Probability Distribution as in (2.5), after specifying the conditional probabilities

$\mathbb{P}(X_i|\text{Pa}(X_i))$ for each node X_i . As in Section 6, we might want to represent the natural logarithm of the full Joint Probability Distribution as a pseudo-Boolean function. Observe that

$$\ln \mathbb{P}(X_1, \dots, X_n) = \sum_{i=1}^n \ln \mathbb{P}(X_i|\text{Pa}(X_i)),$$

where for each i , $\mathbb{P}(X_i|\text{Pa}(X_i))$ is a function of the Random Variables in $\{X_i\} \cup \text{Pa}(X_i)$. Thus, if we know the pseudo-Boolean representation of $\ln \mathbb{P}(X_i|\text{Pa}(X_i))$ for each i , their sum equals the natural logarithm of the full Joint Probability Distribution.

Given an index i , we write the pseudo-Boolean representation of the conditional probability distribution $\mathbb{P}(X_i|\text{Pa}(X_i))$ as

$$\ln \mathbb{P}(X_i|\text{Pa}(X_i)) = \sum_{\lambda \in \mathcal{S}_i} \gamma_i^\lambda \prod_{X_k \in \lambda} X_k,$$

where

$$\mathcal{S}_i = 2^{\{j \mid X_j \in \{X_i\} \cup \text{Pa}(X_i)\}}.$$

Note that here, the full representation is chosen, that is, we sum over the power set of $\{X_i\} \cup \text{Pa}(X_i)$. Also, from here, we let the coefficients of the pseudo-Boolean representation of probability distributions be denoted by γ s, in fact for the conditional distribution $\mathbb{P}(X_i|\text{Pa}(X_i))$, we use $\{\gamma_i^\lambda\}_\lambda$.

Now, we assume that the nodes represent binary Random Variables, that is $X_i \in \{0, 1\}$. Observe, that the coefficients γ_i^λ can be calculated recursively, in order of increasing size of λ . In fact, first let

$$\gamma_i^\emptyset = \ln \mathbb{P}(X_i = 0 \mid X_k = 0 \ \forall X_k \in \text{Pa}(X_i)).$$

Then, for any subset $\Lambda \subseteq \text{Pa}(X_i)$,

$$\gamma_i^\Lambda = \ln \mathbb{P}(X_i = 0 \mid X_k = 1 \ \forall X_k \in \Lambda \text{ and } X_j = 0 \ \forall X_j \in \text{Pa}(X_i) \setminus \Lambda) - \sum_{\lambda \subsetneq \Lambda} \gamma_i^\lambda.$$

Similarly, for any set $\Lambda = \{X_i\} \cup \Psi$ where $\Psi \subseteq \text{Pa}(X_i)$,

$$\gamma_i^\Lambda = \ln \mathbb{P}(X_i = 1 \mid X_k = 1 \ \forall X_k \in \Psi \text{ and } X_j = 0 \ \forall X_j \in \text{Pa}(X_i) \setminus \Psi) - \sum_{\lambda \subsetneq \Lambda} \gamma_i^\lambda.$$

3.3.1 An example Network

This example will illustrate the Joint Probability Distribution for the tests presented in Sections 8 and 9. We have a Bayesian Network over Random Variables that are either on or off, corresponding to the values 1 and 0, respectively. Assume we are given a probability p_{X_i} for each root node X_i , and then let

$$p_{X_i} = \mathbb{P}(X_i = 1) = 1 - \mathbb{P}(X_i = 0).$$

This implies the following formula for the root nodes X_i ,

$$\ln \mathbb{P}(X_i) = \ln(1 - p_{X_i}) + X_i(\ln p_{X_i} - \ln(1 - p_{X_i})),$$

indicating

$$\begin{aligned}\gamma_i^\emptyset &= \ln(1 - p_{X_i}) \\ \gamma_i^{\{i\}} &= \ln p_{X_i} - \ln(1 - p_{X_i}).\end{aligned}$$

For each edge e_{X_i, X_j} from node X_i to node X_j , assume we are given two probabilities, p_{X_i, X_j}^0 and p_{X_i, X_j}^1 , respectively. In fact, let

$$p_{X_i, X_j}^0 = \mathbb{P}(X_j \text{ receives signal from } X_i | X_i = 0)$$

and

$$p_{X_i, X_j}^1 = \mathbb{P}(X_j \text{ receives signal from } X_i | X_i = 1).$$

Whether a child X_j receives a signal from one of its parents, say X_i , is assumed to be conditionally independent of whether a signal is received in any other pair of child and parent, given the value of the parent X_i . Assume that for any non root node, its value is 1 if it has received a signal from at least one of its parents, and 0 otherwise.

Given a node X_j and an assignment $X_i = x_i$ to each of its parents $X_i \in \text{Pa}(X_j)$, we let

$$\text{Pa}(X_j) = \{X_i \mid i \in \Lambda \cup \Psi\},$$

where $\Lambda \cup \Psi$ is a decomposition of X_j 's parents such that

$$x_i = \begin{cases} 1 & \text{if } X_i \in \Lambda, \\ 0 & \text{if } X_i \in \Psi. \end{cases}$$

Observe that

$$\begin{aligned} \ln \mathbb{P}(X_j = 0 | \text{Pa}(X_j)) &= \ln \left(\prod_{i \in \Lambda} (1 - p_{X_i, X_j}^1) \prod_{i \in \Psi} (1 - p_{X_i, X_j}^0) \right) \\ &= \sum_{i \in \Lambda} \ln (1 - p_{X_i, X_j}^1) + \sum_{i \in \Psi} \ln (1 - p_{X_i, X_j}^0), \end{aligned}$$

which implies

$$\gamma_j^\Lambda = \sum_{i \in \Lambda} \ln (1 - p_{X_i, X_j}^1) + \sum_{i \in \Psi} \ln (1 - p_{X_i, X_j}^0) - \sum_{\lambda \subsetneq \Lambda} \gamma_j^\lambda \quad (3.4)$$

and

$$\gamma_j^{\Lambda \cup \{j\}} = \ln \left(1 - \prod_{i \in \Lambda} (1 - p_{X_i, X_j}^1) \prod_{i \in \Psi} (1 - p_{X_i, X_j}^0) \right) - \sum_{\lambda \subsetneq \Lambda \cup \{j\}} \gamma_j^{\lambda \cup \{j\}}.$$

Observe from (3.4) that for a non root node X_j , we get

$$\gamma_j^\emptyset = \sum_{X_i \in \text{Pa}(X_j)} \ln(1 - p_{X_i, X_j}^0),$$

and if $X_k \in \text{Pa}(X_j)$,

$$\begin{aligned} \gamma_j^{\{k\}} &= \ln(1 - p_{X_k, X_j}^1) + \sum_{X_i \in \text{Pa}(X_j) \setminus \{X_k\}} \ln(1 - p_{X_i, X_j}^0) - \gamma_j^\emptyset \\ &= \ln(1 - p_{X_k, X_j}^1) - \ln(1 - p_{X_k, X_j}^0). \end{aligned}$$

But then also, for $X_k, X_\ell \in \text{Pa}(X_j)$ where $k \neq \ell$, (3.4) reduces to

$$\gamma_j^{\{k, \ell\}} = 0.$$

In fact, we get

$$\gamma_j^\Lambda = 0 \text{ when } |\Lambda| \geq 2$$

by induction on the size of Λ .

As an example, we let $p_{X_i, X_j}^0 = \epsilon$ for all edges e_{X_i, X_j} and calculate the representation of the corresponding conditional distribution for some of the variables in Figure 1. For example, we get

$$\ln \mathbb{P}(K_1) = \ln(1 - p_{K_1}) + K_1(\ln p_{K_1} - \ln(1 - p_{K_1})),$$

$$\ln \mathbb{P}(P_1|K_1) = \ln(1 - \epsilon) + K_1 \ln \frac{1 - p_{K_1, P_1}}{1 - \epsilon} + P_1 \ln \frac{\epsilon}{1 - \epsilon} + K_1 P_1 \ln \frac{(1 - \epsilon)p_{K_1, P_1}}{(1 - p_{K_1, P_1})\epsilon}.$$

and

$$\ln \mathbb{P}(S_1|P_1) = \ln(1 - \epsilon) + P_1 \ln \frac{1 - p_{P_1, S_1}}{1 - \epsilon} + S_1 \ln \frac{\epsilon}{1 - \epsilon} + P_1 S_1 \ln \frac{(1 - \epsilon)p_{P_1, S_1}}{(1 - p_{P_1, S_1})\epsilon}.$$

Note that the formulas for $\ln \mathbb{P}(S_2|P_2)$ and $\ln \mathbb{P}(S_3|P_2)$ are similar to those for $\ln \mathbb{P}(P_1|K_1)$ and $\ln \mathbb{P}(S_1|P_1)$.

4 Approximate Forward-Backward Algorithm

From Tjelmeland and Austad (2011), we have an approximate forward-backward algorithm for probability distributions of the form

$$\mathbb{P}(X_1, \dots, X_n) = \frac{1}{C} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right)$$

That is, the algorithm is constructed to sum out Random Variables, and return the value of the sum. This is useful for example when working with Markov Random Fields, where the normalization constant C might be unknown. From Section 2.3, we recall that

$$C = \sum_{X_1, \dots, X_n} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right).$$

Another application is for calculating the expected value $\mathbb{E}h$ for some function h on a set of Random Variables. Let

$$h(X_1, \dots, X_n) = \exp \left(\sum_{\lambda \in S} \beta^\lambda \prod_{k \in \lambda} X_k \right).$$

Then,

$$\begin{aligned} \mathbb{E}h &= \sum_{X_1, \dots, X_n} h(X_1, \dots, X_n) \mathbb{P}(X_1, \dots, X_n) \\ &= \sum_{X_1, \dots, X_n} \exp \left(\sum_{\lambda \in S} (\beta^\lambda + \gamma^\lambda) \prod_{k \in \lambda} X_k \right). \end{aligned}$$

Also, note that if h is a function which depends on more variables than our Random Variables X_1, \dots, X_n , say

$$h = h(X_1, \dots, X_n, X_{n+1}, \dots, X_{n+m}),$$

we would have

$$h(X_1, \dots, X_{n+m}) = \exp \left(\sum_{\lambda \in S} \beta^\lambda \prod_{k \in \lambda} X_k \right).$$

for a collection S of index sets λ where some of the indexes $k \in \lambda$ might have $k > n$. We get

$$\begin{aligned} \mathbb{E}h &= \sum_{X_1, \dots, X_n} h(X_1, \dots, X_{n+m}) \mathbb{P}(X_1, \dots, X_n) \\ &= \sum_{X_1, \dots, X_n} \exp \left(\sum_{\lambda \in S} (\beta^\lambda + \gamma^\lambda) \prod_{k \in \lambda} X_k \right), \end{aligned}$$

which corresponds to looking for a function $C(X_{n+1}, \dots, X_{n+m})$. Observe how these three problems are essentially the same, and thus we can focus on how to find the normalization constant C .

The exact forward-backward algorithm works by summing out one variable at the time. That is, assume we want to calculate

$$\begin{aligned} C &= \sum_{X_n, X_{n-1}, \dots, X_1} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \\ &= \sum_{X_n} \dots \sum_{X_2} \sum_{X_1} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \\ &= \sum_{X_n} \dots \sum_{X_2} \left(\sum_{X_1} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \right). \end{aligned}$$

First, X_1 is summed out, to get

$$\sum_{X_1} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right) = \exp \left(\sum_{\lambda \in \hat{S}} \hat{\gamma}^\lambda \prod_{k \in \lambda} X_k \right)$$

for some \hat{S} and some collection of coefficients $\{\hat{\gamma}^\lambda\}_{\lambda \in \hat{S}}$. Note that

$$X_1 \notin \lambda \quad \forall \lambda \in \hat{S}.$$

After this step,

$$C = \sum_{X_2, \dots, X_n} \exp \left(\sum_{\lambda \in \dot{S}} \hat{\gamma}^\lambda \prod_{k \in \lambda} X_k \right).$$

Continuing like this, we finally end up with some \bar{S} and some collection of coefficients $\{\bar{\gamma}^\lambda\}_{\lambda \in \bar{S}}$ such that

$$C = \sum_{X_n} \exp \left(\sum_{\lambda \in \bar{S}} \bar{\gamma}^\lambda \prod_{k \in \lambda} X_k \right).$$

We observe that in this stepwise procedure, each step is essentially the same. Therefore, it is sufficient to consider how to calculate

$$C^1(X_2, \dots, X_n) = \sum_{X_1} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right).$$

Observe that for any $\dot{S} \subseteq S$,

$$\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k = \sum_{\lambda \in \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k + \sum_{\lambda \in S \setminus \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k.$$

Now, let

$$\dot{S} = \{\lambda \in S \mid 1 \in \lambda\}$$

Then,

$$\begin{aligned} C^1(X_2, \dots, X_n) &= \sum_{X_1} \exp \left(\sum_{\lambda \in \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k + \sum_{\lambda \in S \setminus \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \\ &= \sum_{X_1} \exp \left(\sum_{\lambda \in \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \exp \left(\sum_{\lambda \in S \setminus \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \\ &= \exp \left(\sum_{\lambda \in S \setminus \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \sum_{X_1} \exp \left(\sum_{\lambda \in \dot{S}} \gamma^\lambda \prod_{k \in \lambda} X_k \right). \end{aligned}$$

Thus, essentially, what needs to be calculated is

$$\sum_{X_1} \exp \left(\sum_{\lambda \in S} \sum_{\text{s.t. } 1 \in \lambda} \gamma^\lambda \prod_{k \in \lambda} X_k \right). \quad (4.1)$$

Let

$$\text{Neigh}(X_i, S) = \{X_j \mid \exists \lambda \in S \text{ such that } \{i, j\} \subseteq \lambda\}.$$

Then, $\text{Neigh}(X_1, S)$ is the set of nodes with indexes appearing in (4.1). In fact, if we look at

$$\frac{1}{\bar{C}} \sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k$$

as the probability distribution for a Markov Random Field, then, $\text{Neigh}(X_1, S)$ is the set of neighboring nodes of X_1 . In a visualization of this Markov Random Field, this would be illustrated by an undirected arrow between each pair of neighbors. Also, this implies that for this Markov Random Field $G = (V, E)$, there is an edge e between the nodes X_i and X_j , denoted

$$e = \{X_i, X_j\} \in E$$

for each node $X_j \in \text{Neigh}(X_i, S)$. Recall from Section 2.3, that we can only have conditional dependencies between nodes X_i and X_j given all other Random Variables if X_i and X_j are neighbors.

Calculating (4.1) has a complexity of

$$\mathcal{O}(2^{|\text{Neigh}(X_i, S)|}).$$

Tjelmeland and Austad (2011) presents an algorithm which finds an approximate value of the sum with a complexity of $\mathcal{O}(2^\nu)$, for some input constant ν . That is, if

$$|\text{Neigh}(X_i, S)| \leq \nu,$$

the exact value of the sum is calculated. Otherwise, if

$$|\text{Neigh}(X_i, S)| > \nu,$$

the approximation

$$f \equiv \sum_{X_1} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right) \approx \sum_{X_1} \exp \left(\sum_{\lambda \in \tilde{S}} \tilde{\gamma}^\lambda \prod_{k \in \lambda} X_k \right) \equiv \tilde{f}$$

is made for some \tilde{S} and some collection of coefficients $\{\tilde{\gamma}^\lambda\}_{\lambda \in \tilde{S}}$ such that

$$|\text{Neigh}(X_1, \tilde{S})| \leq \nu.$$

The approximating function \tilde{f} for the function f is chosen to minimize the error sum of squares

$$SSE(f, \tilde{f}) = \sum_X \left(f(X) - \tilde{f}(X) \right)^2.$$

Now, the operation of summing out a Random Variable has a complexity which is bounded by $\mathcal{O}(2^\nu)$ for the input constant ν .

Also, in Tjelmeland and Austad (2011), a couple of possible ways to calculate bounds on the error are discussed. These bounds together with the approximate value of the sum, gives upper and lower bounds on the exact value. What is referred to as "Optimal bounds for pseudo-Boolean functions" in Tjelmeland and Austad (2011) is already implemented, and will be used for the tests later in this report. This computer program, implemented by Tjelmeland and Austad, will be referred to as "The AMRF Programs", where AMRF is an abbreviation for Approximate Markov Random Field. For a short user oriented presentation of "The AMRF Programs", the reader is referred to Appendix B.

5 Mathematical Formulation of the Problem

Given a graphical model with n nodes numbered from 1 to n , we assume that each node i is associated with a Boolean Random Variable X_i . Following the notation from Tjelmeland and Austad (2011), let

$$N = \{1, 2, \dots, n\},$$

and also, let

$$\Omega = \{0, 1\}^n$$

be the set of all Boolean vectors of length n . The vector

$$X = [X_i]_{i=1}^n,$$

containing all the Random Variables X_i in our graphical model, is then a Random Variable in the probability space

$$(\Omega, 2^\Omega, \mathbb{P}),$$

where \mathbb{P} is the Joint Probability Distribution on the graph.

Observe that the function

$$\dot{j} : \Omega \rightarrow 2^N \quad \text{such that} \quad \dot{j}(v) = \{ j \mid v_j = 1 \}$$

is one-to-one and onto, and thus defines a bijection between vectors in Ω and subsets of N . This allows us to associate Ω with the power set 2^N , and talk about their corresponding elements. That is, a given subset of N contains the indexes of the 1s in the corresponding vector in Ω .

Definition 5.1. *A decision is a deterministic Boolean vector Z of length n , that is, with each entry corresponding to one of the Random Variables X_i in our graphical model.*

Let u be an integer, and also, let $\Gamma \subseteq N$ be the indexes of a subset of all nodes in the graph. Define

$$\Omega^{\Gamma,u} = \left\{ Z \in \Omega : Z_i = 0 \ \forall i \notin \Gamma, \quad \text{and} \quad \sum_{i=1}^n Z_i \leq u \right\},$$

to be our decision space. That is, for a decision Z , we only allow the entries Z_i with index $i \in \Gamma$ to take on the value 1, i.e. all other entries have to be 0. Also, no more than u entries are allowed to have the value 1. Note that $\Omega^{\Gamma,u} \subseteq \Omega$.

Given a function

$$f : \Omega^{\Gamma,u} \rightarrow \mathbb{R},$$

we look for a decision $Z^* \in \Omega^{\Gamma,u}$ that maximizes the value of $f(Z)$. That is,

$$Z^* = \arg \max_{Z \in \Omega^{\Gamma,u}} \{f(Z)\}.$$

We will refer to such a decision Z^* as the optimal decision according to the function f .

Assume we want to look for nodes i where $X_i = 1$, and that it is not possible to observe all nodes. In fact, assume that there is only a subset of the nodes for which it is possible to observe the value of X_i , and also that there is an upper limit of how many nodes we can observe. We let the value of the entries Z_i in the decision vector Z denote whether we are going to check the value of the corresponding Random Variable, as in

$$Z_i = \begin{cases} 1 & \text{if we observe the value of } X_i, \\ 0 & \text{if we don't observe the value of } X_i. \end{cases}.$$

Then, $\Gamma \subseteq N$ contains the indexes of the nodes that are observable, and u is the maximum number of nodes we are allowed to check. Also, for our graphical

model which encodes probabilistic information about the unknown variables in X , we assume

$$f(Z) = \mathbb{E}h(X, Z) = \int_{\Omega} h(X, Z)\mathbb{P}(dX)$$

for some function

$$h : \Omega \times \Omega^{\Gamma, u} \rightarrow \mathbb{R}.$$

Let

$$c : \Omega^{\Gamma, u} \rightarrow \mathbb{R}^+$$

represent the cost of a decision. That is, $c(Z)$ is the cost associated with observing X_i for each i such that $Z_i = 1$. Also, let

$$w : \Omega \times \Omega^{\Gamma, u} \rightarrow \mathbb{R}^+$$

be the income of a decision for a specific assignment to the Random Variables in the graph. That is $w(X, Z)$ is the income associated with the observations according to the decision Z if the Random Variables had the assignment as in X . Now, we let $h(X, Z) = w(X, Z) - c(Z)$ and get

$$\begin{aligned} f(z) &= \mathbb{E}(\text{profit} |_{\text{following } Z}) \\ &= \mathbb{E}(w(X, Z) - c(Z)) \\ &= \mathbb{E}w(X, Z) - c(Z) \\ &= \sum_X w(X, Z)\mathbb{P}(X) - c(Z). \end{aligned}$$

This is the function f for which we want to find the optimal decision Z^* , and we will refer to this as the optimal decision Z^* , and assume this choice of f to be implicit.

As in Martinelli et al. (2011), the graphical model could be a Directed Acyclic Graph which denotes the causal conditional distributions for the presence of hydrocarbons in certain areas. We would let $X_i = 1$ denote presence of hydrocarbons in the area associated with node X_i , and correspondingly, $X_i = 0$ would denote absence of hydrocarbons. The nodes with indexes in Γ would represent the Potential Drilling Sites, while the other nodes are used to impose geological causal dependencies. Then, u is the maximum number of nodes we are allowed to drill. Also, $c(Z)$ would be the cost of drilling the Potential Drilling Sites with a nonzero entry in Z . Correspondingly, $w(X, Z)$ would be the net income after the finds of hydrocarbons according to X and Z . That is, we would find hydrocarbons in node X_i if and only if $X_i = Z_i = 1$.

6 Mathematical Formulation of the Solution

Typically, it is too time consuming to calculate the exact value of $f(Z)$ for all Z s in our decision space $\Omega^{\Gamma,u}$. That is why we have to look at approximations. Since we still want to make sure that we actually find the optimal decision, we look at upper and lower bounds for the value of $f(Z)$ for each Z . An algorithm for finding upper and lower bounds for the normalization constant C of a given probability distribution

$$\mathbb{P}(X_1, \dots, X_n) = \frac{1}{C} \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right),$$

as in Tjelmeland and Austad (2011), is already implemented. This will be used as a "Black Box" for finding upper and lower bounds on the value of $f(Z) = \mathbb{E}h(X, Z)$ for a given Z . Recall from Section 4, that finding the expected value of a function h on some Random Variables X is essentially the same problem as finding the normalization constant for a distribution $P(X)$ that is not normalized. Letting some set of upper and lower bounds be implicit, this lets us define the functions

$$f^+ : \Omega^{\Gamma,u} \rightarrow \mathbb{R}$$

and

$$f^- : \Omega^{\Gamma,u} \rightarrow \mathbb{R}.$$

We let $f^+(Z)$ denote the upper bound for the expected profit, and correspondingly, $f^-(Z)$ denote the lower bound.

Given a set of bounds, we have to use the values of $f^+(Z)$ and $f^-(Z)$ for all Z s to separate out as few candidates for the optimal decision as possible. By the definition of the optimal decision Z^* , we know that

$$f(Z^*) \geq f(Z) \quad \forall Z \in \Omega^{\Gamma,u}.$$

Also, for the upper and lower bounds, $f^+(Z)$ and $f^-(Z)$, respectively, we know that

$$f^-(Z) \leq f(Z) \leq f^+(Z) \quad \forall Z \in \Omega^{\Gamma,u}. \quad (6.1)$$

From the two above equations, we deduce that

$$f(Z^*) \geq f^-(Z) \quad \forall Z \in \Omega^{\Gamma,u},$$

which again implies

$$f(Z^*) \geq \max_{Z \in \Omega^{\Gamma, u}} f^-(Z).$$

Since (6.1) is true for any decision $Z \in \Omega^{\Gamma, u}$, more specifically we also have

$$f^-(Z^*) \leq f(Z^*) \leq f^+(Z^*),$$

which finally leaves us with

$$f^+(Z^*) \geq \max_{Z \in \Omega^{\Gamma, u}} f^-(Z).$$

Thus, we know that any decision \tilde{Z} with

$$f^+(\tilde{Z}) < \max_{Z \in \Omega^{\Gamma, u}} f^-(Z).$$

is not a candidate for the optimal decision. This also tells us that if we could find a \dot{Z} such that

$$f^-(\dot{Z}) \geq f^+(Z) \quad \forall Z \in \Omega^{\Gamma, u} \setminus \{\dot{Z}\},$$

we would know that $Z^* = \dot{Z}$. In general, the upper and lower bounds are not tight enough to separate out only one candidate for Z^* , but hopefully we would be able to exclude a significant fraction of the possible decisions Z . Note that it is sufficient to observe that

$$f^+(\tilde{Z}) < f^-(\dot{Z}).$$

for some decisions $\tilde{Z}, \dot{Z} \in \Omega^{\Gamma, u}$ to rule out \tilde{Z} as a candidate for the optimal decision.

Now, we have a plan for how to deal with a set of evaluations of the upper and lower bounds. Our focus will now be on how to obtain such bounds. As defined in Section 5, we assume that we are given a cost function $c(Z)$ and an income function $w(X, Z)$. Recall from Section 3, that the cost function can be represented as a pseudo-Boolean function.

$$c(Z) = \sum_{\lambda \in T_{cost}} \alpha_{cost}^\lambda \prod_{k \in \lambda} Z_k, \quad (6.2)$$

where T_{cost} is some subset of the power set $2^{\Omega^{\Gamma, \nu}}$. Since we only observe the values of X in the nonzero entries of Z , it is natural to assume that w is a

function of the scalar product $X \cdot Z$, where $(X \cdot Z)_i = X_i \cdot Z_i$. Then, we can write w as a pseudo-Boolean function of the following form,

$$w(X, Z) = \sum_{\lambda \in T_{inc}} \alpha_{inc}^\lambda \prod_{k \in \lambda} X_k \cdot Z_k. \quad (6.3)$$

Since we have assumed the income function to be a positive function $w(X, Z) > 0$ of the decision Z and the Random Variables X , we also let S_{inc} denote a collection of sets of indexes and $\{\beta_{inc}^\lambda\}_{\lambda \in S_{inc}} \subset \mathbb{R}$ denote coefficients such that

$$w(X, Z) = \exp \left(\sum_{\lambda \in S_{inc}} \beta_{inc}^\lambda \prod_{k \in \lambda} X_k \cdot Z_k \right). \quad (6.4)$$

Recall that Section 3.2 presented an algorithm for how to convert between the representations in (6.3) and (6.4). As in Section 3.3, we also write the probability distribution of the Random Variables X in our graphical model as

$$\mathbb{P}(X) = \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} X_k \right). \quad (6.5)$$

We introduce a variable \tilde{X} , which holds information from both the Random Variable X and the decision Z . In fact, we let \tilde{X} be a vector of length $2n$, where the first n entries are the entries in X , and the next n entries are the entries in Z . That is, for the Random Variable X and a decision Z , element-wise we define

$$\tilde{X}_i = \begin{cases} X_i & \text{if } i \leq n, \\ Z_{i-n} & \text{if } i > n. \end{cases} \quad (6.6)$$

Observe for a decision Z , $\tilde{X}(X, Z)$ is a Random Variable where

$$\mathbb{P} \left(\tilde{X} = [x, z] \right) = \begin{cases} \mathbb{P}(X = x) & \text{if } z \neq Z, \\ 0 & \text{if } z = Z, \end{cases}$$

for Boolean vectors x, z of length n . Thus, implicitly letting the choice of Z deterministically fill out the last n entries of \tilde{X} , and also letting $\tilde{X}_k(X, Z)$ denote the k th entry of $\tilde{X}(X, Z)$, (6.5) can be rewritten as

$$\mathbb{P} \left(\tilde{X}(X, Z) \right) = \exp \left(\sum_{\lambda \in S} \gamma^\lambda \prod_{k \in \lambda} \tilde{X}_k(X, Z) \right),$$

where the collection S of index sets λ is unchanged.

Correspondingly, (6.4) can be translated to

$$w(X, Z) = \exp \left(\sum_{\lambda \in S_{inc}} \beta_{inc}^\lambda \prod_{k \in \lambda} \left(\tilde{X}_k(X, Z) \cdot \tilde{X}_{n+k}(X, Z) \right) \right).$$

We define

$$\theta : 2^N \rightarrow 2^{\{1, \dots, 2n\}} \quad \text{such that} \quad \theta(\lambda) = \lambda \cup \{j + n \mid j \in \lambda\},$$

and let

$$\tilde{S}_{inc} = \{\theta(\lambda) \mid \lambda \in S_{inc}\}.$$

Also, for each $\lambda \in S_{inc}$, we let

$$\tilde{\beta}_{inc}^{\theta(\lambda)} = \beta_{inc}^\lambda.$$

This allows us to define

$$\tilde{w} : \{0, 1\}^{2n} \rightarrow \mathbb{R} \quad \text{such that} \quad \tilde{w}(\tilde{X}) = \exp \left(\sum_{\lambda \in \tilde{S}_{inc}} \tilde{\beta}_{inc}^\lambda \prod_{k \in \lambda} \tilde{X}_k \right).$$

Note all of this is just to get

$$\tilde{w}(\tilde{X}(X, Z)) = w(X, Z).$$

Assuming $\tilde{S}_{inc} = S$, as we could ensure by including zero-valued coefficients if necessary, the expected value $\mathbb{E}w(X, Z)$ can be written as

$$\begin{aligned} \mathbb{E}w(X, Z) &= \sum_X w(X, Z) \mathbb{P}(X) \\ &= \sum_X \tilde{w}(\tilde{X}(X, Z)) \mathbb{P}(\tilde{X}(X, Z)) \\ &= \sum_X \exp \left(\sum_{\lambda \in S} \left(\tilde{\beta}_{inc}^\lambda + \gamma^\lambda \right) \prod_{k \in \lambda} \tilde{X}_k(X, Z) \right) \end{aligned} \quad (6.7)$$

Assume we have the representation of $w(X, Z)$ as in (6.4) and the representation of $\mathbb{P}(X)$ as in (6.5). After some modifications, as illustrated in the functions defined for the augmented variable \tilde{X} , these can be plugged into "The

AMRF Programs”, which will return upper and lower bounds, respectively, on $\mathbb{E}w(X, Z)$ according to the input constant ν controlling the complexity of the algorithm. The obvious upper and lower bounds, respectively, on $f(Z)$ would then be the corresponding bound on $\mathbb{E}w(X, Z)$ minus the value of $c(Z)$. This way of calculating the value of $f(Z)$ corresponds to the equation

$$f(Z) = \mathbb{E}w(X, Z) - c(Z),$$

as we saw it in Section 5. We will refer to this method as to Method 1.

Section 5 also presented the equivalent formula

$$f(Z) = \mathbb{E}(w(X, Z) - c(Z)),$$

which introduces a slightly different way of finding bounds on $f(Z)$. That is, here, we want ”The AMRF Programs” to work with the function

$$h(X, Z) = w(X, Z) - c(Z)$$

and return bounds on $f(Z)$ directly. However, $h(X, Z)$ is not a positive function, so we have to add a sufficient constant M , to get

$$h(X, Z) + M > 0,$$

and thus be able to write

$$h(X, Z) + M = \exp \left(\sum_{\lambda \in \tilde{S}} \beta^\lambda \prod_{k \in \lambda} \tilde{X}_k(X, Z) \right)$$

for some collection of index sets \tilde{S} and corresponding collection of coefficients $\{\beta^\lambda\}_{\lambda \in \tilde{S}}$. This will be referred to as Method 2. Note that adding the constant M does not change the optimal decision, since

$$\max_{Z \in \Omega^{\Gamma, u}} \{f(Z) + M\} = \max_{Z \in \Omega^{\Gamma, u}} \{f(Z)\} + M,$$

and thus

$$\arg \max_{Z \in \Omega^{\Gamma, u}} \{f(Z) + M\} = \arg \max_{Z \in \Omega^{\Gamma, u}} \{f(Z)\}.$$

7 Generating Bayesian Networks

To test our solution method, which is presented in Section 6, we need Bayesian Networks with corresponding cost and income functions. We want to be able to easily generate several test cases, and we want to be able to do this for different sizes of the graph. We also want to study cases where observing a node is relatively cheap compared to the possible incomes, and cases where it is expensive. We also want to be able to control the sparseness of the graph, i.e. the density of edges between the nodes.

The problem discussed in Section 5 assumes that we are given a graphical model $G = (X, E)$ with n nodes represented as entries in the vector

$$X = [X_1, \dots, X_n],$$

with a corresponding Joint Probability Distribution

$$\mathbb{P}(X_1, \dots, X_n).$$

For this graphical model, we also define a decision vector

$$Z = [Z_1, \dots, Z_n],$$

with each entry Z_i corresponding to the Random variable X_i . It also assumes a cost function $c(Z)$ and an income function $w(X, Z)$. Thus, we need to generate sets

$$\{ G = (X, E), \mathbb{P}(X), c(Z), w(X, Z) \}.$$

to test our solution.

We will assume that our graphical model G is a Bayesian Network, and that the observable nodes are the leaf nodes in the graph. Thus, Γ is the indexes of the leaf nodes. Also, we assume the numbering i of the nodes X_i to be according to some topological ordering where any root node is preceding all non root nodes and any leaf node is succeeding all non leaf nodes. We adapt the terminology from Martinelli et al. (2011), and refer to the root nodes as Kitchens (K), the leaf nodes as Segments (S), and the intermediate nodes as Prospects (P). A Bayesian Network with $\#K$ Kitchens, $\#P$ Prospects and $\#S$ Segments, is therefore a graphical model with

$$n = \#K + \#P + \#S$$

nodes. Our numbering of the nodes plus the above terminology, introduce the following re-naming of the nodes X_i ,

$$X_i \rightarrow \begin{cases} K_i, & 1 \leq i \leq \#K \\ P_{i-\#K}, & \#K < i \leq \#K + \#P \\ S_{i-\#K-\#P}, & \#K + \#P < i \leq \#K + \#P + \#S \end{cases},$$

which results in

$$X = [K_1, K_2 \cdots, K_{\#K}, P_1, P_2, \cdots, P_{\#P}, S_1, S_2, \cdots, S_{\#S}].$$

7.1 The Directed Acyclic Graph

The first step is to generate the graph $G = (X, E)$. We let the number of Kitchens $\#K$, the number of Prospects $\#P$, the number of Segments $\#S$ and the total number of edges $\#E$ in the graph be deterministic. What is generated, is the start and end point of each edge. By definition, each Kitchen has only out-edges, that is, it can only be the start point of an edge. Also, we assume that any edge from a Kitchen has to enter a Prospect. This implies that

$$\forall K_i \quad \text{Ch}(K_i) \subseteq \{ P_1, P_2, \cdots, P_{\#P} \}.$$

We also assume

$$\forall K_i \quad P_i \in \text{Ch}(K_i), \tag{7.1}$$

which implies that we need to have

$$\#K \leq \#P.$$

Correspondingly, by definition, the Segments can only have in-edges, that is, they can only be end points of an edge. More specifically, we assume that each Segment has exactly one parent, which needs to be a Prospect. That is,

$$\forall S_i \quad \exists P_j \text{ such that } \text{Pa}(S_i) = \{P_j\}. \tag{7.2}$$

We also assume that among the children of a Prospect, there is at least one Segment. That implies that we need to have

$$\#P \leq \#S.$$

The fact that each Prospect has at least one Segment among its children, ensures that no Prospects are leaf nodes. The property in (7.1) ensures that no Kitchens

are leaf nodes either. Correspondingly, the property in (7.2) ensures that the Segments are not root nodes. To ensure that no Prospects are root nodes, in addition to the edges in (7.1), we need at least one parent for each Prospect P_i with index $i > \#K$. By ensuring that all Prospects and Segments are not root nodes, we get that

$$\#E \geq \#P + \#S.$$

Also, there is an upper limit to the possible number of edges. Actually,

$$\#E = \text{The number of edges with a Prospect as end point} + \#S.$$

Since the numbering of the nodes is assumed to follow some topological ordering, we know that

$$\forall P_i \quad \text{Pa}(P_i) \subseteq \{ K_1, K_2, \dots, K_{\#K}, P_1, P_2, \dots, P_{i-1} \},$$

which implies that

$$|\text{Pa}(P_i)| \leq \#K + i - 1.$$

Thus, the number of edges with a Prospect as end point is bounded,

$$\sum_{i=1}^{\#P} |\text{Pa}(P_i)| \leq \sum_{i=1}^{\#P} (\#K + i - 1) = \#P \left(\#K + \frac{\#P - 1}{2} \right).$$

To sum up the discussion, we get for the number of edges $\#E$

$$\#P + \#S \leq \#E \leq \#P \left(\#K + \frac{\#P - 1}{2} \right) + \#S,$$

and for the number of nodes of each kind, $\#K$, $\#P$, $\#S$, we have

$$1 \leq \#K \leq \#P \leq \#S = n - \#K - \#P.$$

An example graph according to these rules, can be seen in Figure 6.

Generating the edge set E sums up to the following algorithm. Start with $E = \emptyset$, then add edges as follows

- Add edge (K_i, P_i) for each $i \leq \#K$.
- Add edge (X, P_i) for an X drawn uniformly from

$$\{ K_1, K_2, \dots, K_{\#K}, P_1, P_2, \dots, P_{i-1} \}$$

for each $\#K < i \leq \#K + \#P$.

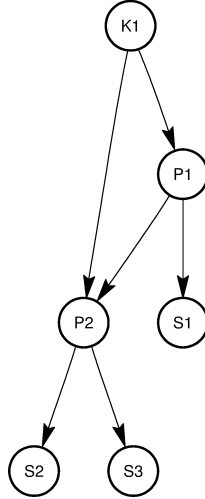


Figure 6: An example plot of a Directed Acyclic Graph with one Kitchen, two Prospects and three Segments.

- Add edge (X,Y) , for an X drawn uniformly from all Kitchens and Prospects

$$\{ K_1, K_2, \dots, K_{\#K}, P_1, P_2, \dots, P_{\#P} \}$$

and a Y drawn uniformly from the Prospects succeeding X in the ordering. Repeat until the current edge set E has size $\#E - \#S$.

- Start by assuming that each Prospect has one Segment as a child. Then draw a Prospect uniformly, and add one to its number of Segment children. Repeat until the total number of Segment children is $\#S$. Then for each $j \leq \#S$, add edge (P_i, S_j) where i starts at $i = 1$ and is increased by 1 whenever P_i has the correct number of Segment children.

7.2 The Joint Probability Distribution

The Joint Probability Distribution $\mathbb{P}(X_1, \dots, X_n)$ for the Bayesian Network over the Random Variables X_1, \dots, X_n , is assumed to be of the same form as in Section 3.3.1. We also assume $p_{X_i, X_j}^0 = \epsilon$ for each edge e_{X_i, X_j} . That is, what we need to specify to get the full Joint Probability Distribution, is the probability p_{K_i} for each Kitchen K_i , and the probability p_{X_i, X_j}^1 for each edge e_{X_i, X_j} , in

addition to the value of the parameter ϵ . The probabilities are all drawn from a beta-distribution. The parameters for the beta-distribution, namely α, β , are constant within all kitchen probabilities p_{K_i} drawn, and correspondingly, the parameters are constant within all edge probabilities p_{X_i, X_j}^1 drawn.

For the hydrocarbon example, as in Martinelli et al. (2011), we let a node X_i correspond to some geographical area. The Kitchen nodes represent places where hydrocarbons can be generated. Then, p_{K_i} is the probability that it has been generated hydrocarbons there. That is, $K_i = 1$ corresponds to the event that hydrocarbons have been generated in the given area, and $K_i = 0$ corresponds to no hydrocarbons generated there. The edges in the graph corresponds to possible directed paths for the hydrocarbons to flow. The probability p_{X_i, X_j}^1 corresponding to edge e_{X_i, X_j} , is then the probability for hydrocarbons to flow from X_i to X_j assuming that there are hydrocarbons in X_i . The Segment nodes are the Potential Drilling Sites, and they will contain hydrocarbons, corresponding to $S_j = 1$, if hydrocarbons have arrived along at least one directed path in the graph, starting at some Kitchen K_i with $K_i = 1$.

The Joint Probability Distribution for the hydrocarbon example coincides with the one discussed in Section 3.3.1. However, the above discussion makes it tempting to set $p_{X_i, X_j}^0 = 0$, since no hydrocarbon in a node makes it impossible to have hydrocarbons flowing from it. But actually assuming $p_{X_i, X_j}^0 = 0$ for some edge e_{X_i, X_j} , introduces assignments to the Random Variables of probability 0. Then, the Joint Probability Distribution cannot be expressed as the exponential of a pseudo-Boolean function, as we want. Thus, we introduce a small $\epsilon > 0$, and deterministically set $p_{X_i, X_j}^0 = \epsilon$ for all edges e_{X_i, X_j} . Then, the pseudo-Boolean representation of the Joint Probability Distribution is calculated as for the example Network in Section 3.3.1.

7.3 The Cost and Income functions

With the graph $G = (V, E)$ and the Joint Probability Distribution $\mathbb{P}(X)$ in hand, the last step is to generate the cost function $c(Z)$ and the income function $w(X, Z)$. We will assume a pseudo-Boolean representation of the cost function and the income function, as in (6.2) and (6.3), respectively. Note that in our setting, we have

$$\Gamma = \{\#K + \#P + 1, \#K + \#P + 2, \dots, \#K + \#P + \#S\},$$

that is, the observable nodes are exactly the leaf nodes, which equals the set of Segments, placed in the end in the topological ordering by choice. Since we

study decisions $Z \in \Omega^{\Gamma, u}$, we let

$$T_{inc}, T_{cost} \subseteq 2^{\Gamma}.$$

Define

$$\vartheta : \mathbb{N} \rightarrow \mathbb{N} \quad \text{such that} \quad \vartheta(m) = \#K + \#P + m,$$

so that for each Segment S_i , we have $S_i \equiv X_{\vartheta(i)}$. Then, we assume a model where observing a Segment S_i is associated with an exploring cost $\alpha_{cost}^{\{\vartheta(i)\}}$, and also where observing two sibling Segments S_i, S_j is associated with some savings $\alpha_{cost}^{\{\vartheta(i), \vartheta(j)\}}$. That is, exploring only S_i has a total cost of

$$\alpha_{cost}^{\emptyset} + \alpha_{cost}^{\{\vartheta(i)\}},$$

and exploring both S_i and S_j has a total cost of

$$\alpha_{cost}^{\emptyset} + \alpha_{cost}^{\{\vartheta(i)\}} + \alpha_{cost}^{\{\vartheta(j)\}} - \alpha_{cost}^{\{\vartheta(i), \vartheta(j)\}}.$$

Correspondingly, observing $S_i = 1$ is associated with some gross income $\alpha_{inc}^{\{\vartheta(i)\}}$, and due to savings we associate some extra gross income $\alpha_{cost}^{\{\vartheta(i), \vartheta(j)\}}$ to observing $S_i = S_j = 1$ for two sibling Segments S_i, S_j . That is, we assume that there is a discount effect corresponding to the costs of extracting the gain from two sibling Segments, which results in a higher gross income. This, and also the savings for the cost function, could represent savings caused by a partly shared infrastructure. However, we also have to ensure that no combination of any size corresponds to a free exploration. Higher order interactions are assumed not to be present, for simplicity.

We have assumed that for each Segment S_i

$$\alpha_{cost}^{\{\vartheta(i)\}}, \alpha_{inc}^{\{\vartheta(i)\}} \geq 0,$$

and also, that for each pair of sibling Segments S_i, S_j

$$\min\{\alpha_{cost}^{\{\vartheta(i)\}}, \alpha_{cost}^{\{\vartheta(j)\}}\} \geq -\alpha_{cost}^{\{\vartheta(i), \vartheta(j)\}} \geq 0, \quad \alpha_{inc}^{\{\vartheta(i), \vartheta(j)\}} \geq 0.$$

The model assumes that cost and income value interactions occur only between Segments S_i, S_j that are siblings in the graph representation, i.e. children of the same Prospect P_k ,

$$\{S_i, S_j\} \subseteq \text{Ch}(P_k).$$

That is, we set

$$T_{inc} = T_{cost} = \{\emptyset\} \cup \{\{i\} \mid i \in \Gamma\} \cup \{\{i, j\} \mid i, j \in \Gamma, \text{Pa}(X_i) = \text{Pa}(X_j)\}.$$

Without loss of generality, we set $\alpha_{inc}^{\emptyset} = \alpha_{cost}^{\emptyset} = 1$. Then, for each Segment S_i , the coefficient $\alpha_{cost}^{\{\vartheta(i)\}}$ is drawn from a Gamma distribution with mean μ_{cost} and standard deviation σ_{cost} , and correspondingly, the coefficient $\alpha_{inc}^{\{\vartheta(i)\}}$ is drawn from a Gamma distribution with mean μ_{inc} and standard deviation σ_{inc} . That is,

$$\alpha_{cost}^{\{\vartheta(i)\}} \sim \text{Gamma}(\mu_{cost}, \sigma_{cost}),$$

and

$$\alpha_{inc}^{\{\vartheta(i)\}} \sim \text{Gamma}(\mu_{inc}, \sigma_{inc}).$$

Then, for each pair of sibling Segments S_i, S_j with k other Segment siblings, we assume

$$u_{i,j} \sim \frac{1}{k+1} \text{Unif}(0, 1),$$

and set

$$\alpha_{cost}^{\{\vartheta(i), \vartheta(j)\}} = -u_{i,j} \cdot \min\{\alpha_{cost}^{\{\vartheta(i)\}}, \alpha_{cost}^{\{\vartheta(j)\}}\}$$

and we also draw $\alpha_{inc}^{\{\vartheta(i), \vartheta(j)\}}$ from a Gamma distribution, according to

$$\alpha_{inc}^{\{\vartheta(i), \vartheta(j)\}} \sim \text{Gamma}\left(\frac{\mu_{inc}}{10}, \frac{\sigma_{inc}}{10}\right).$$

8 First Set of Tests: BN1

The first set of tests were done according to Method 1, as introduced in Section 6, on a test case generated as described in Section 7. We represent the income function $w(X, Z)$ as the exponential of a pseudo-Boolean function, by use of formulas from Section 3.2. Combining this with the representation of the probability distribution, we get a representation for $w(X, Z)\mathbb{P}(X)$. Then, we calculate upper and lower bounds, respectively, on $\mathbb{E}w(X, Z)$ as in (6.7), by use of "The AMRF Programs" introduced in Section 4. This computer program sums out the Random Variables according to the numbering of the nodes. The first Random Variable to be summed out is K_1 and the last is $S_{\#S}$. From the resulting bound functions, we try to rule out candidates for the optimal decision, as described in Section 6.

We will present a Bayesian Network with 4 Kitchens, 7 Prospects and 10 Segments. A visualization of the graph can be seen in Figure 7. Corresponding

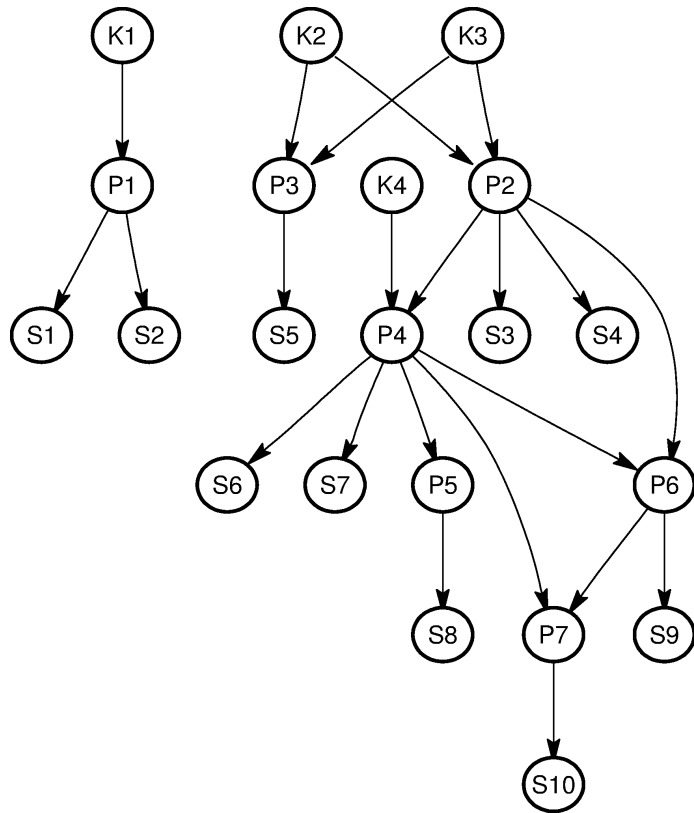


Figure 7: BN1, the Bayesian Network for first set of tests

to this graph, we have a Joint Probability Distribution, which is characterized by the numbers in Table 14 in Section C.1. This graph, with its corresponding Joint Probability Distribution, will be named BN1. Two sets consisting of a cost function $c(Z)$ and an income function $w(X, Z)$, are also generated, and can be found in Tables 15 and 16, respectively, also in Section C.1. For the first set, denoted by a , the parameters of the distribution for generating the coefficients were set to

$$\mu_{\text{cost},a} = 2000, \quad \sigma_{\text{cost},a} = 2000, \quad \mu_{\text{inc},a} = 4500, \quad \sigma_{\text{inc},a} = 1200.$$

Correspondingly, for the second set, denoted by b , the parameters were set to

$$\mu_{\text{cost},b} = 4500, \quad \sigma_{\text{cost},b} = 4000, \quad \mu_{\text{inc},b} = 4500, \quad \sigma_{\text{inc},b} = 4000.$$

When we refer to BN1a, we think of BN1 with the cost and income functions denoted by a , and correspondingly for BN1b. The parameters for drawing the Joint Probability Distribution were set to

$$\alpha_{K_i} = 6.0, \quad \beta_{K_i} = 2.0, \quad \alpha_{e_{i,j}} = 5.0, \quad \beta_{e_{i,j}} = 2.0,$$

and the ϵ parameter was given the value $\epsilon = 0.01$. Recall that the Joint Probability Distribution is exactly the same for test a and b .

Given a cost function and an income function, we calculate bounds on $f(Z)$ corresponding to an input constant ν for "The AMRF Programs". Actually, we calculate bounds for each

$$\nu \in \{11, 12, 13, 14, 15, 16, 17, 18, 19\}, \quad (8.1)$$

and thus we get nine pairs consisting of upper bound functions f^+ and lower bound functions f^- , respectively, for each set a, b of cost function and income function. We let $f_{\nu,i}^+$ denote the upper bound function obtained for the given value of ν for cost and income function i , and correspondingly for $f_{\nu,i}^-$. That is, ν is as in (8.1), and $i \in \{a, b\}$. For each pair $f_{\nu,i}^+, f_{\nu,i}^-$, we evaluate the values $f_{\nu,i}^+(Z), f_{\nu,i}^-(Z)$ for each

$$Z \in \Omega^{\Gamma,u} \quad \text{for} \quad u = 5.$$

That is, we evaluate the upper and lower bounds for each decision that involves observing a maximum of five Segments. For each pair (ν, i) , the resulting number of candidates for the optimal decision can be found in Tables 1 and 2 for a and b , respectively.

Decision size	Total number of decisions	Number of accepted decisions for $f_{\nu,a}^+, f_{\nu,a}^-$ for $\nu =$								
		11	12	13	14	15	16	17	18	19
0	1	0	0	0	0	0	0	0	0	0
1	10	6	6	6	3	2	1	0	0	0
2	45	40	42	39	35	30	24	17	9	0
3	120	119	119	116	111	100	85	64	36	0
4	210	210	210	209	205	196	175	140	84	0
5	252	252	252	252	251	246	232	197	127	1
Sum	638	627	629	622	608	574	517	388	256	1

Table 1: Number of accepted decisions as a function of ν for the income function and cost function from Table 15.

Decision size	Total number of decisions	Number of accepted decisions for $f_{\nu,b}^+, f_{\nu,b}^-$ for $\nu =$								
		11	12	13	14	15	16	17	18	19
0	1	1	0	0	0	0	0	0	0	0
1	10	6	7	7	3	2	1	0	0	0
2	45	41	42	39	36	31	25	18	10	1
3	120	119	119	116	110	100	85	64	36	0
4	210	210	210	209	205	195	175	140	84	0
5	252	252	252	252	251	246	231	196	126	0
Sum	638	629	630	623	605	517	418	388	256	1

Table 2: Number of accepted decisions as a function of ν for the income function and cost function from Table 16.

Note that even though this is a quite small graph, we need $\nu = 19$ to find the optimal decision, that is, for the bounds $f_{\nu,i}^+, f_{\nu,i}^-$ to leave out only one candidate. This is also the value of ν that ensures exact calculation of $f(Z)$. That is, $\nu_{exact} = 19$ corresponds to no approximations for this graph, and we get

$$f_{19,i}^+ \equiv f_{19,i}^- \quad \text{for} \quad i = a, b.$$

Also note that exact calculation will only leave candidates that actually are optimal decisions. That is, if we are left with more than one decision after exact calculations, these must all correspond to the same value of f , that is, the maximal value of f within $\Omega^{\Gamma,u}$. Thus, all decisions left are optimal. Hence, needing $\nu = 19$ to separate out one candidate is actually the worst case for this graph. Also, note that $\nu = 18$ leaves 256 candidates for the optimal decision in both cases $i = 1, 2$, from a total of 638 decisions in $\Omega^{\Gamma,u}$. That is, reducing the value of ν with 1, and thus reducing the asymptotic complexity in run time from $\mathcal{O}(2^{\nu_{exact}})$ to $\mathcal{O}(2^{\nu_{exact}-1})$, leaves us with roughly half of our initial decision space still. A further reduction of the value of ν leads to at least 418 candidates in all cases, which corresponds to a reduction by less than 35% of the candidates. That is, we observe that we need a quite large value of ν to get useful results for these examples.

Also note that for the first test, denoted a , the optimal decision implies observing 5 nodes. In the next, b , the costs are higher, which results in an optimal decision with only 3 observations.

The examples presented illustrates the general observations from tests according to Method 1. As soon as approximations were made on our function $w(X, Z)\mathbb{P}(X)$, we end up with a large fraction of the initial decision space $\Omega^{\Gamma,u}$ left. Thus, we had to come up with a revised strategy to be able to find the optimal decision more effectively on larger graphs.

9 Second Set of Tests

The tests of Method 1 indicated that we needed to rethink our strategy for finding the optimal decision. First of all, we had to look for any unnecessary approximations. Observe that we can take advantage of (2.7) in Section 2.1. That is, previously, when we calculated the expected value of the function $w(X, Z)$, we took the expected value with respect to the full Joint Probability Distribution. That is, if the biggest maximal clique in the graph had a size greater than the current value of ν in the calculations, in general, the approximations made would introduce error terms for the whole graph.

However, the form we assume for the cost and income functions, presented in Section 7.3, allows us to divide the cost and income functions into parts corresponding to each Prospect. Let X_{B_j} denote Segment children of Prospect P_j , that is

$$X_{B_j} = \{S_i \in \text{Ch}(P_j) \mid \vartheta(i) \in \Gamma\},$$

and also, let Z_{B_j} denote the corresponding entries in the decision vector. This allows us to write the cost function as

$$c(Z) = \sum_{j=1}^{\#P} c_j(Z_{B_j})$$

and the income function as

$$w(X, Z) = \sum_{j=1}^{\#P} w_j(X_{B_j}, Z_{B_j}).$$

In fact, let

$$T_k = \{\emptyset\} \cup \{\{i\} \mid i \in \Gamma, X_i \in \text{Ch}(P_k)\} \cup \{\{i, j\} \mid i, j \in \Gamma, X_i, X_j \in \text{Ch}(P_k)\}.$$

Then,

$$T_{inc} = T_{cost} = \cup_{j=1}^{\#P} T_j,$$

and for positive integers $j, k \leq \#P$

$$j \neq k \quad \Leftrightarrow \quad T_k \cap T_j = \{\emptyset\}.$$

Thus,

$$w(X, Z) + (\#P - 1)\alpha_{inc}^{\emptyset} = \sum_{j=1}^{\#P} \sum_{\lambda \in T_j} \alpha_{inc}^{\lambda} \prod_{k \in \lambda} X_k Z_k,$$

and

$$c(Z) + (\#P - 1)\alpha_{cost}^{\emptyset} = \sum_{j=1}^{\#P} \sum_{\lambda \in T_j} \alpha_{cost}^{\lambda} \prod_{k \in \lambda} Z_k.$$

If we ignore the multiples of $\alpha_{cost}^{\emptyset}$ and α_{inc}^{\emptyset} added, respectively, we get

$$w_j(X_{B_j}, Z_{B_j}) = \sum_{\lambda \in T_j} \alpha_{inc}^{\lambda} \prod_{k \in \lambda} Z_k$$

and

$$c_j(Z_{B_j}) = \sum_{\lambda \in T_j} \alpha_{cost}^\lambda \prod_{k \in \lambda} Z_k.$$

Also note that our choice of

$$\alpha_{cost}^\emptyset = \alpha_{inc}^\emptyset = 1$$

actually implies

$$w(X, Z) - c(Z) \equiv \sum_{j=1}^{\#P} (w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j})).$$

Thus,

$$\begin{aligned} f(Z) &= \mathbb{E}(w(X, Z) - c(Z)) \\ &= \mathbb{E}\left(\sum_{j=1}^{\#P} (w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j}))\right) \\ &= \sum_{j=1}^{\#P} \mathbb{E}(w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j})). \end{aligned} \tag{9.1}$$

Assume that $j \leq \#P$ is a positive integer. It is when calculating

$$\mathbb{E}(w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j})),$$

we get to use the observations from (2.7). Actually, let

$$A_j = \{P_j\} \cup \text{Anc}(P_j) \cup B_j,$$

and observe that A_j meets the requirements set in (2.2). Thus,

$$\mathbb{E}(w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j})) = \sum_{X_i \in A_j} (w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j})) \mathbb{P}(X_{A_j}).$$

Let

$$M_j = 1 + \max_{Z_{B_j}} c_j(Z_{B_j}),$$

define

$$S_j = \left\{ \lambda \mid \{\tilde{X}_i \mid i \in \lambda\} \in 2^{X_{B_j} \cup Z_{B_j}} \right\},$$

and let $\{\beta_j^\lambda\}_{\lambda \in S_j}$ be constants such that

$$0 < w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j}) + M_j = \exp\left(\sum_{\lambda \in S_j} \beta_j^\lambda \prod_{k \in \lambda} \tilde{X}_k\right).$$

Note that we identify each variable \tilde{X}_i with an entry in X or Z , respectively, according to (6.6).

Finally, let

$$M = \sum_{j=1}^{\#P} M_j,$$

and note that (9.1) implies

$$\begin{aligned} f(Z) + M &= \mathbb{E}(w(X, Z) - c(Z) + M) \\ &= \mathbb{E}\left(\sum_{j=1}^{\#P} (w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j}) + M_j)\right) \\ &= \sum_{j=1}^{\#P} \mathbb{E}(w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j}) + M_j) \\ &= \sum_{j=1}^{\#P} \left(\sum_{X_i \in A_j} \mathbb{P}(X_{A_j}) \exp\left(\sum_{\lambda \in S_j} \beta_j^\lambda \prod_{k \in \lambda} \tilde{X}_k\right)\right). \end{aligned}$$

Note that the core idea of Method 2 was introduced by adding the constants M_j . Also note that, in practice, the above equation tells us to calculate $f(Z) + M$ by adding the contributions corresponding to each prospect. That is, each of those terms are obtained similarly as for Method 1 in Section 8, but now from the distribution $\mathbb{P}(X_{A_j})$ and the function

$$w_j(X_{B_j}, Z_{B_j}) - c_j(Z_{B_j}) + M_j.$$

Also note that for any positive integer $j \leq \#P$,

$$w_j(X_{B_j}, \vec{0}) = \alpha_{inc}^\emptyset = 1 = \alpha_{cost}^\emptyset = c_j(\vec{0}),$$

and thus

$$f(\vec{0}) = 0.$$

The tests for Method 2 show that we need the graphs to be sparse for our upper and lower bounds to be tight enough to eliminate a significant fraction of the possible decisions $Z \in \Omega^{\Gamma,u}$. This section presents tests where the approximations work well, but it also presents a description of how it fails for an example Bayesian Network. Also in these tests, "The AMRF Programs" sum out the Random Variables according to our numbering of the nodes, and the numbers we present for each test is as before. The strategy for finding the optimal decision is still the same, we just calculate the bounds in a slightly different way.

9.1 Sparse graphs: BN2-BN4

First, a set of tests on sparse graphs will be presented. Here, this means that the number of edges in the graph is kept close to the minimum according to our definitions in Section 7. These graphs will encode more conditional independence assumptions than if more edges were added, and the clique sizes in the corresponding undirected graph will be smaller than for a graph with more edges added. That means that "The AMRF Programs" will do less approximations, and it is more likely that our strategy will work as we hope.

Three single Bayesian Networks of different sizes will be presented. For each of these, we will study the number of accepted decisions Z as a function of the approximation constant ν . To help the discussion of these numbers, we also present the value of $\nu = \nu_{exact}(P_i)$ that ensures exact calculation of the term of $f(Z)$ corresponding to Prospect P_i .

For each of these three examples, the test is repeated 1000 times. That is, for each example, 1000 Bayesian Networks with cost function and income function are generated from the same distribution as for the corresponding example. From each of these replications, we will study the value of ν needed to find the optimal decision by our strategy, that is

$$\nu_{one} = \min_{\nu} \{ \nu \mid \text{Number of accepted decisions}(\nu) = 1 \}$$

We will also look at the value of ν needed to separate out just a few candidates. In fact, we study

$$\nu_{acceptable} = \min_{\nu} \{ \nu \mid \text{Number of accepted decisions}(\nu) \leq 7 \}.$$

These two numbers will be compared to the value of ν that ensures exact calculation of the expected gross income $f(Z)$ for the graph, that is

$$\nu_{exact} = \max_{\text{Prospect } P_i} \nu_{exact}(P_i).$$

Then also, for a given value of ν , and corresponding upper bounds f^+ and lower bounds f^- , respectively, let \tilde{Z} be the decision with the highest lower bound for expected income. That is,

$$\tilde{Z} = \arg \max_{Z \in \Omega^{f,u}} f^-(Z).$$

Define

$$L_\nu = \frac{f(Z^*) - f(\tilde{Z})}{f(Z^*)},$$

the percentage loss in exact expected income, when following the approximate optimal decision \tilde{Z} instead of the (exact) optimal decision Z^* . In each of the three sets of 1000 tests, the value of ν corresponding to the loss L_ν is chosen so that we expected (according to our best guesses) that about half of the tests in the set would have $\nu_{exact} > \nu$. That is, so that our program would have to make a guess for the optimal decision in about half the cases.

9.1.1 BN2

The following test describes a Bayesian Network with 7 Kitchens, 15 Prospects and 40 Segments. The number of extra edges added after the minimum value of 55, is 10. That is, the result is a graph with 62 nodes and 65 edges, and it is visualized in Figure 8. A corresponding visualization where only the Kitchens and the Prospects are shown, can be found in Figure 9. Observe that the undirected version of this Bayesian Network consists of two connected components, that is, the nodes $\{K2, P2, D2\}$ do not have edges to or from any of the other nodes not in this set. The Bayesian Network with corresponding cost and income functions is generated as described in Section 7. The parameters for drawing cost and income coefficients were set to

$$\mu_{\text{cost}} = 4000, \quad \sigma_{\text{cost}}^2 = 2500, \quad \mu_{\text{inc}} = 5000, \quad \sigma_{\text{inc}}^2 = 2500.$$

The parameters for drawing the Joint Probability Distribution were set to

$$\alpha_{K_i} = 6.0, \quad \beta_{K_i} = 2.0, \quad \alpha_{e_{i,j}} = 5.0, \quad \beta_{e_{i,j}} = 2.0,$$

and the ϵ parameter was given the value $\epsilon = 0.1$. For numbers fully describing the probability distribution and the cost and income functions, see Tables 17-32 in Appendix C.2.

Observe from Table 3 that we need $\nu = 11$ to get exact numbers, but Table 4 tells us that already from $\nu = 6$, we get only one candidate for the optimal

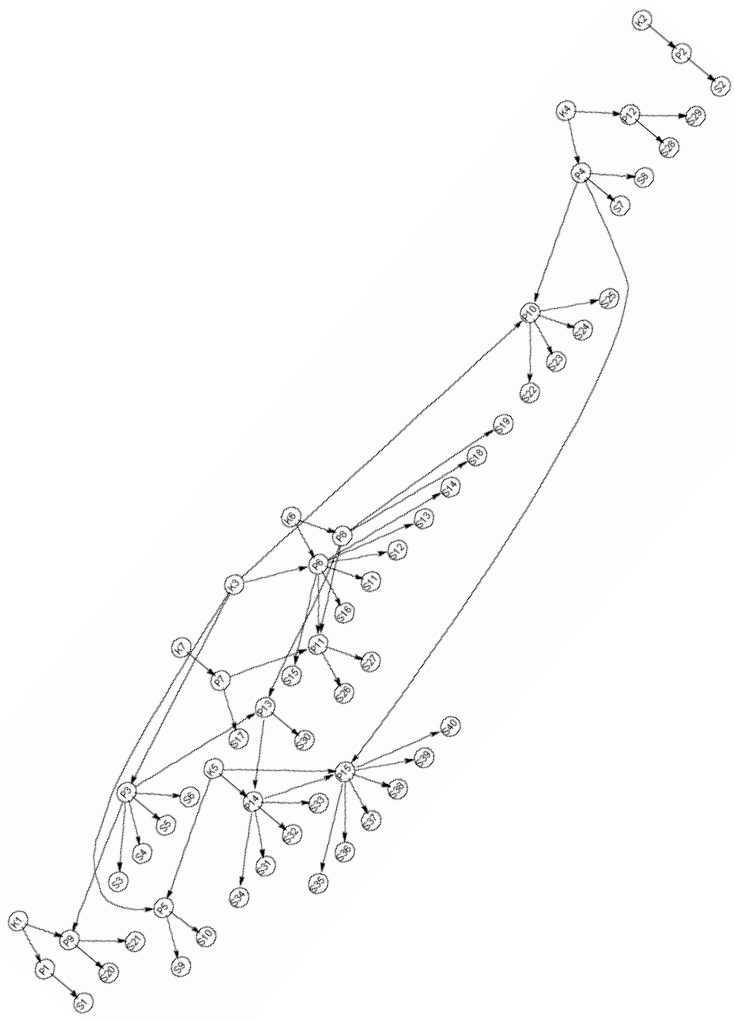


Figure 8: The Bayesian Network 2BN.

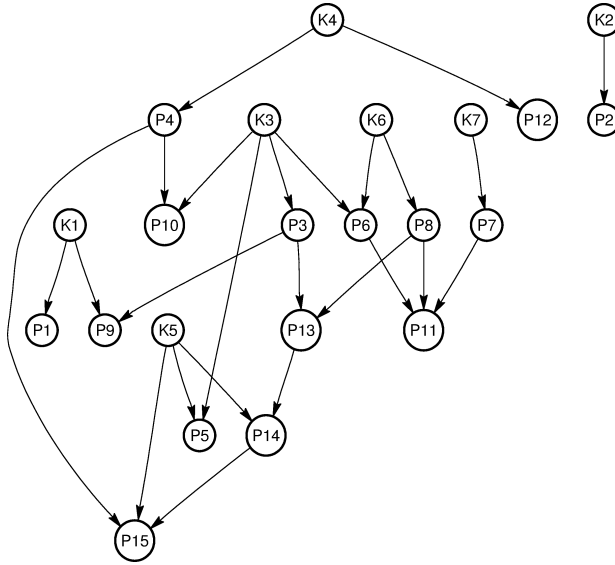


Figure 9: BN2: A visualization of only the Kitchens, the Prospects and the edges between them.

Prospect P_i	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
ν	1	1	7	3	3	11	1	3	3	7	3	3	2	7	11

Table 3: BN2: Value of ν that ensures exact calculation of expected gross income function $w_i - c_i + M_i$ corresponding to prospect P_i for the Bayesian Network in Figure 8.

Decision size	Total number of decisions	Number of accepted decisions for $\nu =$			
		1 – 3	4	5	6 – 11
0	1	1	1	0	0
1	40	40	40	0	0
2	780	780	770	1	1
3	9880	9880	9336	8	0
4	91390	91390	82081	28	0
5	658008	658008	567072	58	0
Sum	760099	760099	65930	95	1

Table 4: BN2: Number of accepted decisions as a function of ν .

decision. Even for $\nu = 5$, we get pretty good results. However, for $\nu \leq 3$, we are not able to eliminate any decisions, and $\nu = 4$ only allows us to eliminate 13% of the decisions. Observe from Table 3, that $3 \leq \nu \leq 6$ all introduces approximations for the functions corresponding to Prospects P_3, P_6, P_{10}, P_{14} and P_{15} , so the stepwise improvements of the results from $\nu = 3$ to $\nu = 6$ are all due to better approximations, and not to approximations on a smaller number of functions.

This test was replicated 1000 times. That is, 1000 Bayesian Networks with 7 Kitchens, 15 Prospects and 40 Segments were generated, with Joint Probability Distributions and cost and income functions drawn from the same distributions as those for BN2. Recall from Section 9.1, that ν_{exact} is the smallest value of the input constant ν that ensures "The AMRF Programs" to do exact calculations for each replication. Correspondingly, $\nu_{acceptable}$ is the smallest value of ν that leaves us with a maximum of 7 candidates for the optimal decision, and ν_{one} is the smallest value of ν that leaves us with only the optimal decision Z^* . The results of the 1000 replications are summed up in Figures 10, 11, 13, 12 and 14.

Observe from Figure 10, how the values for ν_{exact} are dominated by odd numbers. Actually, 931 tests have $\nu_{exact} \in \{7, 9, 11\}$. At the same time, Figure 11 tells us that for 70.5% of the tests, it is sufficient with $\nu = 6$ to separate out only one candidate for the optimal decision. In fact, 625 tests had $\nu_{one} \in \{5, 6\}$, while no tests had $\nu_{exact} \in \{5, 6\}$. Only one test had $\nu_{exact} = 4$, but actually, in 80 of the 1000 tests, $\nu = 4$ was sufficient to find the optimal decision. We observe from Figure 12 that 728 of the 1000 tests had a reduction of 3 or more from the value of ν_{exact} to the value of ν_{one} . There is even a test where the

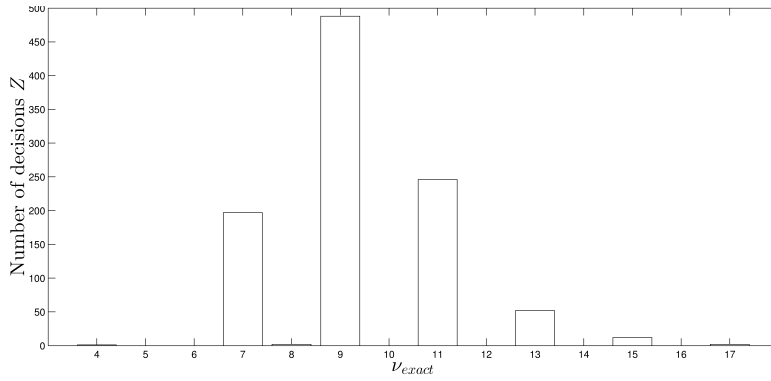


Figure 10: BN2: The value of ν needed to do exact calculations for the 1000 tests.

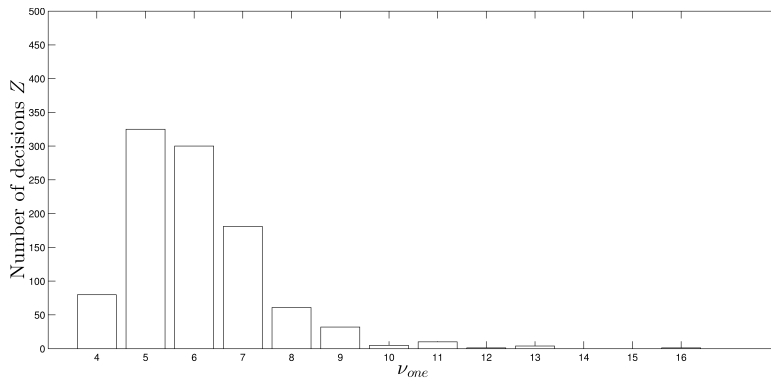


Figure 11: BN2: The value of ν needed to separate out only one candidate in each of the 1000 tests.

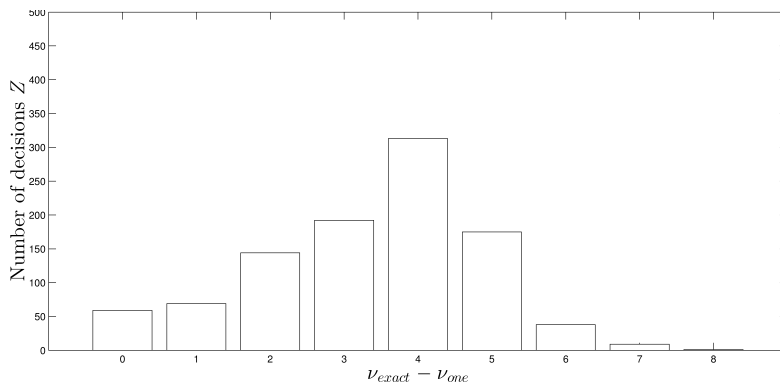


Figure 12: BN2: The difference between ν_{exact} and ν_{one} in each of the 1000 tests.

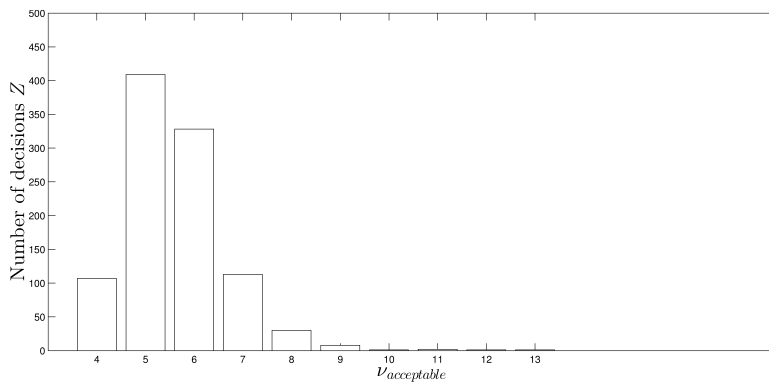


Figure 13: BN2: The value of ν needed to separate out a maximum of seven candidates in each of the 1000 tests.

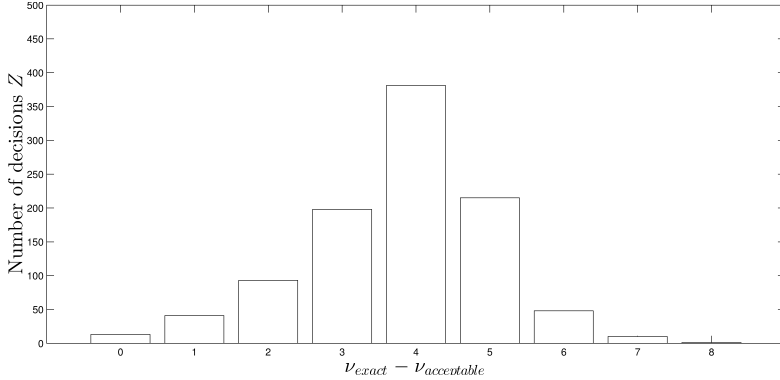


Figure 14: BN2: The difference between ν_{exact} and $\nu_{acceptable}$ in each of the 1000 tests.

value of this reduction is 8, and as much as 48 had a reduction of at least 6. Similarly, we observe from Figure 13, that for as much as 84.4% of the tests, it is sufficient with $\nu = 6$ to reduce the number of candidates for the optimal decision to seven or less. Actually, 737 tests had $\nu_{acceptable} \in \{5, 6\}$, and as much as 107 tests had $\nu_{acceptable} = 4$. Figure 12 shows how 853 of the 1000 tests had a reduction of 3 or more from ν_{exact} to the value of $\nu_{acceptable}$. As few as 13% of the tests had $\nu_{acceptable} = \nu_{exact}$, and 59 of the 1000 tests had a reduction of 6 or more.

For these 1000 replications, we chose $\nu = 6$, and studied the loss fraction L_6 . We got 40 cases where the loss fraction L_6 was nonzero. These came from the in total 295 cases where $\nu = 6$ was not enough to leave only one candidate for the optimal decision. Of these 295 cases, where a guess of the optimal decision had to be made, the average of L_6 was 0.019, with a standard deviation of 0.081. The smallest nonzero loss fraction was $3.905 \cdot 10^{-5}$, and the greatest loss fraction was as high as 0.8318. Actually, there are five cases where the loss fraction is higher than 0.3, and thus we get a large standard deviation. For a listing of all nonzero loss fractions, the reader is referred to Table 33 in Section C.2. In total, on all 1000 cases, that left us with an average loss $\bar{L}_6 = 0.0055$, with a standard deviation of 0.045.

Prospect P_i	1	2	3	4	5	6	7	8	9	10
ν	3	7	3	3	1	3	1	3	1	3
Prospect P_i	11	12	13	14	15	16	17	18	19	20
ν	3	3	1	3	3	1	3	3	1	3

Table 5: BN3: Value of ν that ensures exact calculation of expected gross income function $w_i - c_i + M_i$ corresponding to prospect P_i for the Bayesian Network in Figure 15.

9.1.2 BN3

The following test describes a Bayesian Network with 10 Kitchens, 20 Prospects and 40 Segments. The number of extra edges added after the minimum value of 60, is 10. That is, the result is a graph with 70 nodes and 70 edges, and it is visualized in Figure 15. Observe that the undirected version of this Bayesian Network consists of four connected components. The nodes in the leftmost connected component is visualized in Figure 16, and correspondingly, we get a closer look on the three rightmost connected components in Figure 17. The Bayesian Network with corresponding cost and income functions is generated as described in Section 7. The parameters for drawing cost and income coefficients were set to

$$\mu_{\text{cost}} = 3000, \quad \sigma_{\text{cost}}^2 = 2500, \quad \mu_{\text{inc}} = 5000, \quad \sigma_{\text{inc}}^2 = 2500.$$

The parameters for drawing the Joint Probability Distribution were set to

$$\alpha_{K_i} = 6.0, \quad \beta_{K_i} = 2.0, \quad \alpha_{e_{i,j}} = 5.0, \quad \beta_{e_{i,j}} = 2.0,$$

and the ϵ parameter was given the value $\epsilon = 0.01$. For numbers fully describing the probability distribution and the cost and income functions, see Tables 34-54 in Appendix C.3.

Observe from Table 5 that we need $\nu = 7$ to get exact numbers, but Table 6 tells us that already from $\nu = 3$, we are able to eliminate 99.8% of the possible decisions. In addition, we only need $\nu \geq 5$ to get just one candidate for the optimal decision. Also for $\nu = 4$, we are able to remove most of the candidates for the optimal decision. Note from Table 5 that for $\nu \in \{1, 2\}$, we calculate exact values for the functions corresponding to $P_5, P_7, P_9, P_{13}, P_{16}$ and P_{19} , but that is not enough to eliminate any candidates. However, for $\nu \geq 3$, approximations are only made on the function corresponding to P_2 , and as a result, the number

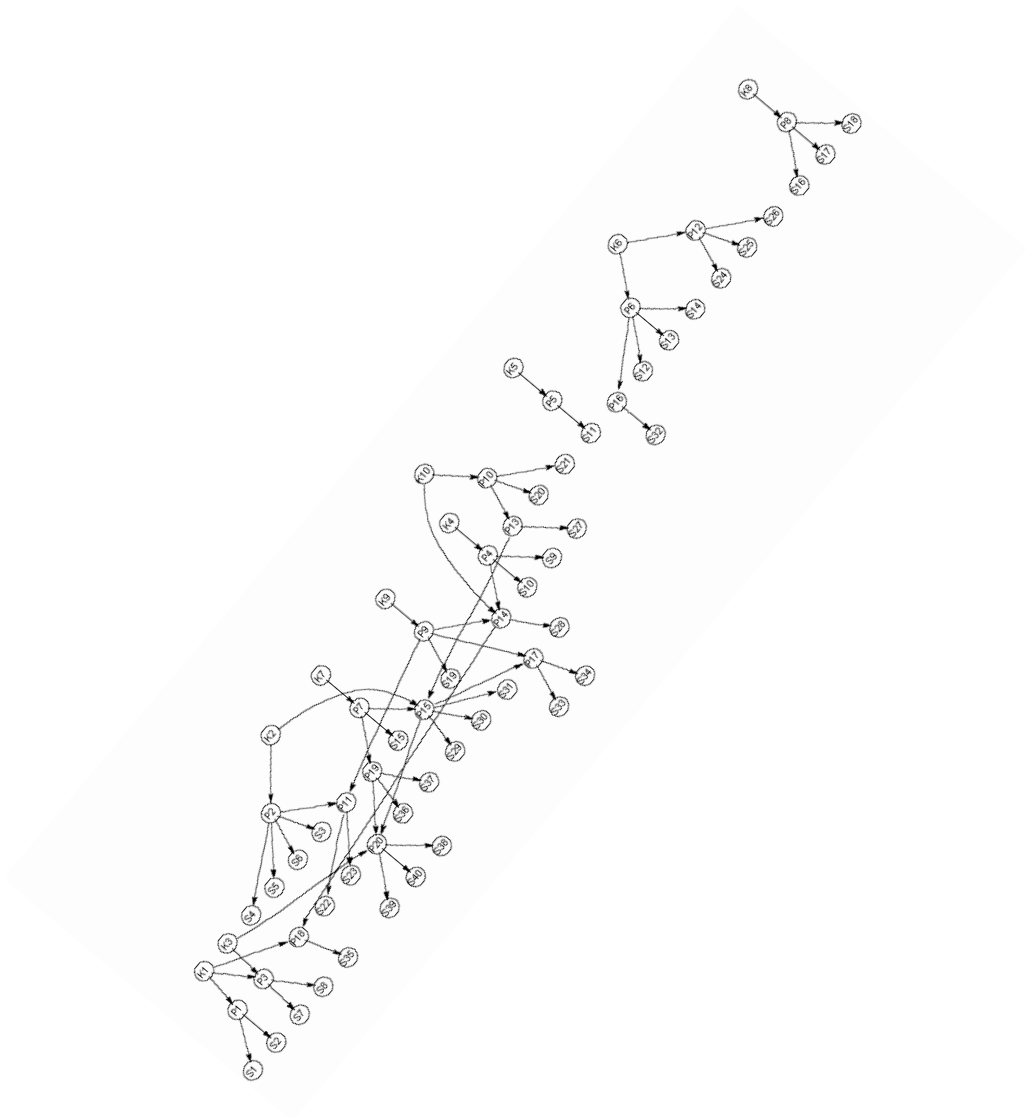


Figure 15: The Bayesian Network 3BN

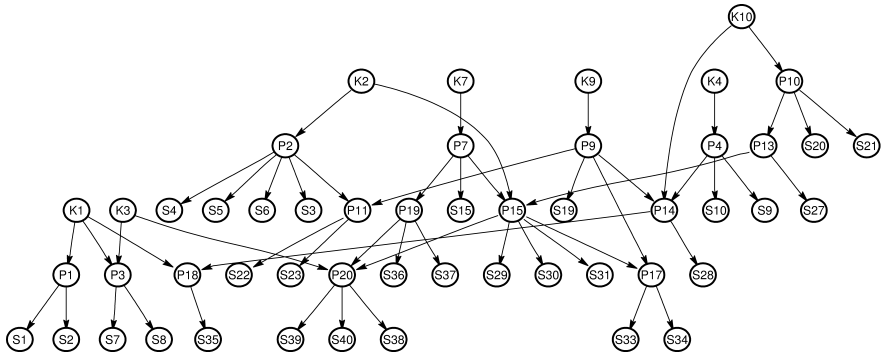


Figure 16: BN3: The left connected component of the Bayesian Network in Figure 15.

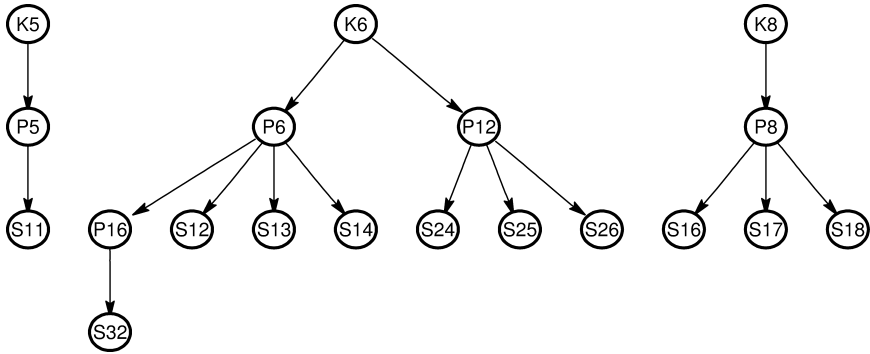


Figure 17: BN3: The right three connected components of the Bayesian Network in Figure 15.

Decision size	Total number of decisions	Number of accepted decisions for $\nu =$			
		1 – 2	3	4	5 – 7
0	1	1	0	0	0
1	40	40	0	0	0
2	780	780	1	0	0
3	9880	9880	22	0	0
4	91390	91390	212	1	0
5	658008	658008	1331	4	1
Sum	760099	760099	1566	5	1

Table 6: BN3: Number of accepted decisions as a function of ν .

of candidates are dropping rapidly from $\nu = 2$ to $\nu = 3$. However, we note that we still find the optimal decision for ν with value $\nu_{exact}(P_2) - 2$.

This test was replicated 1000 times, and the results are summed up in Figures 18, 19, 21, 20 and 22. Note that the Figures in this Section visualize the results for the same variables as the corresponding Figures in Section 9.1.1. Observe from Figure 18, how the values of ν_{exact} still are dominated by odd numbers. Actually, 814 tests have $\nu_{exact} \in \{7, 9\}$. At the same time, Figure 19 tells us that for 69.5% of the tests, it is sufficient with $\nu = 5$ to separate out only one candidate for the optimal decision. In fact, 782 tests had $\nu_{one} \in \{4, 5, 6\}$, while only 90 tests had $\nu_{exact} \in \{4, 5, 6\}$. We observe from Figure 20 that 541 of the 1000 tests had a reduction of 3 or more from the value of ν_{exact} to the value of ν_{one} . 63.4% of the cases had a reduction of either 2, 3 or 4. Also, 37 of the 1000 tests had a reduction of at least 5, but there are also 243 tests with no reduction. Similarly, we observe from Figure 21, that for as much as 80.5% of the tests, it is sufficient with $\nu = 5$ to reduce the number of candidates for the optimal decision to seven or less. Actually, 907 tests had $\nu_{acceptable} \in \{4, 5, 6\}$. Figure 20 shows how 651 of the 1000 tests had a reduction of 3 or more from the value of ν_{exact} to the value of $\nu_{acceptable}$. Also, 148 of the tests had $\nu_{acceptable} = \nu_{exact}$, a significant reduction from the 243 tests with $\nu_{one} = \nu_{exact}$. Also note that 45 of the tests had a reduction of 5 or more from the value of ν_{exact} to the value of $\nu_{acceptable}$.

For these 1000 tests, we chose $\nu = 5$, and got 27 cases where L_5 was nonzero. These came from the in total 305 cases where $\nu = 5$ was not enough to leave only one candidate for the optimal decision. Of these 305 cases, where a guess of the optimal decision had to be made, the average of L_5 was 0.0059, with a

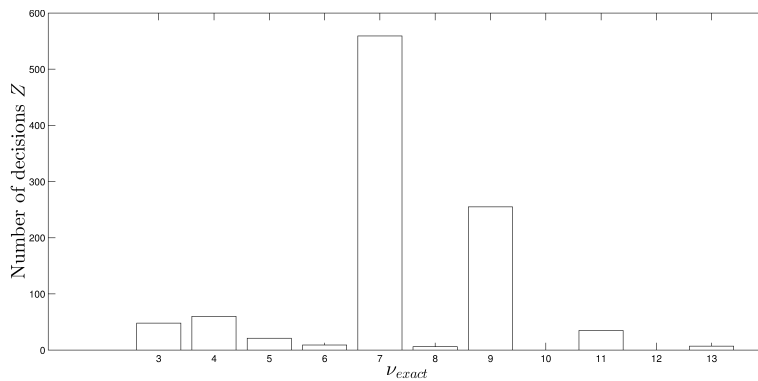


Figure 18: BN3: The value of ν needed to do exact calculations for the 1000 tests.

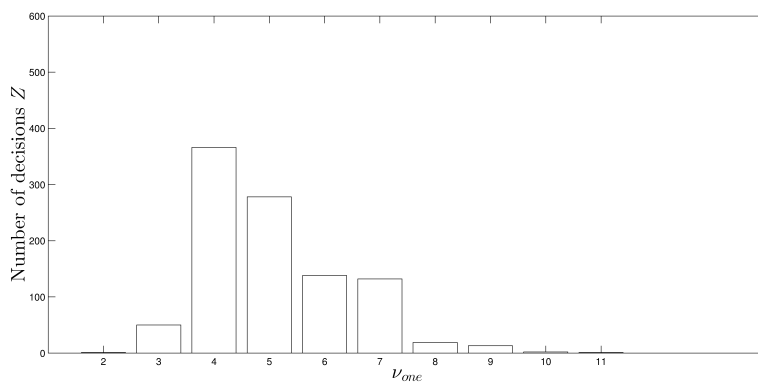


Figure 19: BN3: The value of ν needed to separate out only one candidate in each of the 1000 tests.

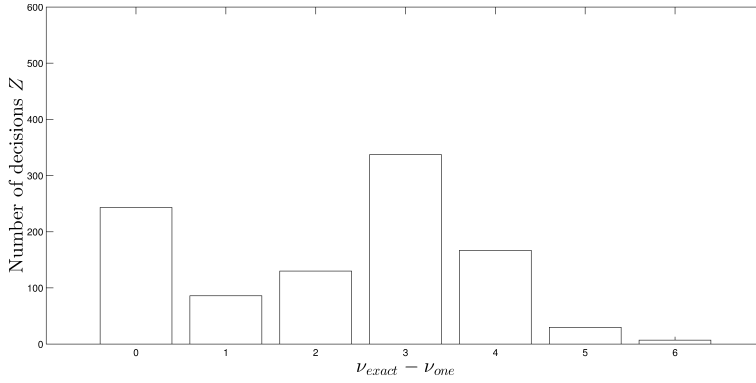


Figure 20: BN3: The difference between ν_{exact} and ν_{one} in each of the 1000 tests.

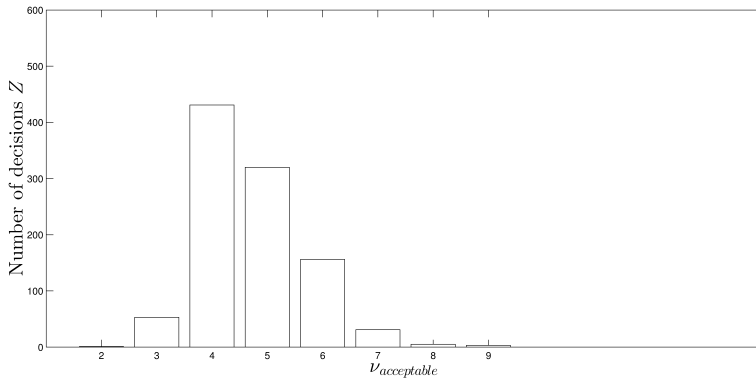


Figure 21: BN3: The value of ν needed to separate out a maximum of seven candidates in each of the 1000 tests.

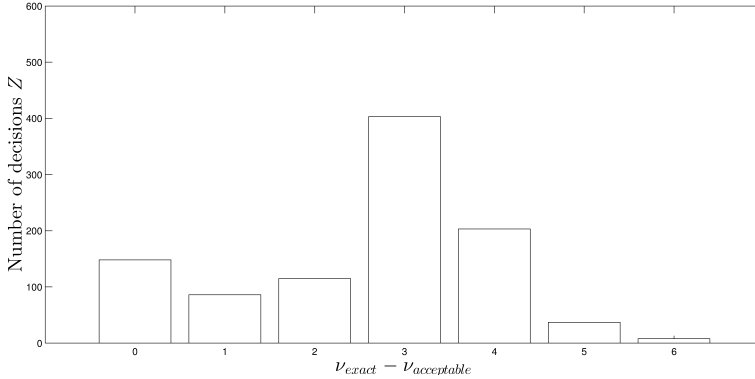


Figure 22: BN3: The difference between ν_{exact} and $\nu_{acceptable}$ in each of the 1000 tests.

standard deviation of 0.0354. The smallest nonzero loss fraction was $8.46 \cdot 10^{-4}$, and the greatest loss fraction was 0.4682. There was only three cases where the loss fraction was higher than 5%. For a listing of all nonzero loss fractions, the reader is referred to Table 55 in Section C.3. In total, on all 1000 cases, that left us with an average loss $\bar{L}_5 = 0.0018$, with a standard deviation of 0.0197.

9.1.3 BN4

The following test describes a Bayesian Network with 10 Kitchens, 20 Prospects and 60 Segments. The number of extra edges added after the minimum value of 60, is 20. That is, the result is a graph with 90 nodes and 80 edges, and it is visualized in Figure 23. A corresponding visualization where only the Kitchens and the Prospects are shown, can be found in Figure 24. Observe that the undirected version of this Bayesian Network is fully connected. That is, assuming all edges undirected, it is possible to find a path from any node to any other node. The Bayesian Network with corresponding cost and income functions is generated as described in Section 7. The parameters for drawing cost and income coefficients were set to

$$\mu_{\text{cost}} = 4000, \quad \sigma_{\text{cost}}^2 = 2500, \quad \mu_{\text{inc}} = 5000, \quad \sigma_{\text{inc}}^2 = 2500.$$

The parameters for drawing the Joint Probability Distribution were set to

$$\alpha_{K_i} = 6.0, \quad \beta_{K_i} = 2.0, \quad \alpha_{e_{i,j}} = 5.0, \quad \beta_{e_{i,j}} = 2.0,$$

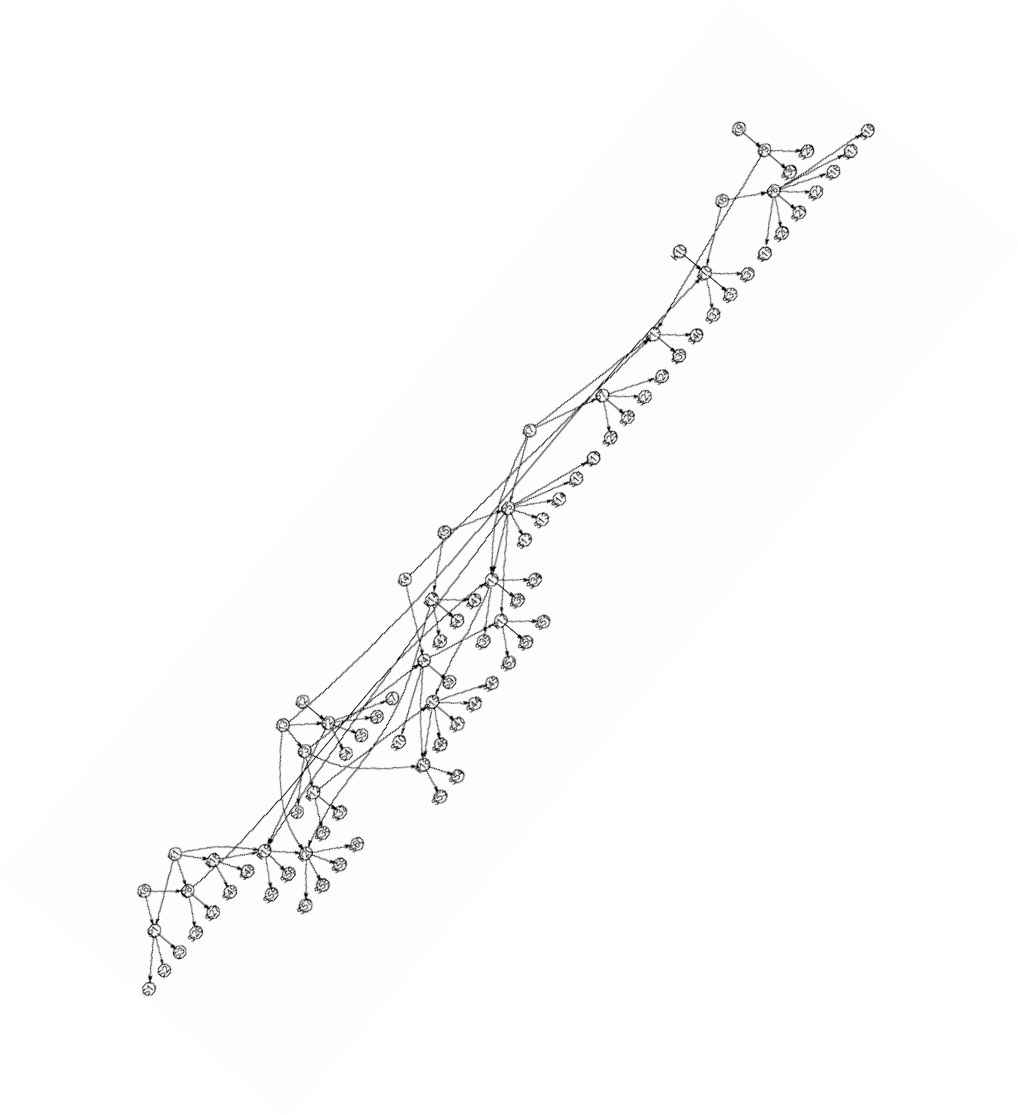


Figure 23: The Bayesian Network 4BN.

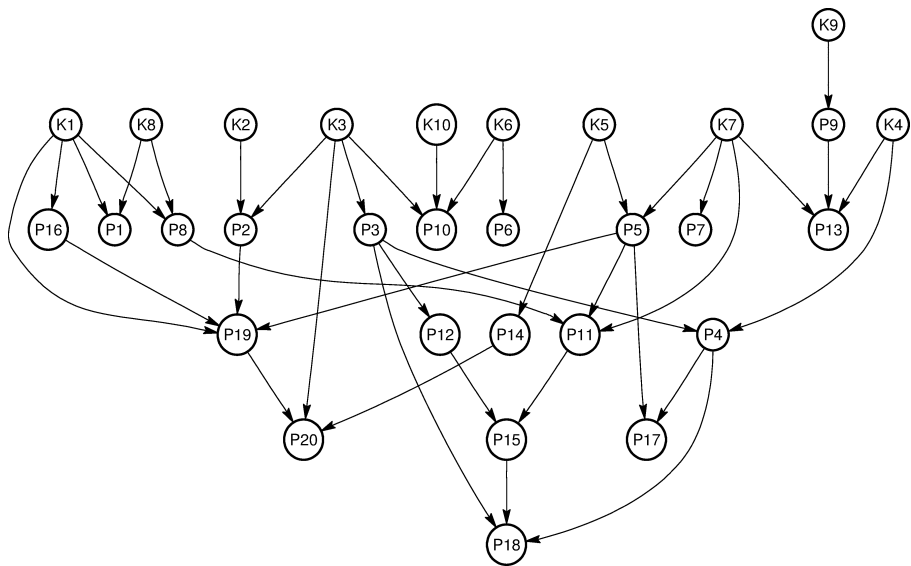


Figure 24: BN4: A visualization of only the Kitchens, the Prospects and the edges between them.

Prospect P_i	1	2	3	4	5	6	7	8	9	10
ν	3	7	1	3	9	13	7	3	3	3
Prospect P_i	11	12	13	14	15	16	17	18	19	20
ν	3	3	3	3	7	3	3	4	4	7

Table 7: BN4: Value of ν that ensures exact calculation of expected gross income function $w_i - c_i + M_i$ corresponding to prospect P_i for the Bayesian Network in Figure 23.

Decision size	Total number of decisions	Number of accepted decisions for $\nu =$			
		1 – 6	7	8	9 – 13
0	1	1	0	0	0
1	60	60	0	0	0
2	1770	1770	0	0	0
3	34220	34220	2	2	1
4	487635	487635	2	1	0
Sum	523686	523686	4	3	1

Table 8: BN4: Number of accepted decisions as a function of ν .

and the ϵ parameter was given the value $\epsilon = 0.01$. For numbers fully describing the probability distribution and the cost and income functions, see Tables 56-76 in Appendix C.4.

Observe from Table 7 that we need $\nu = 13$ to get exact numbers, but Table 8 tells us that already from $\nu = 7$, we are able to eliminate a great fraction of the possible decisions. In addition, we only need $\nu \geq 9$ to eliminate all non optimal decisions. Observe from Table 7 that for $\nu \geq 7$, we calculate exact values for the functions corresponding to all Prospects except P_5 and P_6 . Also, for $\nu \geq 9$, approximations are only made on the function corresponding to P_6 , and these are the same ν s that allows to eliminate all non optimal decisions.

This test was replicated 1000 times, and the results are summed up in Figures 25, 26, 28, 27 and 29. Note that the Figures in this Section visualizes the same variables as the corresponding Figures in Sections 9.1.1 and 9.1.2. Observe from Figure 25, that also here, the values for ν_{exact} are dominated by odd numbers. Actually, 899 tests have $\nu_{exact} \in \{9, 11, 13\}$. At the same time, Figure 26 tells us that for 47.9% of the tests, it is sufficient with $\nu = 7$ to separate out only one candidate for the optimal decision. In fact, 658 tests had $\nu_{one} \leq 8$, while

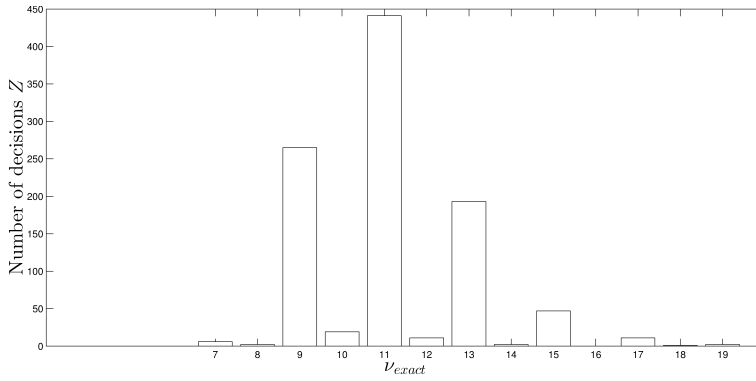


Figure 25: BN4: The value of ν needed to do exact calculations for the 1000 tests.

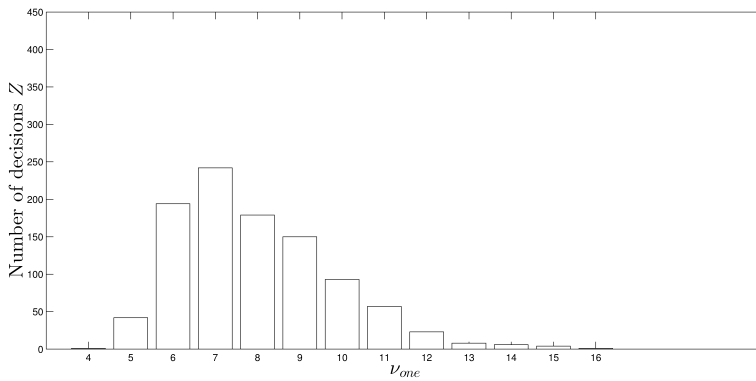


Figure 26: BN4: The value of ν needed to separate out only one candidate in each of the 1000 tests.

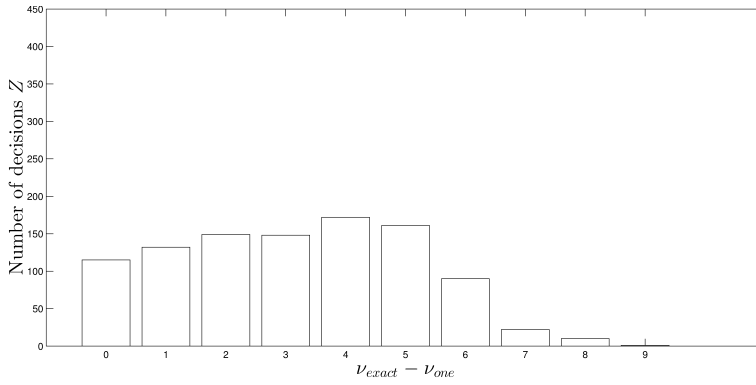


Figure 27: BN4: The difference between ν_{exact} and ν_{one} in each of the 1000 tests.

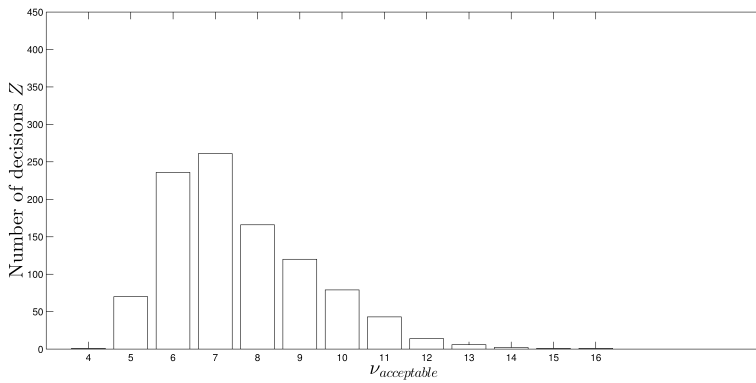


Figure 28: BN4: The value of ν needed to separate out a maximum of seven candidates in each of the 1000 tests.

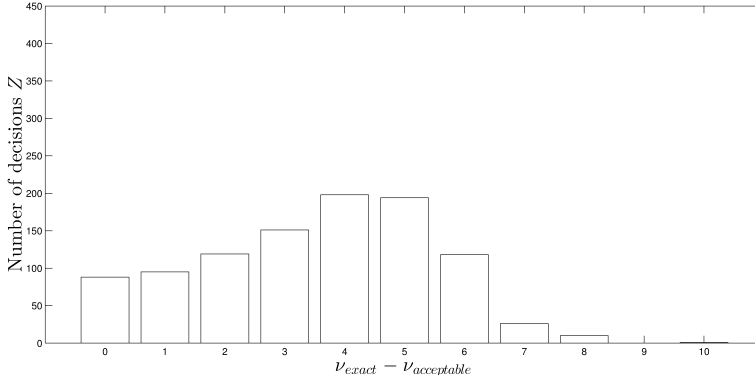


Figure 29: BN4: The difference between ν_{exact} and $\nu_{acceptable}$ in each of the 1000 tests.

only 8 tests had $\nu_{exact} \leq 8$. We observe from Figure 27 that 604 of the 1000 tests had a reduction of 3 or more from the value of ν_{exact} to the value of ν_{one} . 48.1% of the cases had a reduction of either 3, 4 or 5. Of the 1000 replications, 33 had a reduction of at least 7, but there are also 115 tests with no reduction from ν_{exact} to ν_{one} . Similarly, we observe from Figure 28, that for as much as 56.8% of the tests, it is sufficient with $\nu = 7$ to reduce the number of candidates for the optimal decision to seven or less. Actually, 734 tests had $\nu_{acceptable} \leq 8$. Figure 27 shows how 698 of the 1000 tests had a reduction of 3 or more from the value of ν_{exact} to the value of $\nu_{acceptable}$. We observe that 88 of the tests had $\nu_{acceptable} = \nu_{exact}$, a significant reduction from the 115 tests with $\nu_{one} = \nu_{exact}$. Note that 37 of the 1000 tests had a reduction of 7 or more from the value of ν_{exact} to the value of $\nu_{acceptable}$. Also note that for one of the tests, the difference between ν_{exact} and $\nu_{acceptable}$ is as high as 10.

Also, for the 1000 tests, we chose $\nu = 7$, and got 67 cases where L_7 was nonzero. These came from the in total 521 cases where $\nu = 7$ was not enough to leave only one candidate for the optimal decision. Of these 521 cases, where a guess of the optimal decision had to be made, the average of L_7 was 0.032, with a standard deviation of 0.12. The smallest nonzero loss fraction was $7.955 \cdot 10^{-6}$, and the greatest loss fraction was 0.9579. In about half of the nonzero cases, the loss fraction was higher than 15%, and actually in 13 cases, the loss fraction is higher than 50%. For a listing of all nonzero loss fractions, the reader is referred to Table 77 in Section C.4. In total, on all 1000 cases, that left us with

Prospect P_i	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
ν	2	3	3	7	3	3	7	3	7	7	3	7	6	16	17

Table 9: BN5: Value of ν that ensures exact calculation of expected gross income function corresponding to this prospect for the Bayesian Network in Figure 30.

an average loss $\bar{L}_7 = 0.017$, with a standard deviation of 0.092.

9.2 A less sparse graph: BN5

To put our solution to a bigger challenge, we increase the number of edges, in order to observe how and when the solution method fails. Now, we look at a Bayesian Network with 7 Kitchens, 15 Prospects, 40 Segments, and 40 edges more than the minimum of 55 edges for this set of nodes. That means a graph with in total 62 Nodes and 95 edges. A visualization of the Bayesian Network can be seen in Figure 30. Notice the web of edges between the nodes, both in Figure 30 and in Figure 31. The Bayesian Network with corresponding cost and income functions was generated as described in Section 7. The parameters for drawing cost and income coefficients were set to

$$\mu_{\text{cost}} = 4000, \quad \sigma_{\text{cost}}^2 = 2500, \quad \mu_{\text{inc}} = 5000, \quad \sigma_{\text{inc}}^2 = 2500.$$

The parameters for drawing the Joint Probability Distribution were set to

$$\alpha_{K_i} = 6.0, \quad \beta_{K_i} = 2.0, \quad \alpha_{e_{i,j}} = 5.0, \quad \beta_{e_{i,j}} = 2.0.$$

For numbers fully determining the probability distribution and the cost and income functions, see Tables 78-93 in Section C.5.

From Table 9, we observe that for $7 \leq \nu \leq 15$, only approximations on $\mathbb{E}w_{14} - c_{14}$ and $\mathbb{E}w_{15} - c_{15}$ are made. Correspondingly, for $\nu = 16$, only approximations on $\mathbb{E}w_{15} - c_{15}$ are made. For the fraction of accepted decisions for a given ν , see Table 10. The table clearly states that our approximations are not good enough. Any $\nu \leq 16$ implies getting a lower bound function that easily is strictly bounded by the smallest upper bound value. Hence, we are not able to eliminate any decisions, except when all calculations are done exactly. To study how and why this happens, we investigate the behavior of $\mathbb{P}(X_{A_{15}})$ as we sum out the Random Variables, one at a time. First, we calculate upper bounds on the log normalization constant for $\mathbb{P}(X_{A_{15}})$ for each $\nu \in \{10, \dots, 17\}$, to obtain the data found in Table 11. Observe from this Table, that for all ν s

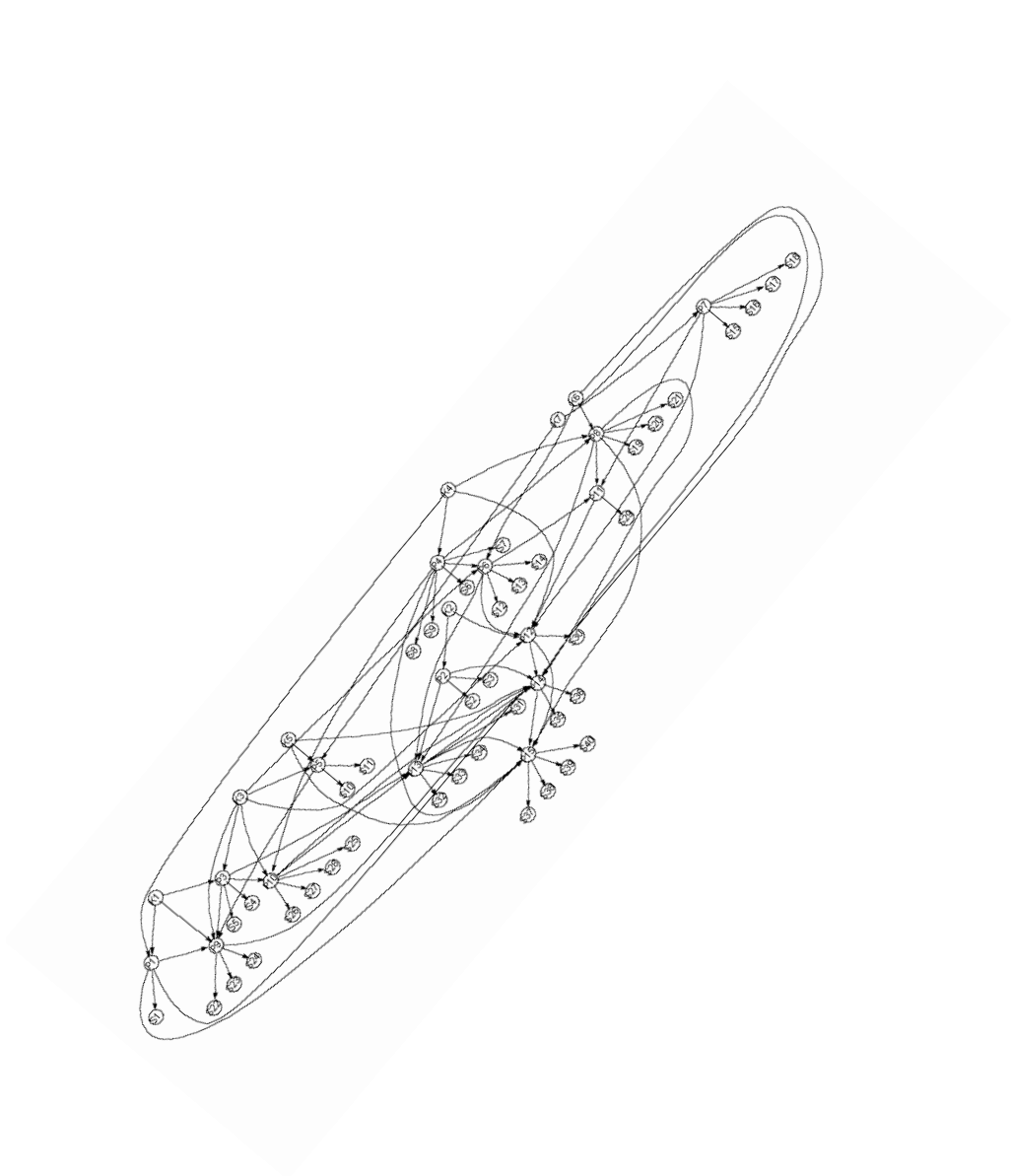


Figure 30: The Bayesian Network BN5.

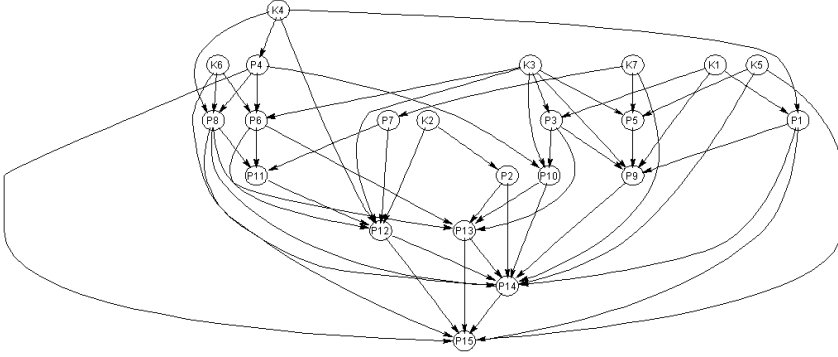


Figure 31: BN5: A visualization of only the Kitchens, the Prospects and the edges between them.

Decision size	Total number of decisions	Decisions accepted, $\nu =$	
		1 – 16	17
0	1	1	0
1	40	40	0
2	780	780	1
3	9880	9880	0
4	91390	91390	0
5	658008	658008	0
Sum	760099	760099	1

Table 10: BN5: Number of accepted decisions as a function of ν . Observe from Table 9 that we need $\nu = 17$ to get exact numbers, and that is the same value that is needed to eliminate any candidates for the optimal decision.

ν	Log normalization constant
10	193.930367
11	200.448060
12	231.803170
13	217.573379
14	243.259810
15	386.116269
16	92.7247710
17	-0.0000000

Table 11: BN5: Upper bounds on the log normalization constant for the probability distribution π_{15} calculated with different ν s. 17 is the smallest value of ν that ensures no approximations to be made, and hence $\nu = 17$ gives the exact value $0 = \ln 1$. For all other values of ν , the upper bounds calculated are useless.

except $\nu = 17$, we get upper bounds with enormous values. Note that $\nu = 17$ corresponds to exact calculations and thus, we obtain the exact normalization constant $1 = e^0$. Hence, the upper bounds are useless in this case. Also note that the normalization constant corresponds to some constant function

$$\exp(\beta^\emptyset),$$

and thus, that the bounds that failed to give good results, had resulted in a too large value of this β^\emptyset .

Still focusing on Prospect P_{15} , we observe how the value of β^\emptyset changes as we sum out Random Variables in the expression for

$$\mathbb{E}(w_{15}(X_{B_{15}}, Z_{B_{15}}) - c(Z_{B_{15}})).$$

This test is done for $\nu = 10$, and the result is to be found in Table 12. We observe that the value of β^\emptyset is slowly growing at first, and that the blow-up starts at the 11th ancestor we sum out. Note that Prospect P_{15} has 13 ancestors, which means that we have to sum out a total of 14 nodes, including P_{15} itself. After summing out all of those Random Variables, the resulting upper bound is a function of the decision Z , with $\beta^\emptyset = 1968$. Since the logarithm of the exact value for no decision is even close to this number, this is an upper bound function where no single evaluation of the lower bound function can compete with even the minimum of the upper bound. Hence, no candidates for the optimal decision can be eliminated.

Number of Random Variables summed out	Value of β^\varnothing in current representation
0	-1.023
1	-1.006
2	-1.005
3	-1.005
4	-0.838
5	-0.821
6	-0.819
7	-0.811
8	0.715
9	1.450
10	1.702
11	8.014
12	17.48
13	169.7
14	196.8

Table 12: BN5: Value of β^\varnothing for the upper bound function determined in each step, summing out a Random Variable at a time. Here, $\nu = 10$ and $\epsilon = 0.01$.

ν	$\epsilon = 0.01$	$\epsilon = 0.05$	$\epsilon = 0.1$	$\epsilon = 0.2$
15	386.1	86.34	15.90	1.409
16	92.72	19.51	5.190	0.733
17	-0.000	-0.000	-0.000	-0.000

Table 13: BN5: Upper bounds on the log normalization constant for the probability distribution π_{15} calculated with different ν s and different ϵ s. 17 is the smallest value of ν that ensures no approximations to be made, and hence $\nu = 17$ gives the exact value $0 = \ln 1$.

Recall that we let $\epsilon > 0$ ensure that any assignment of the Random Variables has a positive probability. For the results in Table 11 and Table 12, $\epsilon = 0.01$ has been used. Taking the same Bayesian Network, with the same edge probabilities, we can vary ϵ and observe if that has any effect on this seemingly hopeless example. Again, we study only the probability distribution $\mathbb{P}(X_{A_{15}})$. Table 13 shows the resulting log normalization constants. Note that an increase in the value for ϵ introduces less error, but even high values of ϵ give useless results. Note that the results for $\epsilon = 0.2$ are not too bad, but 0.2 is definitely a questionable approximation to 0. Recall that $\epsilon > 0$, by definition, should be a small probability, hence much closer to the value 0 than the value 1.

10 Some Analysis on Complexity

First, we want to study the complexity of calculating $\mathbb{E}h(X, Z)$ with a naïve, and also exact, method, for some function h depending on the entries in X, Z . If the function h and the probability distribution $\mathbb{P}(X)$ is on the exponential pseudo-Boolean form, we can write

$$h(X, Z)\mathbb{P}(X) = \exp\left(\sum_{\lambda \in S} \beta^\lambda \prod_{k \in \lambda} \tilde{X}_k(X, Z)\right).$$

We could assume that S has size $2^{n+|\Gamma|}$. That corresponds to a full representation of $\mathbb{P}(X)$, that is, 2^n possible combinations of X_1, \dots, X_n , and also a full representation for $h(X, Z)$ for the variables X_1, \dots, X_n and Z_1, \dots, Z_n with indexes in Γ . That is, calculating the value of $h(X, Z)\mathbb{P}(X)$ for an assignment to X, Z has complexity $\mathcal{O}(2^{n+|\Gamma|})$. To get the expected value $\mathbb{E}h(X, Z)$ for a given Z , we would have to sum the value of $h(X, Z)\mathbb{P}(X)$ for 2^n different X s. That

implies a complexity of $\mathcal{O}(2^{2n+|\Gamma|})$ for finding the expected value for a given Z . Our decision space $\Omega^{\Gamma,u}$ is of size

$$|\Omega^{\Gamma,u}| = \sum_{i=1}^u \binom{|\Gamma|}{i} \in \mathcal{O}(|\Gamma|^u),$$

since we assume $u \ll |\Gamma|$. To evaluate the expected value $\mathbb{E}h(X, Z)$ for each Z in our decision space would then have a total complexity of

$$\mathcal{O}(|\Gamma|^u \cdot 2^{2n+|\Gamma|}). \quad (10.1)$$

However, note that this is based on a very naïve approach. In fact, we assume that the expected value is calculated as

$$\sum_{X_n} \left(\sum_{X_{n-1}} \cdots \left(\sum_{X_1} \exp \left(\sum_{\lambda \in S} \beta^\lambda \prod_{k \in \lambda} \tilde{X}_k(X, Z) \right) \right) \right),$$

where the expression inside each parenthesis pair is calculated without checking whether any factors can be moved outside the sum.

Let us again assume that we want to do exact calculations for the expected value $\mathbb{E}h(X, Z)$, now for a function h and a probability distribution $\mathbb{P}(X)$ not in the exponential pseudo-Boolean form. It is natural to assume that we find the value of $\mathbb{P}(X)$ with $\mathcal{O}(n)$ calculations, and that the complexity of calculating $h(X, Z)$ is within this, and usually even simpler. The resulting total complexity would then be

$$\mathcal{O}(n \cdot 2^n \cdot |\Gamma|^u). \quad (10.2)$$

Now, we have two cases to compare to, and we want to study the complexity for our method, using bounds. Again, first, assume that we start with the function h and the probability distribution $\mathbb{P}(X)$ on the exponential pseudo-Boolean form. For the input constant ν , recall that calculating $\mathbb{E}h(X, Z)$ as a function of Z has complexity $\mathcal{O}(n \cdot 2^\nu)$, plus the time spent on calculating the Approximate Markov Random Fields, which we assume is within the given complexity. This function $\mathbb{E}h(X, Z)$ is of the form

$$\mathbb{E}h(X, Z) = \exp \left(\sum_{\lambda \in \tilde{S}} \tilde{\beta}^\lambda \prod_{k \in \lambda} Z_k \right),$$

where for each $\lambda \in \tilde{S}$, we have $\lambda \subseteq \Gamma$. Thus, evaluating the function $\mathbb{E}h(X, Z)$ for a given Z requires a sum over $2^{|\Gamma|}$ terms. Repeating this for each Z in our

decision space, yields a total complexity of

$$\mathcal{O}(n \cdot 2^\nu + 2^{|\Gamma|} \cdot |\Gamma|^u). \quad (10.3)$$

Comparing (10.1) and (10.3), we observe a significant reduction of complexity. However, note that to get this reduction, we both assumed that we start with the exponential pseudo-Boolean forms and that the approximations "The AMRF Program" makes are made in $\mathcal{O}(n \cdot 2^\nu)$ time.

Looking at the case as in Section 8, we start with a function $h(X, Z)$ of the form

$$h(X, Z) = \sum_{\lambda \in T} \alpha^\lambda \prod_{k \in \lambda} X_k Z_k,$$

and then we need to calculate the form

$$h(X, Z) = \exp \left(\sum_{\lambda \in S} \beta_h^\lambda \prod_{k \in \lambda} X_k Z_k \right).$$

We can assume S to be the power set of Γ , and hence that we need to calculate $2^{|\Gamma|}$ coefficients β_h^λ . Using formulas from Section 3.2 and the coefficient graph structure and formulas from Section A, we get that calculating each β_h^Λ is of complexity $\mathcal{O}(|\lambda|^2)$. That is, we have assumed the β_h^Λ s are calculated in order according to increasing size of Λ , which ensures that all β_h^λ s with $\lambda \subsetneq \Lambda$ are available when we are calculating β_h^Λ . Then, calculating the wanted representation for the function h has complexity

$$\mathcal{O} \left(\sum_{\lambda \subseteq \Gamma} |\lambda|^2 \right) = \mathcal{O} \left(\sum_{i=1}^{|\Gamma|} i^2 \binom{|\Gamma|}{i} \right) \subseteq \mathcal{O} \left(|\Gamma|^2 \cdot 2^{|\Gamma|} \right).$$

We also need to calculate the exponential pseudo-Boolean representation of the probability distribution

$$\mathbb{P}(X_1, \dots, X_n) = \prod_{i=1}^n \mathbb{P}(X_i | \text{Pa}(X_i)).$$

Assuming that

$$r = \max_{1 \leq i \leq n} |\text{Pa}(X_i)|,$$

we get the pseudo-Boolean representation of $\mathbb{P}(X_i | \text{Pa}(X_i))$ in $\mathcal{O}(2^{r+1})$ calculations. Assuming r is not growing with n , which could be a reasonable assumption

if we assume a certain "sparseness" to be constant as the size of the Bayesian Network grows, this is a constant running time. Thus, calculating the representation of the full Joint Probability Distribution has a linear complexity in the total number n of Random Variables. This is because we calculate something of constant complexity for each Random Variable X_i . That would leave a total complexity of

$$\mathcal{O}(|\Gamma|^2 \cdot 2^{|\Gamma|} + n + n \cdot 2^\nu + 2^{|\Gamma|} \cdot |\Gamma|^u) \subseteq \mathcal{O}(n \cdot 2^\nu + 2^{|\Gamma|} \cdot |\Gamma|^u), \quad (10.4)$$

assuming $u \geq 2$. Comparing (10.2) and (10.4), we still see a reduction of complexity. Note that the calculation of each of the pseudo-Boolean representations disappeared in the complexity of "The AMRF Programs" and the evaluation for all Z s, respectively. Also note that for the tests in Section 9, we would calculate the representation of $w_j - c_j$ for each P_j with a complexity of $\mathcal{O}(t_j^2 2^{t_j})$, where t_j is twice the number of P_j 's Segment children. If we assume that the biggest such t_j does not grow as we increase the size n of our Bayesian Network, this would again mean that each $w_j - c_j$ representation is calculated in constant time. Thus, in total, all parts of h is calculated with with a complexity linear in the number of Prospects, which could be assumed to grow linearly with the total number n of Random Variables. In other words, this does not reduce the complexity of the full calculations, but it would reduce some of the constant factors.

11 Closing Remarks

We have studied five Boolean Bayesian networks; BN1-BN5, with Boolean decision vectors. First of all, from Section 8, we realized that our initial idea for how to do the calculations did not lead to good results. After, we could easily say that these calculations were done in a naïve way. But after some improvements, we observed that our solution method worked well on sparse graphs, as in BN2, BN3 and BN4. We also observed that usually, for a given ν , the loss fraction L_ν for an approximate optimal decision \tilde{Z} had a value equal to or close to zero. That means that on average, the approximate optimal decision \tilde{Z} worked well. Note that the value of ν was chosen to be fairly close to the expected size of the maximal clique in the graph. However, in Table 77 in Section C.4, we find several loss fractions L_ν which clearly implies choosing a decision with a low exact value. This is why we get a high value for the standard deviation, especially for BN4. On average, the performance is still good, because the algorithm finds the optimal decision Z^* , or makes the right guess, in most cases, also with

approximate values. To get less cases with high loss fractions, the algorithm could be modified to choose the exact best decision from the top few, according to the lower bounds, for example the top 5 or top 10. This would still allow us to do approximate calculations on most decisions.

We also observed how the solution method failed when the sparseness of the graph decreased, that is, when more edges were added. A further study of when this starts to happen, and how we could avoid this, would be interesting. Then, it would be natural to take a closer look on how the "optimal bounds" from Tjelmeland and Austad (2011) are calculated, and look for improvements or one's own version of such bounds. One could also look for improvement on the order of which the Random Variables are summed out, when calculating some expected value. Also here, a broader understanding of how the error bounds are calculated would be helpful.

This solution method could also be broadened by looking at upper and lower bounds on the value for a decision $Z \in \Omega^{\Gamma,u}$ where we only specify some of the entries. That is, for example, if we could find that any decision

$$Z^1 \in \{Z \in \Omega^{\Gamma,u} | Z_i = 1\}$$

would do worse than the best decision $Z^2 \in \Omega^{\Gamma,u}$ with $Z_i^2 = 0$, we would know that the optimal decision Z^* does not have $Z_i = 1$. This could be done by some approximate Viterbi algorithm.

As mentioned in the beginning of this section, we have assumed Boolean Random Variables and Boolean decision vectors. However, there are no reason why this solution method could not be expanded. We could assume Random Variables that take on values in a set A with $|A| > 2$. Then, our Random Vector X would be living in $\Omega = A^n$. Correspondingly, there could be more than one choice for what to do with each Random Variable, that is, more choices for the values of the entries of the decision vector. The challenge is then in how to define functions and probability distributions so that we still can do approximations on these. In this report, we have assumed the approximations from Tjelmeland and Austad (2011), which are defined on pseudo-Boolean functions. Of course, each Random Variable $X_i \in A$ could be represented by a collection of Boolean Variables, and correspondingly for the decision entries Z_i . But probably, it would be better to use other upper and lower bounds for that specific case. In either case, the method for eliminating decisions as described in Section 6 is still valid.

Also, we recall from Section 10, that, of course, this solution method with "The AMRF Program" bounds, has the greatest reduction in complexity if

we start with functions that already are on exponential pseudo-Boolean form. Thus, according to the complexity of the algorithm, this solution method would be most useful for cases where it is natural to start with such a formulation of the Joint Probability Distribution. However, note that for the dimensions in the tests for this report, most computational time was spent on calculating the expected value of the function $w - c$, and compared to that, the calculation of the exponential pseudo-Boolean forms was done in no time. Anyway, it could be interesting to look for applications, and study the performance on problems where the exponential pseudo-Boolean form is more natural. It is worth noting that in this report, we have only done tests for Bayesian Networks. Other graphical models could be of great interest. Note that for a Markov Random Field with Random Variables in X , and an unknown normalization constant C for the probability distribution $\frac{1}{C}P(X)$, an upper bound on the expected value of a function $h(X)$ could be expressed as

$$\frac{\sum^+_X h(X)P(X)}{\sum^-_X P(X)},$$

where we let \sum^+_X denote the upper bounds on the sum over X , and correspondingly, \sum^-_X for the lower bound. Note that the Ising model is an example of such a distribution, and so is also any other Markov Random Field where the Joint Probability Distribution

$$\mathbb{P}(X) = \frac{1}{C} \prod_{E(X)} \exp(E(X)),$$

is defined by a collection of energy functions $\{E(X)\}$. Also, it could be interesting to look for applications where it is more natural to have the value function as the exponential of a pseudo-Boolean function. That is, where the value function behaves more like a product.

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APPENDIX

A Implementation of Graph Structure

A graph like the one described in Section 3.1 is implemented by a C++ class. An alternative way of storing the coefficients α^λ would be in a vector of length 2^n . That is, each subset $\lambda \subseteq N$ corresponds to entry number $E(\lambda) = \sum_{k \in \lambda} 2^k$ in the vector. Note that if we only want to look at sets λ of size $|\lambda| \leq L$ with $L \ll m$, this alternative would imply a lot of unused entries in the vector. For the graph alternative, we are able to only create the first $L + 1$ layers of the graph, including the root node. However, the main argument for the graph alternative, is the efficient way of calculating sums over subsets, as in

$$\sum_{\lambda \subseteq \Lambda} \alpha^\lambda. \tag{A.1}$$

The number of layers in the graph, $L + 1$, together with the number of different values appearing in the sets, give a unique description of the graph structure. Also, each node is an instance of a struct holding a vector of pointers to its parents, a vector of pointers to its children and a vector holding its corresponding values. Associate each node with its index set λ , that is, in this discussion, we will not distinguish between the index set and its corresponding node. Let the root node be layer 0, and name each layer outwards layer 1, \dots , L .

We introduce an ordering on the children and parents of node a , where child b precedes child c if the element added to a to obtain b has a lower value than the element added to obtain c . Correspondingly for the parents, parent c precedes parent d if the element removed from a to obtain c has a lower value than the element removed to obtain d . That is, the listings of children and parents in Section 3.1 follows the ordering described above. In the graph, we let each node λ hold pointers to its children and parents, respectively, according to this order.

The length of the value vector may depend on the problem, and what we want to calculate for each node. Since we often want to calculate expressions of the kind in (A.1), this length is a multiple of the layer number r . If our purpose is to convert between the linear and exponential pseudo-Boolean representations of a function, as in Section 3.2, we would let the value vector length be twice the layer number, and let the root node hold two values. If we just want to store coefficients, or calculate sums over subsets, it is sufficient with a value vector length equal to the layer number, and the root node would hold one value. That is, call the value vector v , and assume for simplicity that each node has only

one coefficient associated with it, i.e. we work with one set of coefficients α^λ . Then, the value vector has length r in layer r , and the first value element in the node Λ holds the value of α^Λ . The k th element, $k \geq 2$, is the sum of the coefficients α^λ where $\lambda \subset \Lambda$ and $|\Lambda \setminus \lambda| = k - 1$. That is, the sum over the first value element for each node in the layer k steps down, reachable from Λ by k steps of going from a node to one of its parents. Then, if we let $\lambda \rightarrow v[k]$ denote the k th entry in the value vector for the node λ , we obtain the following two easy formulas for our graph structure,

$$\sum_{\lambda \subsetneq \Lambda} \alpha^\lambda = \emptyset \rightarrow v[0] + \sum_{k=2}^{|\lambda|-1} \lambda \rightarrow v[k], \quad (\text{A.2})$$

and

$$\sum_{\lambda \subseteq \Lambda} \alpha^\lambda = \emptyset \rightarrow v[0] + \sum_{k=1}^{|\lambda|-1} \lambda \rightarrow v[k]. \quad (\text{A.3})$$

Assume that we are given two nodes λ_1 and λ_2 , where λ_1 is an ancestor of λ_2 , k layers up. There are $k!$ different routes from y_1 to y_2 , going from a node to one of its children in each of the k steps. That is, in the first step, you have k different coordinates you can turn on, each of which in the next step gives $k - 1$ choices, and so on.

That results in the following formula, for $1 < k < |\lambda| - 1$,

$$\lambda \rightarrow v[k] = \frac{1}{k} \sum_{\tilde{\lambda} \in \text{Pa}(\lambda)} \tilde{\lambda} \rightarrow v[k - 1]. \quad (\text{A.4})$$

The resulting algorithm first creates the graph structure of the dimensions wanted. Then, the first entry $\lambda \rightarrow v[1]$ in the value vector for each node λ is assigned the value α^λ . After that, layer by layer, starting with the grand children of the root node and continuing outwards, the rest of the values in each value vector is filled out according to (A.4). In the end, the wanted sums as in (A.1), are easily calculated by the formulas in (A.2) and (A.3).

B A short User Oriented Introduction to "The AMRF Programs"

The programs referred to as "The AMRF Programs", are already implemented algorithms from Tjelmeland and Austad (2011), as mentioned in Section 4. For

more information and the theory behind it, the reader is referred to Tjelmeland and Austad (2011).

B.1 A pseudo-Boolean function as input

In addition to input parameters specific for each algorithm, a pseudo-Boolean function of the form

$$f(X) = \exp \left(\sum_{\lambda \in S} \beta^\lambda \prod_{k \in \lambda} X_k \right)$$

is given in an input file. In the model assumed, the nodes are numbered from N_0 to N_{end} , and each node k , with $N_0 \leq k \leq N_{end}$, has a corresponding coefficient $\beta^{\{k\}}$ in the coefficient set $\{\beta^\lambda\}_{\lambda \in S}$. The input file needs to be of the following format. The first line holds the start node number N_0 , and the second line holds the end node number N_{end} . Then, on each line, a set coefficient β^λ is presented, with no restriction on the order of the lines. But within each line, the following criteria need to be met.

- The first number on each line, is the size $|\lambda|$ of the coefficient set λ .
- The next numbers are the indexes $k \in \lambda$, in increasing order.
- Last, is the value of the corresponding coefficient β^λ .

As an example, the input file

```
1
3
0 -2.5
1 1 -2.8
1 2 -4.0
2 1 2 1.8
```

represents the function

$$f([X_1, X_2]) = \exp(-2.5 - 2.8X_1 - 4.0X_2 + 1.8X_1X_2).$$

B.2 Calculating an approximate normalizing constant

The approximate normalizing constant, or an approximation of a marginal, for a pseudo-Boolean function as in Section B.1 will be found by use of the function call

```
amrf_graph_v3  
(graphfile, simulation, likelihood, sim, limit, sstop, thev),
```

where the input parameters are

- **graphfile** is the name of the file where the pseudo-Boolean function is stored, as described in Section B.1.
- **simulation** is a file name for a file describing a realization for which the likelihood is to be evaluated (we set `sim=1` to get one).
- **likelihood** is the name of the file where the output will be stored.
- We set **sim** = 1, since the program requires a realization to evaluate the likelihood of.
- We set **limit** = 0, since we don't want to do approximations corresponding to their ϵ parameter.
- **sstop** is the number of the last node to be summed out, i.e. we obtain the marginal for the nodes $x_{sstop+1}$ to $x_{N_{end}}$. Default is **sstop** = 0, which corresponds to all nodes being summed out, i.e. finding the normalizing constant of the distribution.
- **thev** is the maximum number of neighbors for each node for not making approximations before its summed out. That is, a small value of **thev** usually means more approximations, while a high enough value means no approximations.

B.3 Calculating bounds for the normalizing constant

The upper and lower bounds for the normalizing constant, or bounds of a marginal, for a pseudo-Boolean function as in Section B.1, will be found by use of the function call

```
amrf_graph_bound(graphfile,  
simulation, likelihood, sim, limit, sstop, thev, maximum),
```

where the first seven input parameters are as for **amrf_graph.v3**, and the last parameter is set to be 1 for the upper bound, and 0 for the lower bound.

B.4 An approximate Viterbi algorithm

The upper or lower bounds for the maximum function value, or a function that bounds the maximum over a subset of the variables, for a pseudo-Boolean function as in Section B.1, will be found by use of the function call

```
amrf_graph_viterbi(graphfile,  
simulation, likelihood, sim, limit, sstop, thev, maximum),
```

where the input parameters are as for **amrf_graph_bound**.

C Probability Distribution, Cost and Income Functions for Section 8 and Section 9

C.1 Distribution, Cost and Income functions BN1

First, Table 14 provides all edge probabilities $p_{e_{i,j}}$ to fully determine the Joint Probability Distribution for the graph in Figure 7 in Section 8. Recall that we have assumed the distribution to be as in Section 3.3.1. Then, Tables 15 and 16 provides the linear coefficients to fully determine the two sets of cost and income functions, a and b , respectively. Recall that $\alpha_{cost}^{\vartheta(i)}$ represents the cost of drilling Segment S_i , while $-\alpha_{cost}^{\vartheta(i),\vartheta(j)}$ represents the savings in the costs when drilling both Segment siblings S_i, S_j , and correspondingly for the income coefficients.

Kitchen nr K_i	$\mathbb{P}(K_i = 1)$
K1	0.926
K2	0.728
K3	0.817
K4	0.735
Node pair (i, j)	$p_{e_{i,j}}$
(K1, P1)	0.217
(P1, D1)	0.840
(P1, D2)	0.637
(K2, P2)	0.773
(K3, P2)	0.916
(P2, D3)	0.805
(P2, D4)	0.515
(K2, P3)	0.608
(K3, P3)	0.788
(P3, D5)	0.909
(K4, P4)	0.471
(P2, P4)	0.575
(P4, D6)	0.818
(P4, D7)	0.762
(P4, P5)	0.616
(P5, D8)	0.613
(P2, P6)	0.717
(P4, P6)	0.797
(P6, D9)	0.583
(P4, P7)	0.609
(P6, P7)	0.409
(P7, D10)	0.524

Table 14: BN1: Probabilities for the Bayes Net in Figure 7.

Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D1	4552	5660
D2	2222	4693
D3	5382	3996
D4	1769	6829
D5	501.9	3534
D6	44.11	7291
D7	3170	3868
D8	365.9	2871
D9	917.7	3342
D10	1997	5326
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D1, D2)	871.3	236.8
(D3, D4)	405.8	430.5
(D6, D7)	4.140	83.43

Table 15: BN1: Cost function a and income function a .

Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D1	4729	4696
D2	4962	3830
D3	2419	2871
D4	5094	3342
D5	5758	5211
D6	2189	6161
D7	1928	5946
D8	3611	4478
D9	5277	7135
D10	1673	4884
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D1, D2)	3482	83.43
(D3, D4)	2180	139.3
(D6, D7)	649.8	288.9

Table 16: BN1: Cost function b and income function b .

C.2 Distribution, Cost and Income functions BN2

In this subsection, in addition to Sections C.3, C.4 and C.5, we will present the edge probabilities p_e , and also the cost and income coefficients in groups after which Prospect P_i they belong to. That is, we say that p_e belong to Prospect P_i if Prospect P_i is the end point of edge e , or if e is an edge from Prospect P_i to one of its Segment children. The cost and income coefficients under Prospect P_i , belongs to Prospect P_i 's Segment children. Then, the numbers appear in groups as we use them to calculate the representation of $w_j - c_j$ and $\mathbb{P}(X_{A_j})$, respectively. Note that the calculation of the latter also uses calculations corresponding to previous Prospects, however. All numbers listed in this subsection, corresponds to BN2 from Section 9.1.1, visualized in Figure 8. Sections C.2, C.3 and C.4 also provides a table over the nonzero loss fractions L_ν in the 1000 replications test for each, respectively.

Kitchen K_i	$\mathbb{P}(K_i = 1)$
K1	0.730
K2	0.778
K3	0.790
K4	0.836
K5	0.887
K6	0.732
K7	0.454

Table 17: Probabilities corresponding to each of Kitchens K_i .

Node pair (i, j)	$p_{e_{i,j}}$	
(K1, P1)	0.616	
(P1, D1)	0.358	
PDS D_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D1	3975	5068

Table 18: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_1 .

Node pair (i, j)	$pe_{i,j}$	
(K2, P2)	0.813	
(P2, D2)	0.838	
PDS D_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D2	4005	5037

Table 19: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_2 .

Node pair (i, j)	$pe_{i,j}$	
(K3, P3)	0.982	
(P3, D3)	0.700	
(P3, D4)	0.805	
(P3, D5)	0.496	
(P3, D6)	0.781	
PDS D_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D3	4036	5037
D4	4002	5034
D5	4012	5007
D6	3984	4951
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D3, D4)	345.6	522.8
(D3, D5)	153.9	490.1
(D3, D6)	603.1	493.7
(D4, D5)	1104	459.2
(D4, D6)	139.1	480.4
(D5, D6)	264.8	496.7

Table 20: BN2: Probabilities describing the probability distribution for Prospect P_3 .

Node pair (i, j)	$pe_{i,j}$	
(K4, P4)	0.937	
(P4, D7)	0.928	
(P4, D8)	0.548	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D7	3947	5031
D8	4014	5037
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D7, D8)	3614	501.3

Table 21: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_4 .

Node pair (i, j)	$pe_{i,j}$	
(K3, P5)	0.797	
(K5, P5)	0.550	
(P5, D9)	0.762	
(P5, D10)	0.956	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D9	4046	5001
D10	4024	4945
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D9, D10)	3897	505.0

Table 22: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_5 .

Node pair (i, j)	$pe_{i,j}$	
(K3, P6)	0.672	
(K6, P6)	0.613	
(P6, D11)	0.625	
(P6, D12)	0.789	
(P6, D13)	0.753	
(P6, D14)	0.807	
(P6, D15)	0.664	
(P6, D16)	0.841	
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Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D11	4095	4863
D12	3996	4985
D13	4005	4988
D14	4052	5110
D15	3944	4949
D16	4056	5003
<hr/>		
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D11, D12)	223.4	499.0
(D11, D13)	288.7	496.5
(D11, D14)	358.0	498.4
(D11, D15)	656.3	502.3
(D11, D16)	348.0	491.0
(D12, D13)	567.7	494.2
(D12, D14)	218.1	501.6
(D12, D15)	243.5	507.2
(D12, D16)	576.5	495.3
(D13, D14)	159.1	514.3
(D13, D15)	322.9	500.7
(D13, D16)	720.9	504.7
(D14, D15)	413.2	514.7
(D14, D16)	328.2	491.4
(D15, D16)	672.0	500.2

Table 23: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_6 .

Node pair (i, j)	$pe_{i,j}$	
(K7, P7)	0.803	
(P7, D17)	0.626	
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Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D17	3993	4982

Table 24: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_7 .

Node pair (i, j)	$pe_{i,j}$	
(K6, P8)	0.606	
(P8, D18)	0.934	
(P8, D19)	0.529	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D18	3986	5010
D19	4049	5001
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D18, D19)	3804	510.9

Table 25: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_8 .

Node pair (i, j)	$pe_{i,j}$	
(K1, P9)	0.265	
(P3, P9)	0.723	
(P9, D20)	0.354	
(P9, D21)	0.587	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D20	3996	5032
D21	4017	5084
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D20, D21)	2830	470.1

Table 26: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_9 .

Node pair (i, j)	$Pe_{i,j}$	
(K3, P10)	0.825	
(P4, P10)	0.809	
(P10, D22)	0.934	
(P10, D23)	0.759	
(P10, D24)	0.676	
(P10, D25)	0.718	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D22	3959	5017
D23	3989	4939
D24	4104	4976
D25	4008	4940
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D22, D23)	307.8	490.0
(D22, D24)	910.8	502.8
(D22, D25)	431.8	492.3
(D23, D24)	1096	465.9
(D23, D25)	1005	489.5
(D24, D25)	300.8	496.2

Table 27: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_{10} .

Node pair (i, j)	$Pe_{i,j}$	
(P6, P11)	0.592	
(P7, P11)	0.784	
(P8, P11)	0.844	
(P11, D26)	0.921	
(P11, D27)	0.479	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D26	4066	5022
D27	4065	5009
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D26, D27)	3881	492.0

Table 28: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_{11} .

Node pair (i, j)	$pe_{i,j}$	
(K4, P12)	0.850	
(P12, D28)	0.837	
(P12, D29)	0.855	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D28	4024	5034
D29	4034	5043
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D28, D29)	1370	494.9

Table 29: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_{12} .

Node pair (i, j)	$pe_{i,j}$	
(P3, P13)	0.267	
(P8, P13)	0.567	
(P10, D30)	0.724	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D30	4046	4988

Table 30: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_{13} .

Node pair (i, j)	$pe_{i,j}$	
(K5, P14)	0.892	
(P13, P14)	0.904	
(P14, D31)	0.769	
(P14, D32)	0.793	
(P14, D33)	0.587	
(P14, D34)	0.829	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
(D31, D32)	1054	487.1
(D31, D33)	223.3	502.5
(D31, D34)	184.9	478.8
(D32, D33)	579.3	502.7
(D32, D34)	817.4	517.1
(D33, D34)	420.3	524.8
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D9, D10)	3897	505.0

Table 31: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_{14} .

Node pair (i, j)	$pe_{i,j}$	
(K5, P15)	0.921	
(P4, P15)	0.683	
(P14, P15)	0.587	
(P15, D35)	0.849	
(P15, D36)	0.839	
(P15, D37)	0.658	
(P15, D38)	0.805	
(P15, D39)	0.822	
(P15, D40)	0.981	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D35	3952	4976
D36	3992	4958
D37	3909	5019
D38	3962	4965
D39	4070	4964
D40	4076	4912
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D35, D36)	299.7	511.1
(D35, D37)	744.7	502.8
(D35, D38)	100.5	488.2
(D35, D39)	17.69	499.0
(D35, D40)	548.2	504.2
(D36, D37)	435.0	504.6
(D36, D38)	359.2	489.3
(D36, D39)	358.3	514.5
(D36, D40)	599.7	510.3
(D37, D38)	15.29	511.4
(D37, D39)	129.1	490.8
(D37, D40)	18.45	496.0
(D38, D39)	71.00	502.9
(D38, D40)	534.2	504.3
(D39, D40)	626.4	495.2

Table 32: BN2: Probabilities, cost coefficients and income coefficients, Prospect P_{15} .

Nonzero loss fractions L_6			
0.0039%	2.0994%	6.6597%	17.086%
0.0171%	3.0456%	8.5423%	22.791%
0.0468%	3.0662%	8.7180%	27.024%
0.0596%	3.2426%	8.9417%	29.596%
0.2432%	3.4734%	9.2905%	31.045%
0.3263%	4.3125%	12.990%	37.278%
0.4715%	4.4725%	14.813%	37.991%
0.7584%	5.2099%	15.040%	41.739%
0.7924%	5.3182%	15.099%	61.640%
1.8310%	6.0684%	16.650%	83.181%

Table 33: BN2: Listing of all nonzero loss fractions L_6 for the 1000 tests in Section 9.1.1.

C.3 Distribution, Cost and Income functions BN3

All numbers listed in this subsection, corresponds to BN3 from Section 9.1.1, visualized in Figure 15.

Kitchen nr K_i	$\mathbb{P}(K_i = 1)$
K1	0.695
K2	0.944
K3	0.760
K4	0.638
K5	0.615
K6	0.668
K7	0.870
K8	0.593
K9	0.859
K10	0.821

Table 34: BN3: Probabilities corresponding to each of Kitchens K_i .

Node pair (i, j)	$p_{e_{i,j}}$	
(K1, P1)	0.510	
(P1, D1)	0.813	
(P1, D2)	0.934	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D1	3003	4949
D2	3012	5053
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D1, D2)	1273	478.3

Table 35: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_1 .

Node pair (i, j)	$pe_{i,j}$	
(K2, P2)	0.967	
(P2, D3)	0.878	
(P2, D4)	0.646	
(P2, D5)	0.780	
(P2, D6)	0.586	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D3	3004	4967
D4	2965	4987
D5	2933	5033
D6	2969	5050
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D3, D4)	84.85	518.9
(D3, D5)	950.4	483.4
(D4, D5)	478.8	499.2
(D3, D6)	219.1	532.8
(D4, D6)	805.0	464.1
(D5, D6)	566.4	531.5

Table 36: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_2 .

Node pair (i, j)	$pe_{i,j}$	
(K1, P3)	0.740	
(K3, P3)	0.585	
(P3, D7)	0.830	
(P3, D8)	0.968	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D7	3072	5060
D8	3000	4904
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D7, D8)	1663	488.6

Table 37: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_3 .

Node pair (i, j)	$pe_{i,j}$	
(K4, P4)	0.581	
(P4, D9)	0.589	
(P4, D10)	0.378	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D9	3042	5070
D10	2990	5021
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D9, D10)	2256	510.0

Table 38: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_4 .

Node pair (i, j)	$pe_{i,j}$	
(K5, P5)	0.782	
(P5, D11)	0.576	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D11	3050	4999

Table 39: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_5 .

Node pair (i, j)	$pe_{i,j}$	
(K6, P6)	0.532	
(P6, D12)	0.325	
(P6, D13)	0.692	
(P6, D14)	0.640	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D12	3011	5044
D13	2988	4958
D14	3086	4982
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D12, D13)	586.1	509.0
(D12, D14)	252.3	529.2
(D13, D14)	431.3	490.3

Table 40: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_6 .

Node pair (i, j)	$p_{e_{i,j}}$	
(K7, P7)	0.778	
(P7, D15)	0.687	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D15	2979	4939

Table 41: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_7 .

Node pair (i, j)	$p_{e_{i,j}}$	
(K8, P8)	0.563	
(P8, D16)	0.816	
(P8, D17)	0.790	
(P8, D18)	0.494	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D16	3078	4998
D17	3065	4921
D18	3054	4999
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D16, D17)	1286	467.0
(D16, D18)	901.9	518.6
(D17, D18)	348.5	507.8

Table 42: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_8 .

Node pair (i, j)	$p_{e_{i,j}}$	
(K9, P9)	0.549	
(P9, D19)	0.550	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D19	2932	5109

Table 43: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_9 .

Node pair (i, j)	$pe_{i,j}$	
(K10, P10)	0.676	
(P10, D20)	0.678	
(P10, D21)	0.782	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D20	2987	5052
D21	3010	5033
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D20, D21)	2526	479.1

Table 44: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{10} .

Node pair (i, j)	$pe_{i,j}$	
(P2, P11)	0.852	
(P9, P11)	0.860	
(P11, D22)	0.779	
(P11, D23)	0.842	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D22	2941	4916
D23	2989	5011
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D22, D23)	2823	512.3

Table 45: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{11} .

Node pair (i, j)	$p_{e_{i,j}}$	
(K5, P12)	0.502	
(P12, D24)	0.624	
(P12, D25)	0.823	
(P12, D26)	0.557	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D24	3005	5070
D25	2924	5016
D26	2946	5008
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D24, D25)	1247	511.6
(D24, D26)	641.8	490.1
(D25, D26)	956.3	497.4

Table 46: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{12} .

Node pair (i, j)	$p_{e_{i,j}}$	
(P10, P13)	0.976	
(P13, D27)	0.901	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D27	3072	4992

Table 47: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{13} .

Node pair (i, j)	$p_{e_{i,j}}$	
(K10, P14)	0.978	
(P4, P14)	0.817	
(P9, P14)	0.696	
(P14, D28)	0.659	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D28	2969	5022

Table 48: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{14} .

Node pair (i, j)	$pe_{i,j}$	
(K2, P15)	0.763	
(P7, P15)	0.890	
(P13, P15)	0.836	
(P15, D29)	0.850	
(P15, D30)	0.861	
(P15, D31)	0.776	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D29	2987	4951
D30	3004	5023
D31	3022	5020
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D29, D30)	267.8	497.1
(D29, D31)	894.1	482.6
(D30, D31)	1370	490.2

Table 49: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{15} .

Node pair (i, j)	$pe_{i,j}$	
(P6, P16)	0.719	
(P16, D32)	0.436	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D32	2966	4989

Table 50: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{16} .

Node pair (i, j)	$pe_{i,j}$	
(P9, P17)	0.898	
(P15, P17)	0.617	
(P17, D33)	0.478	
(P17, D34)	0.561	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D33	2913	4938
D34	2933	4980
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D33, D34)	1932	517.8

Table 51: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{17} .

Node pair (i, j)	$pe_{i,j}$	
(K1, P18)	0.246	
(P14, P18)	0.863	
(P18, D35)	0.634	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D35	2956	4952

Table 52: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{18} .

Node pair (i, j)	$pe_{i,j}$	
(P7, P19)	0.910	
(P19, D36)	0.760	
(P19, D37)	0.867	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D36	3050	5064
D37	2941	5023
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D36, D37)	1833	514.1

Table 53: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{19} .

Node pair (i, j)	$pe_{i,j}$	
(K3, P20)	0.601	
(P15, P20)	0.520	
(P19, P20)	0.498	
(P20, D38)	0.785	
(P20, D39)	0.887	
(P20, D40)	0.702	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D38	2920	5019
D39	2908	5048
D40	3003	4985
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D38, D39)	1395	514.6
(D38, D40)	179.8	523.7
(D39, D40)	289.0	480.6

Table 54: BN3: Probabilities, cost coefficients and income coefficients, Prospect P_{20} .

Nonzero loss fractions L_5		
0.0846%	0.1268%	0.1278%
0.1483%	0.1529%	0.6006%
1.1325%	1.7083%	1.7236%
1.8870%	2.5999%	2.6884%
3.2948%	3.3345%	3.4698%
3.8284%	4.1870%	4.5679%
5.1716%	5.4370%	6.0701%
6.9327%	10.1125%	20.0158%
20.5417%	25.1488%	45.8189%

Table 55: BN3: Listing of all nonzero loss fractions L_5 for the 1000 tests in Section 9.1.2.

C.4 Distribution, Cost and Income functions BN4

All numbers listed in this subsection, corresponds to BN4 from Section 9.1.1, visualized in Figure 23.

Kitchen nr K_i	$\mathbb{P}(K_i = 1)$
K1	0.719
K2	0.487
K3	0.897
K4	0.759
K5	0.817
K6	0.904
K7	0.351
K8	0.678
K9	0.874
K10	0.680

Table 56: BN4: Probabilities corresponding to each of Kitchens K_i .

Node pair (i, j)	$pe_{i,j}$	
(K1, P1)	0.700	
(K8, P1)	0.833	
(P1, D1)	0.778	
(P1, D2)	0.820	
(P1, D3)	0.815	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D1	4030	5015
D2	4046	4989
D3	4037	5016
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D1, D2)	1889	471.4
(D1, D3)	284.2	488.2
(D2, D3)	457.7	495.3

Table 57: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_1 .

Node pair (i, j)	$pe_{i,j}$	
(K2, P2)	0.678	
(K3, P2)	0.844	
(P2, D4)	0.707	
(P2, D5)	0.504	
(P2, D6)	0.516	
(P2, D7)	0.760	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D4	4042	5028
D5	3989	5013
D6	4001	4999
D7	4000	5029
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D4, D5)	158.8	493.2
(D4, D6)	235.6	490.1
(D5, D6)	1198	504.3
(D4, D7)	178.5	502.7
(D5, D7)	242.3	511.3
(D6, D7)	878.2	515.7

Table 58: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_2 .

Node pair (i, j)	$pe_{i,j}$	
(P3, D8)	0.714	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D8	4056	5050

Table 59: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_3 .

Node pair (i, j)	$Pe_{i,j}$	
(K4, P4)	0.478	
(P3, P4)	0.815	
(P4, D9)	0.621	
(P4, D10)	0.735	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D9	4037	5044
D10	3955	4975
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D9, D10)	727.7	491.7

Table 60: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_4 .

Node pair (i, j)	$Pe_{i,j}$	
(K5, P5)	0.803	
(K7, P5)	0.846	
(P5, D11)	0.676	
(P5, D12)	0.537	
(P5, D13)	0.379	
(P5, D14)	0.523	
(P5, D15)	0.799	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D11	3953	5025
D12	4054	4959
D13	4043	4994
D14	3988	5099
D15	3954	4987
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D11, D12)	92.44	524.1
(D11, D13)	606.0	479.3
(D12, D13)	64.68	530.0
(D11, D14)	361.3	511.1
(D12, D14)	764.5	470.1
(D13, D14)	948.3	505.4
(D11, D15)	929.5	505.9
(D12, D15)	227.3	524.8
(D13, D15)	575.1	502.6
(D14, D15)	475.3	491.8

Table 61: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_5 .

Node pair (i, j)	$p_{e_{i,j}}$	
(K6, P6)	0.937	
(P6, D16)	0.462	
(P6, D17)	0.698	
(P6, D18)	0.877	
(P6, D19)	0.722	
(P6, D20)	0.592	
(P6, D21)	0.395	
(P6, D22)	0.590	
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Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D16	4064	5007
D17	3944	4934
D18	3964	4964
D19	3972	5068
D20	3959	5016
D21	4025	4982
D22	3952	5013
<hr/>		
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i),\vartheta(j)}$	$\alpha_{inc}^{\vartheta(i),\vartheta(j)}$
(D16, D17)	414.1	517.2
(D16, D18)	239.6	509.5
(D17, D18)	288.5	483.8
(D16, D19)	295.4	502.3
(D17, D19)	313.5	497.2
(D18, D19)	364.3	501.3
(D16, D20)	631.8	471.0
(D17, D20)	65.94	499.8
(D18, D20)	528.9	484.6
(D19, D20)	302.6	512.9
(D16, D21)	372.8	480.1
(D17, D21)	400.5	507.2
(D18, D21)	360.7	482.2
(D19, D21)	647.6	501.6
(D20, D21)	184.2	503.5
(D16, D22)	519.2	495.1
(D17, D22)	78.64	494.0
(D18, D22)	345.7	500.9
(D19, D22)	549.0	480.5
(D20, D22)	48.52	502.9
(D21, D22)	495.5	494.7

Table 62: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_6 .

Node pair (i, j)	$pe_{i,j}$	
(K7, P7)	0.894	
(P7, D23)	0.569	
(P7, D24)	0.933	
(P7, D25)	0.885	
(P7, D26)	0.396	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D23	4008	5074
D24	3964	4969
D25	4029	5049
D26	4072	5025
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D23, D24)	155.5	485.7
(D23, D25)	286.2	510.0
(D24, D25)	1048	469.7
(D23, D26)	853.5	489.7
(D24, D26)	399.8	492.3
(D25, D26)	1079	491.1

Table 63: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_7 .

Node pair (i, j)	$pe_{i,j}$	
(K1, P8)	0.801	
(K8, P8)	0.862	
(P8, D27)	0.768	
(P8, D28)	0.943	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D27	4028	5016
D28	4028	4933
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D27, D28)	980.0	510.6

Table 64: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_8 .

Node pair (i, j)	$Pe_{i,j}$	
(K9, P9)	0.838	
(P9, D29)	0.757	
(P9, D30)	0.759	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D29	4042	4938
D30	4078	4945
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D29, D30)	1355	496.9

Table 65: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_9 .

Node pair (i, j)	$Pe_{i,j}$	
(K3, P10)	0.838	
(K6, P10)	0.674	
(K10, P10)	0.753	
(P10, D31)	0.677	
(P10, D32)	0.840	
(P10, D33)	0.776	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D31	4042	5005
D32	3934	4966
D33	3949	5013
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D31, D32)	332.0	499.2
(D31, D33)	931.6	480.5
(D32, D33)	1910	484.4

Table 66: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{10} .

Node pair (i, j)	$pe_{i,j}$	
(K7, P11)	0.813	
(P5, P11)	0.554	
(P8, P11)	0.766	
(P11, D34)	0.623	
(P11, D35)	0.475	
(P11, D36)	0.697	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D34	3930	5065
D35	3969	4944
D36	4058	5007
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D34, D35)	581.2	474.5
(D34, D36)	1263	504.4
(D35, D36)	531.3	484.6

Table 67: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{11} .

Node pair (i, j)	$pe_{i,j}$	
(P3, P12)	0.900	
(P12, D37)	0.858	
(P12, D38)	0.613	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D37	4004	5034
D38	3986	4980
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D37, D38)	3758	491.5

Table 68: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{12} .

Node pair (i, j)	$pe_{i,j}$	
(K4, P13)	0.706	
(K7, P13)	0.653	
(P9, P13)	0.816	
(P13, D39)	0.563	
(P13, D40)	0.895	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D39	3965	5049
D40	4043	4993
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D39, D40)	1186	476.6

Table 69: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{13} .

Node pair (i, j)	$pe_{i,j}$	
(K5, P14)	0.866	
(P14, D41)	0.822	
(P14, D42)	0.593	
(P14, D43)	0.508	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D41	4073	5059
D42	4015	4962
D43	3977	4963
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D41, D42)	292.3	511.7
(D41, D43)	1978	478.8
(D42, D43)	1792	486.1

Table 70: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{14} .

Node pair (i, j)	$p_{e_{i,j}}$	
(P11, P15)	0.605	
(P12, P15)	0.864	
(P15, D44)	0.853	
(P15, D45)	0.572	
(P15, D46)	0.532	
(P15, D47)	0.942	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D44	4059	5024
D45	4045	4921
D46	3992	4975
D47	3992	5049
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D44, D45)	72.11	485.5
(D44, D46)	696.3	475.1
(D45, D46)	449.1	522.0
(D44, D47)	950.7	489.1
(D45, D47)	777.7	458.4
(D46, D47)	1012	516.7

Table 71: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{15} .

Node pair (i, j)	$p_{e_{i,j}}$	
(K1, P16)	0.625	
(P16, D48)	0.590	
(P16, D49)	0.488	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D48	4002	5015
D49	3967	4955
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D48, D49)	2572	484.7

Table 72: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{16} .

Node pair (i, j)	$pe_{i,j}$	
(P4, P17)	0.633	
(P5, P17)	0.826	
(P17, D50)	0.822	
(P17, D51)	0.963	
(P17, D52)	0.555	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D50	4055	4999
D51	3964	4993
D52	4081	4947
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D50, D51)	739.5	501.3
(D50, D52)	701.6	502.1
(D51, D52)	1895	510.7

Table 73: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{17} .

Node pair (i, j)	$pe_{i,j}$	
(P3, P18)	0.841	
(P4, P18)	0.750	
(P15, P18)	0.771	
(P18, D53)	0.711	
(P18, D54)	0.971	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D53	4049	5046
D54	4062	5044
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D53, D54)	146.4	502.4

Table 74: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{18} .

Node pair (i, j)	$pe_{i,j}$	
(K1, P19)	0.695	
(P2, P19)	0.833	
(P5, P19)	0.618	
(P16, P19)	0.351	
(P19, D55)	0.386	
(P19, D56)	0.741	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D55	3944	5013
D56	3923	5003
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D55, D56)	2692	508.8

Table 75: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{19} .

Node pair (i, j)	$pe_{i,j}$	
(K3, P20)	0.669	
(P14, P20)	0.984	
(P19, P20)	0.813	
(P20, D57)	0.509	
(P20, D58)	0.633	
(P20, D59)	0.774	
(P20, D60)	0.789	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D57	4003	4981
D58	4042	4964
D59	4051	4945
D60	3945	4938
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D57, D58)	802.0	509.7
(D57, D59)	304.2	472.6
(D58, D59)	439.7	499.7
(D57, D60)	279.4	530.4
(D58, D60)	951.4	491.4
(D59, D60)	423.4	512.8

Table 76: BN4: Probabilities, cost coefficients and income coefficients, Prospect P_{20} .

Nonzero loss fractions L_7			
0.0008%	0.0143%	0.1081%	0.1408%
0.4298%	0.4729%	1.2513%	1.3379%
1.4173%	1.5933%	1.6141%	1.6901%
2.3847%	2.4245%	3.1822%	3.7893%
3.8178%	4.3210%	4.3220%	4.9154%
5.3173%	5.8367%	6.0373%	6.1426%
6.6618%	7.5549%	8.2433%	9.0723%
9.3761%	9.5243%	10.5092%	12.3766%
14.8005%	15.5166%	15.6522%	17.3257%
17.6869%	17.7826%	19.2693%	20.2625%
20.9264%	21.2441%	23.4648%	24.5481%
26.7271%	27.8775%	29.7215%	36.6901%
36.9098%	39.4034%	40.1745%	44.1414%
47.6808%	49.1176%	50.9428%	53.4218%
55.2268%	63.2664%	66.5337%	66.7269%
69.9822%	72.1796%	73.6152%	76.5762%
78.0995%	90.0164%	95.7869%	

Table 77: BN4: Listing of all nonzero loss fractions L_7 for the 1000 tests in Section 9.1.3.

C.5 Distribution, Cost and Income functions BN5

All numbers listed in this subsection, corresponds to BN5 from Section 9.2, visualized in Figure 30.

Kitchen K_i	$\mathbb{P}(K_i = 1)$
K1	0.730
K2	0.778
K3	0.790
K4	0.836
K5	0.887
K6	0.732
K7	0.454

Table 78: BN5: Probabilities corresponding to each of Kitchens K_i .

Node pair (i, j)	$p_{e_{i,j}}$	
(K1, P1)	0.616	
(K4, P1)	0.813	
(P1, D1)	0.769	
Segment S_i	$\vartheta_{cost}^{(i)}$	$\vartheta_{inc}^{(i)}$
D1	3863	4996

Table 79: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_1 .

Node pair (i, j)	$pe_{i,j}$	
(K2, P2)	0.982	
(P2, D2)	0.793	
(P2, D3)	0.587	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D2	3985	5005
D3	3988	5001
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D2, D3)	313.8	494.2

Table 80: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_2 .

Node pair (i, j)	$pe_{i,j}$	
(K1, P3)	0.937	
(K3, P3)	0.797	
(P3, D4)	0.829	
(P3, D5)	0.849	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D4	4052	5110
D5	3996	5058
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D4, D5)	3511	501.6

Table 81: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_3 .

Node pair (i, j)	$pe_{i,j}$	
(K4, P4)	0.550	
(P4, D6)	0.839	
(P4, D7)	0.658	
(P4, D8)	0.805	
(P4, D9)	0.822	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D6	3981	5052
D7	4046	4944
D8	4091	5048
D9	4002	5036
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D6, D7)	473.2	502.3
(D6, D8)	940.6	484.3
(D7, D8)	1312	517.3
(D6, D9)	1160	495.3
(D7, D9)	1201	504.7
(D8, D9)	540.3	491.4

Table 82: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_4 .

Node pair (i, j)	$pe_{i,j}$	
(K3, P5)	0.672	
(K5, P5)	0.613	
(K7, P5)	0.803	
(P5, D10)	0.981	
(P5, D11)	0.459	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D10	3990	5002
D11	3993	4982
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D10, D11)	516.4	493.2

Table 83: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_5 .

Node pair (i, j)	$pe_{i,j}$	
(K3, P6)	0.606	
(K6, P6)	0.265	
(P4, P6)	0.723	
(P6, D12)	0.677	
(P6, D13)	0.520	
(P6, D14)	0.863	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D12	4054	4928
D13	3956	5036
D14	4009	5034
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D12, D13)	1456	494.1
(D12, D14)	85.20	508.3
(D13, D14)	1244	486.7

Table 84: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_6 .

Node pair (i, j)	$pe_{i,j}$	
(K7, P7)	0.825	
(P7, D15)	0.589	
(P7, D16)	0.996	
(P7, D17)	0.495	
(P7, D18)	0.478	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D15	4017	4989
D16	3939	5077
D17	3961	5021
D18	3978	4948
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D15, D16)	225.9	502.8
(D15, D17)	678.7	465.9
(D16, D17)	117.9	484.0
(D15, D18)	1085	481.0
(D16, D18)	429.6	492.3
(D17, D18)	998.7	489.5

Table 85: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_7 .

Node pair (i, j)	$pe_{i,j}$	
(K4, P8)	0.809	
(K6, P8)	0.592	
(P4, P8)	0.784	
(P8, D19)	0.624	
(P8, D20)	0.603	
(P8, D21)	0.625	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D19	3936	4976
D20	4024	5034
D21	4030	4968
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D19, D20)	480.0	504.2
(D19, D21)	1540	514.3
(D20, D21)	1748	491.0

Table 86: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_8 .

Node pair (i, j)	$pe_{i,j}$	
(K1, P9)	0.844	
(K3, P9)	0.850	
(P1, P9)	0.267	
(P3, P9)	0.567	
(P5, P9)	0.892	
(P9, D22)	0.877	
(P9, D23)	0.635	
(P9, D24)	0.443	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D22	3951	5004
D23	3965	5009
D24	4119	4985
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D22, D23)	1700	506.7
(D22, D24)	963.1	502.5
(D23, D24)	872.3	502.7

Table 87: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_9 .

Node pair (i, j)	$pe_{i,j}$	
(K3, P10)	0.904	
(P3, P10)	0.921	
(P4, P10)	0.683	
(P10, D25)	0.501	
(P10, D26)	0.798	
(P10, D27)	0.707	
(P10, D28)	0.807	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D25	4007	4939
D26	4046	5046
D27	4028	5104
D28	3956	4921
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D25, D26)	599.3	487.0
(D25, D27)	1076	492.0
(D26, D27)	738.7	486.4
(D25, D28)	626.0	470.9
(D26, D28)	452.6	510.1
(D27, D28)	532.4	504.6

Table 88: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_{10} .

Node pair (i, j)	$pe_{i,j}$	
(P6, P11)	0.587	
(P7, P11)	0.358	
(P8, P11)	0.838	
(P11, D29)	0.860	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D29	3962	4965

Table 89: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_{11} .

Node pair (i, j)	$pe_{i,j}$	
(K2, P12)	0.700	
(K3, P12)	0.805	
(K4, P12)	0.496	
(P6, P12)	0.781	
(P7, P12)	0.928	
(P11, P12)	0.548	
(P12, D30)	0.591	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D30	3953	4964

Table 90: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_{12} .

Node pair (i, j)	$pe_{i,j}$	
(P2, P13)	0.762	
(P3, P13)	0.956	
(P6, P13)	0.625	
(P8, P13)	0.789	
(P10, P13)	0.753	
(P13, D31)	0.846	
(P13, D32)	0.795	
(P13, D33)	0.716	
(P13, D34)	0.808	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D31	4037	5070
D32	3964	4989
D33	4010	5076
D34	3885	4979
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D31, D32)	262.9	499.0
(D31, D33)	747.7	497.8
(D32, D33)	644.5	504.2
(D31, D34)	825.0	510.3
(D32, D34)	30.57	496.0
(D33, D34)	873.2	504.3

Table 91: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_{13} .

Node pair (i, j)	$pe_{i,j}$	
(K5, P14)	0.807	
(K6, P14)	0.664	
(K7, P14)	0.841	
(P1, P14)	0.626	
(P2, P14)	0.934	
(P8, P14)	0.529	
(P9, P14)	0.354	
(P10, P14)	0.587	
(P12, P14)	0.934	
(P13, P14)	0.759	
(P14, D35)	0.921	
(P14, D36)	0.750	
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Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D35	3986	4977
D36	4018	4985
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Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D35, D36)	2715	493.5

Table 92: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_{14} .

Node pair (i, j)	$p_{e_{i,j}}$	
(K5, P15)	0.676	
(P1, P15)	0.718	
(P4, P15)	0.921	
(P8, P15)	0.479	
(P12, P15)	0.837	
(P13, P15)	0.855	
(P14, P15)	0.724	
(P15, D37)	0.913	
(P15, D38)	0.650	
(P15, D39)	0.873	
(P15, D40)	0.611	
Segment S_i	$\alpha_{cost}^{\vartheta(i)}$	$\alpha_{inc}^{\vartheta(i)}$
D37	3983	5030
D38	3971	5035
D39	4000	5008
D40	3982	5028
Segment siblings S_i, S_j	$-\alpha_{cost}^{\vartheta(i), \vartheta(j)}$	$\alpha_{inc}^{\vartheta(i), \vartheta(j)}$
(D37, D38)	524.5	505.4
(D37, D39)	972.3	493.8
(D38, D39)	372.8	509.1
(D37, D40)	1251	486.5
(D38, D40)	1257	508.7
(D39, D40)	289.9	495.8

Table 93: BN5: Probabilities, cost coefficients and income coefficients, Prospect P_{15} .