

Bayesian Network Exploratory Tool

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April 17, 2017

1 Introduction

Probabilistic Graphical Models (PGM) are useful constructs for describing complex probability distributions over high dimensional space. Bayesian networks are a common type of PGMs described using a directed graph, in which nodes correspond to random variables and edges correspond to direct dependencies between variables. The power of PGMs comes from the fact that they factor a large complex joint probability distribution into a product of much smaller probability tables. The graph structure allows for an intuitive visual representation of the network of dependencies between variables. Additionally, formulation of a high-dimensional probability distribution using PGMs enables efficient algorithms for inference from data, and even learning the underlying probability.

The main objective of this project is to develop a user-friendly software tool for constructing Bayesian networks that allows interactive exploration of different aspects and inference algorithms. This tool will be designed both for educational use in academic courses and for self exploration by scientists and developers in the industry. The proposed software tool will contain six main functional units (see Section 2 for more detailed description):

1. Unit 1 - conditional independence in Bayesian networks (D separation criterion).
2. Unit 2 - undirected representations of Bayesian networks and the concept of triangular (or chordal) graphs.
3. Unit 3 - elimination orders. Examine the influence of elimination order on the complexity of the underlying inference algorithm.
4. Unit 4 - inference algorithms. Inferring conditional distributions given observed data using elimination and message passing.
5. Unit 5 - parameter inference.
6. Unit 6 - sampling algorithms on Bayesian networks (MCMC, loopy belief propagation).

There are several existing software tools for PGMs:

1. Bayesian DAG learning [1] - Bayesian inference about directed acyclic graph (DAG) structures using dynamic programming and MCMC (Markov Chain Monte Carlo). This tool is free and it's written in Matlab. The main objective is to facilitate computation, designed to run on small amount of variables (up to 20). It's main features are:

- (a) Computes exact edge marginals using Bayesian model averaging with dynamic programming
 - (b) Computes edge marginals approximately using MCMC
2. Probabilistic Graphical Models [4] - Inference of Bayesian networks and Markov chains. This tool includes learning examples as well. Implemented in Matlab with following features:
- (a) Joint distribution calculation
 - (b) Marginal distribution calculation
 - (c) Maximum probability assignment calculation
 - (d) Inference on tree structured network

Other relevant tools, e.g OpenGM [3] (Discrete factor graph models), Hidden Markov Model Toolbox [2] (Hidden Markov Model), MAP estimation of DAG structure [5], are more specialized and do not contain the main features of our proposed tool.

In summary, existing libraries have some aspects in common with our tool, but none of them appear to have the interactive nature that will support friendly exploration and learning. Additionally, none of these tools possess the gradual unit structure we propose, which we believe will help convey complex concepts in statistical learning.

2 Proposed Design

The software tool will consist of six sequential functional units. A Bayesian network is represented via a directed acyclic graph, where each node represents a random variable, with conditional distribution table given its parents (or predecessors) in the graph. The visual representation of the graph allows users to explore the network and observe the different conditional distribution tables.

1. Unit 1 – fundamental concept of conditional independence in Bayesian networks. This unit considers only the graph structure of the network and ignores the probability tables.
 - (a) Marginal independence. The user selects two nodes in the network, and the program specifies whether they are independent or not. If they are dependent, then the connecting path is highlighted.
 - (b) Conditional independence. The user selects two nodes u, v and a set of nodes to condition on A , and the program specifies whether X_u and X_v are conditionally independent given the collection X_A ($X_V \perp\!\!\!\perp X_U | X_A$). We will use the D-separation criterion for this and highlight the path that blocks dependence.
2. Unit 2 – undirected graph representation of Bayesian networks. This unit also considers only the graph structure of the network and ignores the probability tables.

- (a) Convert the directed graph representation to an undirected graph, called the moralized graph. In this undirected representation parents of a node in the original graph will have a new edge between them. Consequently, all variables in the same conditional probability table form a clique.
 - (b) Examine whether the moralized graph is chordal (triangulated) or not. In a chordal graph, every cycle of four or more vertices has a chord. Note that a perfect elimination exists, iff the moralized graph is chordal (see Unit 3).
 - (c) Convert the moralized graph into a factor graph – a bipartite graph representation with one class representing nodes in original graph and the other class representing maximal cliques (probability distribution tables).
3. Unit 3 – Elimination algorithm on undirected graph. Helps choose an elimination order for the purpose of inference without carrying out the complex computations of the probability tables.
- (a) Show elimination algorithm implementation on graphs. User can run elimination algorithm step by step. On every step the user choose a node to eliminate, on the graph the user can see the new edges added to the graph.
 - (b) Try to find a prefect elimination order, choose one simplicial node and start elimination from there, write the order on the screen, on every step find the next simplicial node, if there is not such one, the user will be able to choose the next node to eliminate.
4. Unit 4 – Inference of hidden variables using observed variables. In this unit the main objective is to show calculation of joint, marginal and max probability of random variables. Show inference on graph representation (undirected, factor) of Bayesian network along with probability tables. Inference starts by the user selecting as first step the user selects the observed nodes on the network, for each one sets the node value (for elimination algorithm choose the nodes to inference).

Inference then proceeds using one of three possible procedures:

- (a) Simple elimination - find marginal distribution for a set of variables given another set of observed variables.
- (b) Message Passing - compute marginal distribution for several variables separately given a set of observed variables.
- (c) Parallel Message Passing - same computation as in the last item but run the procedure in parallel (cannot run the step by step execution mode)

The message passing algorithm (sequential or parallel) can be used to make one of two possible calculations:

- (a) Marginal distribution - calculates marginal distribution for every node
- (b) Max probability assignment - computes assignment of unobserved nodes with observed nodes

All inference procedures can be run automatically (user just see the end result), or step-by-step to allow users to follow the calculations.

5. Unit 5 – Parameter inference from data. The networks this unit is operating on are HMM (Hidden Markov Model) and small Bayesian network with many independent copies.
The user has the ability to choose a couple of modes:
 - (a) Inference where all nodes are observed. is data to sufficient statistics table and display it to the user.
 - (b) Inference where some of the nodes are observed and some are hidden using:
 - i. Maximum probability inference
 - ii. EM (expectation maximization)
6. Unit 6 – Other advanced algorithms such as:
 - (a) Markov chain Monte Carlo algorithm
 - (b) Loopy Belief Propagation

3 Implementation Demo

For a demonstration of the tool, we constructed a Bayesian network with 5 binary random variables, and added dependencies between them (see figure 4). Figure 4 illustrates that conditional distribution table of variable (X_4) is displayed when its vertex in the network is selected via mouse click.

The Binary membership table have two dimensions, one shows all the probability functions one random variable participant in, the second shows all random variables of one probability function. Figure 5 shows the binary membership table for this network. Another important feature of the tool is the ability to examine the undirected version of the graph, moralized graph (Figure 6; unit 2). Note that in our network the moralized graph is not chordal, there is a simple cycle of 4 without a chord (X_1, X_2, X_3, X_4). Next we can start an elimination process by selecting one vertex in the graph (X_4), we can see how it is eliminated from the graph and new edges introduced in the undirected graph representation of the network (figure 7; unit 3). Binary membership table also changed, all probability functions this variable participant in replaced by the new probability function introduced in the elimination.

4 Work plan

Next step is implementation of the tool, right now there is a proof of concept of the tool and all screen shots on this proposal taken from there. Implementation of the tool will be sequential by the units (unit 1, unit 2 ...), since every unit adds new capabilities to former one. But before all units, we are going to implement the UI to define a Bayesian network.

Theoretical and user guide documentation for every unit are written along with the unit implementation.

This tool have two part, UI and server (support heavy operations on the network). As part of the tool we will use different open source libraries.

1. UI - Using browser for display, so languages are Javascript, CSS, HTML. The open source project we are using to support our tool are: React, Redux, Vis.js (Graph library), DataTables (HTML table library).

2. Server - Written in Java. Using Spring boot as framework for DI (dependency injection) and to create executable JAR.

This tool will have two predefined networks, the user can use them instead of defining a new network. The structure of the networks are:

1. Small Bayesian network with 6 random variables. This network is not chordal in it's moralized representation.

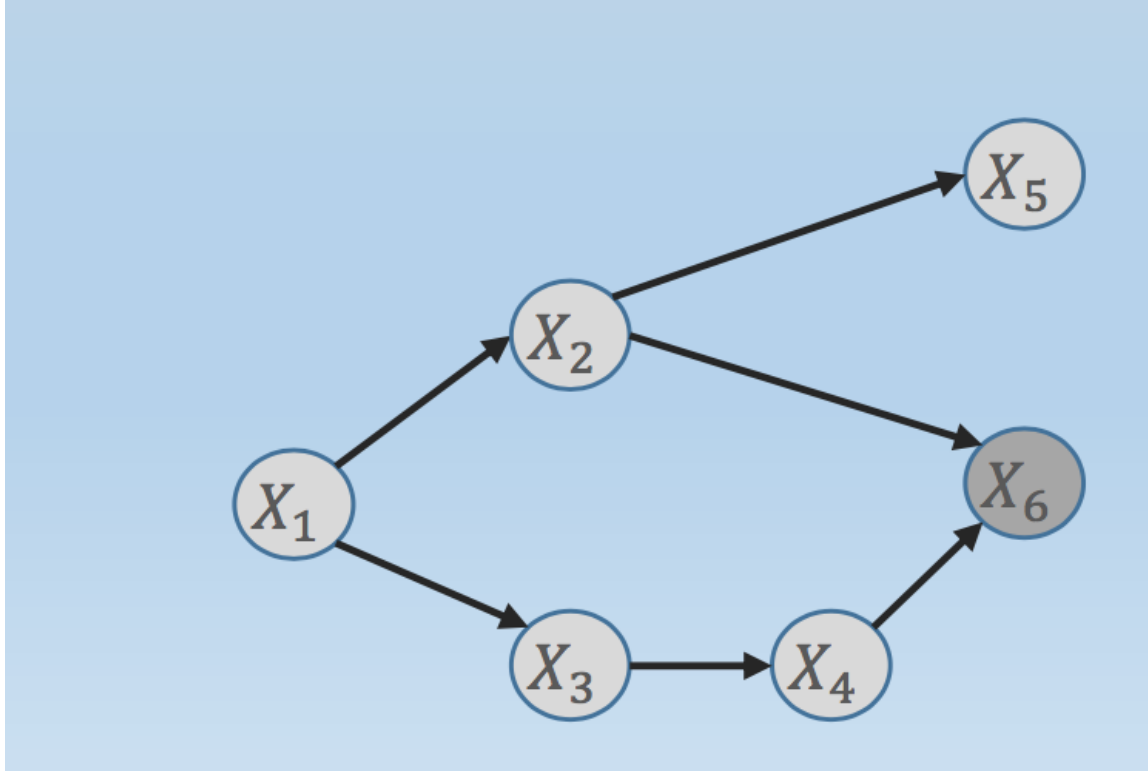


Figure 1: Six random variable Bayesian network representation using directed graph

2. Network with length parameter N , with three variables X, Y, Z , where X and Y dependent on N . Dependencies in the network: $\forall y_i \in Y, i > 0 P(y_i|y_{i-1}), \forall x_i \in X P(x_i|Y_i, Z)$ figure 2. In figure 3 we can see the conditional table.

References

- [1] Kevin Murphy, PGM Matlab libraries,
<http://www.cs.ubc.ca/~murphyk/Software/>

- [2] Kevin Murphy, toolbox for inference on Hidden Markov Models, <http://www.cs.ubc.ca/~murphyk/Software/HMM/hmm.html>
- [3] Bjoern Andre, Thorsten Beier and Joerg H. Kappes, OpenGM, <http://hciweb2.iwr.uni-heidelberg.de/opengm/>

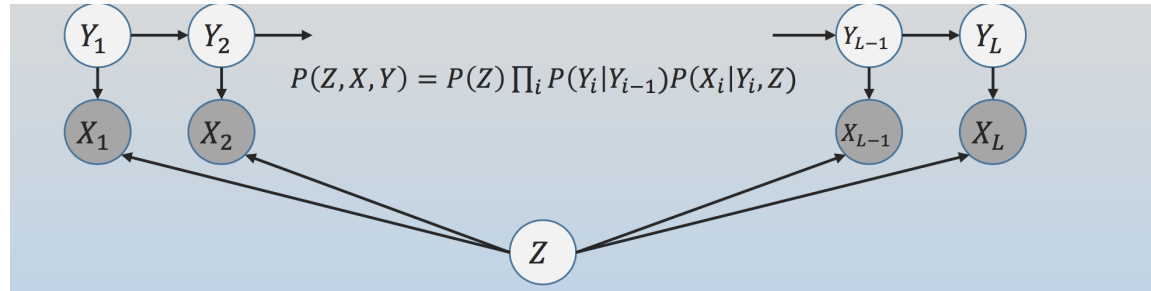


Figure 2: Parameterize network

Y_i	0	1
Y_{i-1}		
0	p_Y	$1 - p_Y$
1	$1 - p_Y$	p_Y

X_i	0	1
(Z, Y_i)		
(0,0)	$1 - p_{X 00}$	$p_{X 00}$
(0,1)	$1 - p_{X 01}$	$p_{X 01}$
(1,0)	$1 - p_{X 10}$	$p_{X 10}$
(1,1)	$1 - p_{X 11}$	$p_{X 11}$

Y_1	0	1
-	0.5	0.5

Z	0	1
-	0.5	0.5

Figure 3: Parameterize network probability tables

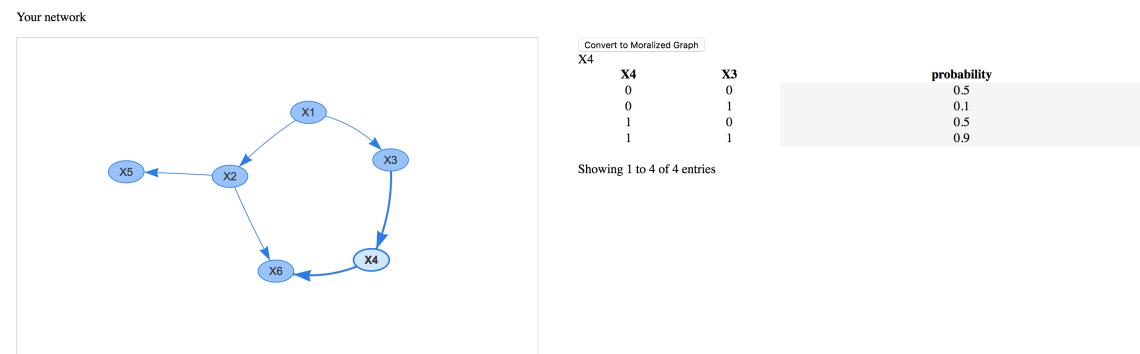


Figure 4: A simple Bayesian network representation using directed graph

- [4] Syllogismos, PGM learning,
<https://github.com/anhncs/Probabilistic-Graphical-Models>
- [5] Mark Schmidt and Kevin Murphy, MAP estimation of DAG structures,
<http://www.cs.ubc.ca/~murphyk/Software/DAGlearn/index.html>

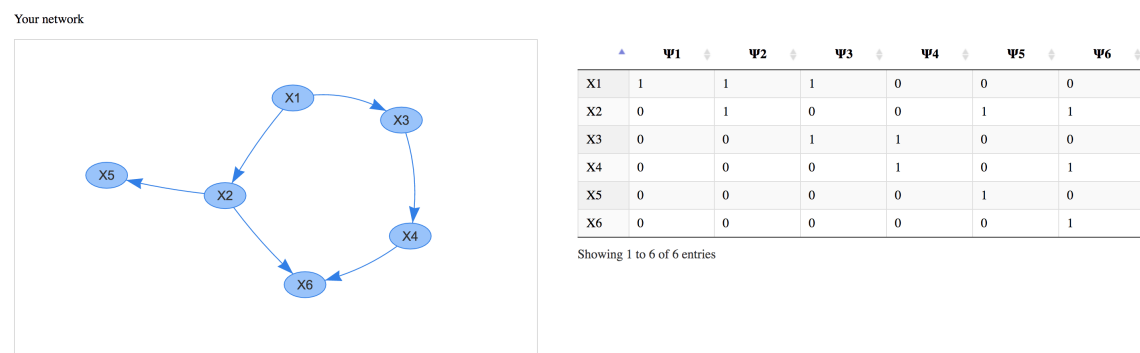


Figure 5: Showing binary membership table of the network

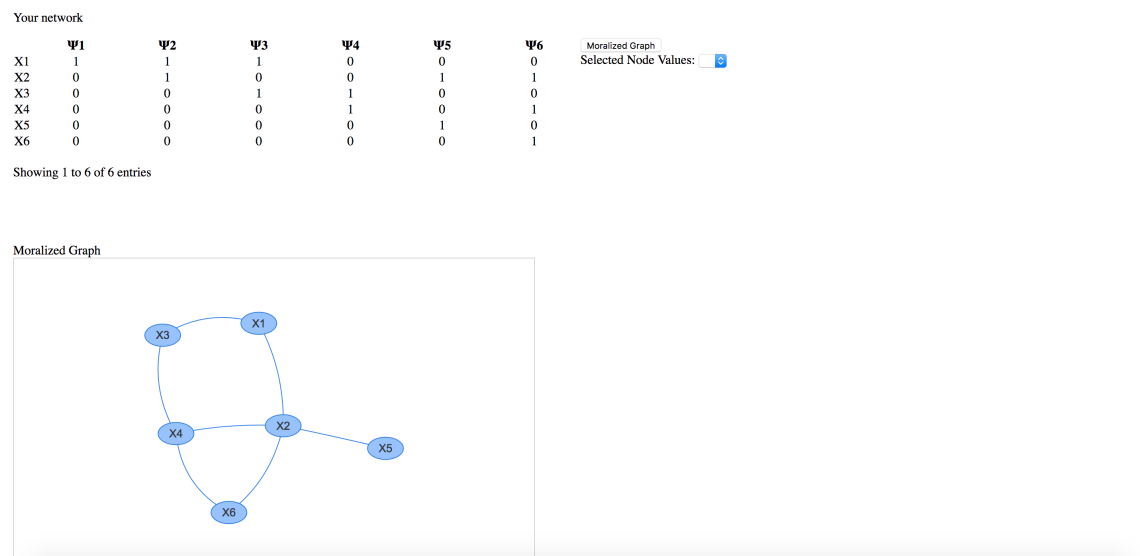


Figure 6: Moralized Graph representation

Your network

	Ψ_1	Ψ_2	Ψ_3	Ψ_5
X1	1	1	1	0
X2	0	1	1	0
X3	0	0	1	0
X5	0	0	0	0

Showing 1 to 4 of 4 entries

Moralized Graph

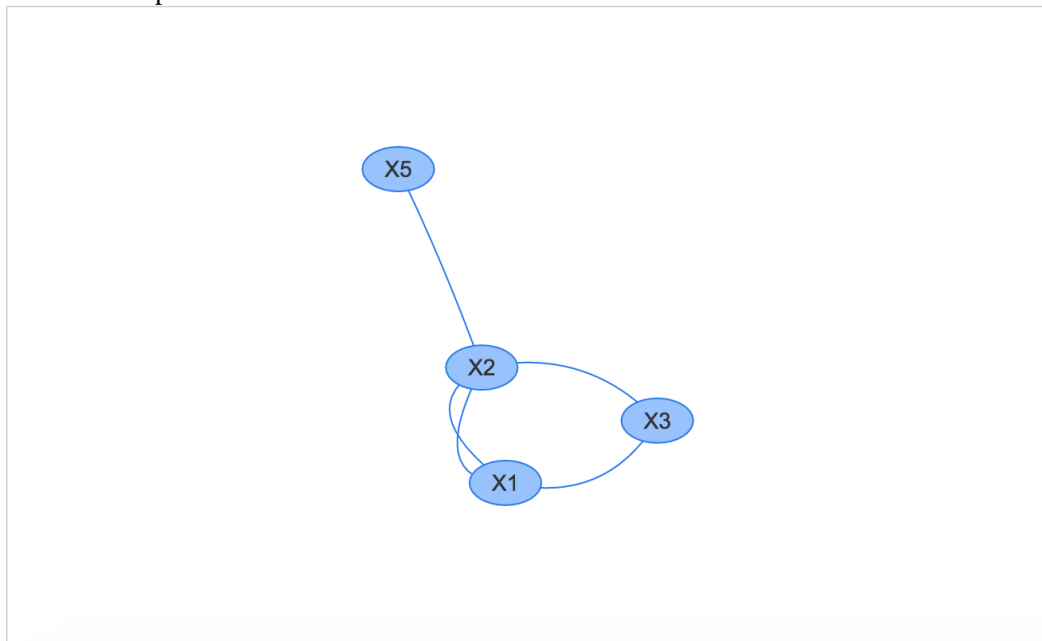


Figure 7: After X4 is eliminated from the graph. We can see a new edge was created between X2 and X3. Now X5 node is chosen to be eliminated next