Switching Activity Estimation of VLSI Circuits Using Bayesian Networks

Sanjukta Bhanja and N. Ranganathan, Fellow, IEEE

Abstract—Switching activity estimation is an important aspect of power estimation at circuit level. Switching activity in a node is temporally correlated with its previous value and is spatially correlated with other nodes in the circuit. It is important to capture the effects of such correlations while estimating the switching activity of a circuit. In this paper, we propose a new switching probability model for combinational circuits that uses a logic-induced directedacyclic graph (LIDAG) and prove that such a graph corresponds to a Bayesian network (BN), which is guaranteed to map all the dependencies inherent in the circuit. BNs can be used to effectively model complex conditional dependencies over a set of random variables. The BN inference schemes serve as a computational mechanism that transforms the LIDAG into a junction tree of cliques to allow for probability propagation by local message passing. The proposed approach is accurate and fast. Switching activity estimation of ISCAS and MCNC circuits with random and biased input streams yield high accuracy (average mean error = 0.002) and low computational time (average elapsed time including CPU, memory access and I/O time for the benchmark circuits = 3.93 s).

Index Terms—Bayesian networks, dynamic power, power estimation, probabilistic modeling, switching activity.

I. INTRODUCTION

POWER consumption in a CMOS circuit can be classified into the following three categories:

- 1) static leakage power;
- 2) short-circuit power;
- 3) dynamic power.

In a well-designed circuit, the total power dissipation is dominated by the dynamic power consumption of the nodes, which arises due to the charging and discharging of the parasitic capacitance during switching [19]. The average power consumption at a gate is given by $P(x) = 0.5 V_{dd}^2 f_{\rm clk} C_{\rm load} {\rm sw}(x)$, where V_{dd} is the supply voltage, $f_{\rm clk}$ is the clock frequency, $C_{\rm load}$ is load capacitance, and ${\rm sw}(x)$ is the average switching activity of the output node x.

It is proven that for a zero-delay model of a combinational circuit, only first-order temporal correlation is exhibited [14] as the signal possesses a Markov property. Hence, we are interested in capturing all higher order spatial correlations and first-order

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- S. Bhanja is with the Department of Electrical Engineering, University of South Florida, Tampa, FL 33620 USA (e-mail: bhanja@eng.usf.edu).
- N. Ranganathan is with the Department of Computer Science and Engineering and the Nanomaterials and Nanoelectronics Research Center, University of South Florida, Tampa, FL 33620 USA (e-mail: ranganat@csee.usf.edu).

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temporal correlation to encapsulate all the spatio-temporal dependencies in the circuit. It is our observation that the Bayesian network (BN) is a powerful tool to model switching activity preserving the various dependencies in a circuit. Further, elegant inference mechanisms exist for BN computation that make the estimation time efficient and, thus, usable for large circuits.

In this study, we model the switching in a combinational circuit using a probabilistic BN [2], [3], which allows us to capture both the temporal and spatial dependencies in a comprehensive manner. BNs are directed acyclic-graph (DAG) representations whose nodes represent random variables and the links denote direct dependencies, which are quantified by conditional probabilities of a node given the states of its parents. This DAG structure essentially models the joint probability function over the set of random variables under consideration in a compact manner. The attractive feature of this graphical representation of the joint probability function is that, not only does it make conditional dependency relationships among the nodes explicit, it also serves as a computational mechanism for efficient probabilistic updating. BNs have traditionally been used in artificial intelligence and image analysis. Their use in power estimation is new.

The core idea is to express the switching activity of a circuit as a joint probability function, which can be mapped one-to-one onto a BN that preserves the dependency model of the probability function. We first construct a logic-induced directed-acyclic graph (LIDAG) based on the logical structure of the circuit. Each signal in the circuit is a random variable in the LIDAG that can have four possible states indicating the transitions from $0 \to 0$, $0 \to 1$, $1 \to 0$, and $1 \to 1$. Directed edges are drawn from the random variables representing switching of the inputs to the random variable for switching at the output of each gate. We prove that the LIDAG, thus obtained, is a BN, which is the minimal representation that captures all the independency relationships in the circuit.

The salient features of the BN model for switching activity estimation are as follows.

- It is able to produce globally correct estimates of switching activity accounting for spatial correlation and by using only local message passing for probability updates, which makes the estimation process quick and efficient.
- 2) The computations are parallelizable and multithreadable.
- 3) It can accommodate input correlation, temporal, and spatial correlations efficiently.
- 4) BNs can accept evidence for any node and propagate the evidence in any direction. For example, it can accept switching activity statistics of an *output* node and update

the switching probabilities at all the other nodes. Thus, the network can give switching probability estimates for each node, conditioned on information about switching activity at the input or output nodes, or even at any internal node.

- 5) After a compilation process that converts the BN into a junction tree of cliques, further computation time is small. Thus, repeated computation of switching activity of the circuit with different input statistics does not require much time.
- 6) Finally, a BN in addition to pair-wise correlations [5] also models conditional independence over a set of variables, which allows it to cover a large class of probabilistic dependencies.

The organization of this paper is as follows: The relevant previous works are discussed in Section II. The proposed BN model is presented in Sections III and IV. Section V describes the computational strategies. We report experimental results in Section VI and provide conclusions and directions for further work in Section VII.

II. RELATED WORK

Nonsimulative power-estimation techniques have been proposed in the past that have reasonable accuracy and extremely fast estimation procedures. They can be purely nonsimulative [4], [6], [7] or they can be statistically simulative [16]. In statistical simulation, the estimated power is highly input sensitive. Although Monte Carlo simulation technique can use input selection efficiently [16], these techniques have problems when the estimation process has to be performed under correlated input streams.

The nonsimulative statistical techniques use knowledge about input statistics to estimate the switching activity of internal nodes. In some of the pioneering works around this idea, Najm et al. [13] estimated the mean and variance of current using probability waveforms. In [7], the concept of transition density is introduced and is propagated throughout the circuit by a Boolean difference algorithm. However, these methods have problems in handling the correlation between nodes and, hence, the estimates are inaccurate when the nodes are highly correlated. An accurate way of switching activity estimation is proposed in [6], which has a high space requirement. Tagged probability simulation is proposed in [9], which is based on the local OBDD propagation. The signal correlations are captured by using local OBDDs. However, spatio-temporal correlation between the signals is not discussed. Kapoor [11] has modeled structural dependencies, Schneider and Schlichtmann [14] used a one-lag Markov model to capture temporal dependence, and Tsui et al. [12] modeled spatial correlation. Pair-wise correlation between circuit lines were first proposed by Ercolani et al. in [8]. Marculescu et al. [18] studied temporal, spatial, and spatio-temporal dependencies by capturing pair-wise correlations. In a later effort to capture higher order correlation approximately, Marculescu et al. [5] handled higher order correlation as a composition of pair-wise correlations. Schneider et al. [15] proposed another approximate technique to model higher order spatial correlations.

The theoretical contribution of our study is that the joint probability function of a set of random variables is exactly mapped capturing higher order correlations between the signals accurately using a BN model. Earlier efforts either treat the distribution as a composition of pair-wise correlated signals between all signals [5], [8] or use an approximate solution for capturing spatial correlation [15]. Moreover, the BN models conditional independence of a subset of signals unlike in [5]. Also, in contrast to [5], the propagation mechanism does not assume any signal isotropy of conditional independence. Results show that the technique has high accuracy with low execution times for combinatorial benchmark circuits.

III. PROPOSED BN MODEL

In this section, we first prove that the modeling of switching activity of a combinational circuit is an exact model for the underlying joint probability distribution of the whole circuit. Hence, any correlation that is present in the joint probability distribution is captured in such modeling. The choice of states that each random switching variable can have captures temporal correlation completely. The way in which the conditional probabilities are provided to the boundaries of the BN is discussed in Section IV.

First, we present mathematical notions that have been proposed to capture dependencies amongst random variables. Second, we list the conditions under which a BN can be constructed to capture these dependencies. Based on these foundations, we prove that all the dependencies amongst the switching variables in a combinational circuit can be captured using a DAG-structured BN that is derived from the underlying circuit structure.

To formalize the concept of dependencies, we first present the concept of conditional independence.¹ The following discussion is fashioned after [3]. We begin with the definition of *conditional* independence among three sets of random variables.

Definition 1: Let $U = \{U_1, U_2, \dots, U_n\}$ be a finite set of variables that can assume discrete values. Let P(.) be the joint probability function over the variables in U, and let X, Y, and Z be any three subsets of U. X, and Y, and Z may or may not be disjoint. X and Y are said to be *conditionally independent* given Z if

$$P(x|y,z) = P(x|z)$$
, whenever $P(y,z) > 0$ (1)

Following Pearl [3], this conditional independence amongst X, Y, and Z is denoted as I(X,Z,Y) in which X and Y are said to be *conditionally independent* given Z. Conditional independence implies that knowledge of Z makes X and Y independent of each other. In Fig. 1, for example, let us denote the switching activity at line i by random variable X_i . U is defined as a set $\{X_1,\ldots,X_9\}$. Switching in a combinational circuit follows directed Markov property since the output of a gate is dependent only on its inputs. Thus, the random variable X_9 is completely independent of X_4 given $\{X_6,X_8\}$. Hence, $I(X_9,\{X_6,X_8\},X_4)$ is one of the many independencies that are present in the circuit.

¹This concept was introduced to the power-estimation community by Marculescu *et al.* [5] in the context of switching estimation.

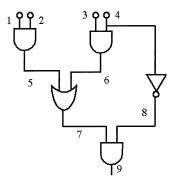


Fig. 1. Small combinational circuit.

A dependency model M of a domain should capture all these triplets, namely, (X, Z, Y) conditional independencies amongst the variables in that domain. The joint probability density function is one such dependency model. The properties involving the notion of independence are axiomatized by the following theorem.

Theorem 1: Let X, Y, and Z be three distinct subsets of U. If I(X, Z, Y) stands for the relation "X is independent of Y given Z" in some probabilistic model P, then I must satisfy the following four independent conditions:

$$I(X, Z, Y) \Rightarrow I(Y, Z, X)$$
 (symmetry) (2)

$$I(X, Z, Y \cup W) \Rightarrow I(X, Z, Y) \& I(X, Z, W)$$

$$I(X, Z, Y \cup W) \Rightarrow I(X, Z \cup W, Y)$$

$$I(X, Z, Y) \& I(X, Z \cup Y, W) \Rightarrow I(X, Z, Y \cup W)$$
 (contradiction). (5)

Proof: For proof, see [2].

Next, we introduce the concept of *d-separation* of variables in a DAG structure, which is the underlying structure of a BN. This notion of *d-separation* is then related to the notion of independence amongst triple subsets of a domain.

Definition 2: If X, Y, and Z are three distinct node subsets in a DAG D, then X is said to be d-separated from Y by $Z \langle X|Z|Y \rangle$ if there is no path between any node in X and any node in Y along which the following two conditions hold: 1) every node on the path with converging arrows is in Z or has a descendent in Z and 2) every other node is outside Z. If there exist such a path where the following two conditions hold, the path is called an active path.

In the example DAG in Fig. 2, let $X=\{X_6\}$, $Y=\{X_8\}$, and $Z=\{X_9\}$. Path $X_6\leftarrow X_4\to X_8$ is active by condition 2 and the path $X_6\to X_7\to X_9\leftarrow X_8$ is active by both conditions 1 and 2. Hence, X_6 is not d-separated from X_8 by X_9 . It is worth mentioning that if any one path is active, even though the other paths are blocked, the nodes are not d-separated. In the same example, X_7 is d-separated from X_2 by X_5 since the only path $X_2\to X_5\to X_7$ is blocked.

Definition 3: A DAG D is said to be an I-map of a dependency model M if every d-separation condition displayed in D corresponds to a valid conditional independence relationship in

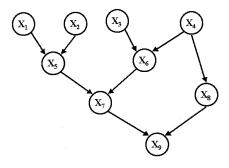


Fig. 2. BN corresponding to the circuit in Fig. 1.

M, i.e., if for every three disjoint sets of vertices X, Y, and Z, we have, $\langle X|Z|Y\rangle \Rightarrow I(X,Z,Y)$. In Fig. 2, for example $\langle X_7|X_5|X_1\rangle$ implies the independence relation $I(X_7,X_5,X_1)$ in the dependency model M formed by the random variables depicting switching activity at a line in the combinational circuit in Fig. 1.

 $\begin{array}{ll} \textit{Definition 4:} & \text{A DAG is a } \textit{minimal I-map of } M \text{ if none of its} \\ \textit{edges can be deleted without destroying its dependency model} \\ M \end{array}$

Definition 5: Given a probability function P on a set of variables U, a DAG D is called a BN of P if D is a minimum I-map of P.

There is an elegant method of inferring the minimal I-map of P that is based on the notion of a Markov blanket and a boundary DAG, which are defined below.

Definition 6: A Markov blanket of element $X_i \in U$ is a subset S of U for which $I(X_i, S, U - S - X_i)$ and $X_i \notin S$. A set is called a Markov boundary, B_i of X_i if it is a minimal Markov blanket of X_i , i.e., none of its proper subsets satisfy the triplet independence relation. For example, considering the domain U in Fig. 1, the random variable X_7 , $S = \{X_5, X_6, X_4\}$ is a Markov blanket, which means given the state of $\{X_5, X_6, X_4\}$, X_7 is independent of any other random variable in the domain that is not in S, namely, X_1 or, say, X_4 . A subset of S, $S_1 = \{X_5, X_6\}$ is a minimal Markov blanket of X_7 and, hence, a Markov boundary of X_7 , which implies that given X_5 and X_6 , X_7 is independent of the rest of the random variables in U and any proper subset of S_1 , namely, singleton sets $\{X_5\}$ or $\{X_6\}$ will not be a Markov blanket for X_7 .

Definition 7: Let M be a dependency model defined on a set $U = \{X_1, \ldots, X_n\}$ of elements, and let d be an ordering $\{X_{d1}, X_{d2}, \ldots\}$ of the elements of U. The boundary strata of M termed as B_M relative to d is an ordered set of subsets of U, $\{B_{d1}, B_{d2}, \ldots\}$ such that each B_{di} is a Markov boundary (defined above) of X_{di} with respect to the set $U_{di}(\subset U) = \{X_{d1}, X_{d2}, \ldots, X_{d(i-1)}\}$, i.e., B_{di} is the minimal set satisfying $B_{di} \subset U$ and $I(X_{di}, B_{di}, U_{di} - B_{di})$. The DAG created by designating each B_{di} as the parents of the corresponding vertex X_{di} is called a boundary DAG of M relative to d. It should be noted here that the only ordering restriction is that the variables in the Markov boundary set (of a particular variable) has to be ordered before the random variable. For the ordering of variables X_1, \ldots, X_9 in Fig. 1, the boundary strata of underlying dependency model M over the domain U is given as

$$B_M = \{ \{\phi\}, \{\phi\}, \{\phi\}, \{\phi\}, \{X_1, X_2\}, \\ \{X_3, X_4\}, \{X_5, X_6\}, \{X_4\}, \{X_7, X_8\} \}.$$

It is easily observed from the definition that the DAG in Fig. 2 is a boundary DAG corresponding to dependency model M over domain U in Fig. 1.

This leads us to the final theorem that relates the BN to I-maps, which has been proven in [3]. This theorem is the key to constructing a BN.

Theorem 2: Let M be any dependency model satisfying the axioms of independence listed in (3)–(5). If the graph structure D is a boundary DAG of M relative to ordering d, then D is a minimal I-map of M. **Proof:** For proof, see [2].

This theorem along with definitions 2–4 above specifies the structure of the BN. We use these to prove our theorem regarding the structure of the BN to capture the switching activity of a combinational circuit.

Let a combinational circuit consist of gates $\{G_1,\ldots,G_N\}$ with n being primary input signals denoted by the set $\{I_1,\ldots,I_n\}$. Let the output of gate G_i be denoted by O_i . The inputs to a gate are either a primary input signal or output of another gate. The switching of these input signal and output lines $\{I_1,\ldots,I_n,O_1,\ldots,O_N\}$ are the random variables of interest. Note that the set of output lines include both the intermediate lines and final output lines. Let X_i be the switching at the ith line, which is either an input or an output line, taking on four possible values $\{x_{00},x_{01},x_{10},x_{11}\}$ corresponding to the possible transitions $0 \to 0$, $0 \to 1$, $1 \to 0$, and $1 \to 1$.

Definition 8: An LIDAG structure LD corresponding to a combinational circuit consists of nodes X_i s representing the switching at each line and links between them is constructed as follows. The parents of a random variable representing the switching at an output line O_i of a gate G_i are the nodes representing switchings at the input lines of that gate. Each input line is either one of $\{I_1, \ldots, I_n\}$ or an output of another gate. The DAG shown in Fig. 2 is an LIDAG corresponding to the combinational circuit shown in Fig. 1.

Theorem 3: The LIDAG structure LD corresponding to the combinational circuit is a minimal I-map of the underlying switching dependency model and, hence, is a BN.

Proof: Let us order the random variables $\{X_1, X_2, \ldots, X_{N+n}\}$ such that: 1) variables representing the switching at the input lines appear first followed by those representing the output lines of the gates and 2) if a line O_i is an input to a gate whose output line is O_k , then the variable corresponding to line O_i appear before O_k .

With respect to this ordering, the Markov boundary of a node X_i is given as follows. If X_i represents switching of an input signal line, then its Markov boundary is the null set. Also, since the switching of an output line is only dependent on the inputs of the corresponding gate, the Markov boundary of a variable representing an output line consists of just those that represent the inputs to that gate. In the LIDAG structure, the parents of each node are its Markov boundary elements; hence, the LIDAG is a boundary DAG. Also, by Theorem 2 above, the LIDAG is a minimal I-map and, thus, a BN.

It is interesting to note that the LIDAG structure corresponds exactly to the DAG structure one would arrive by considering the *principle of causality*, which states that one can arrive at the appropriate BN structure by directing links from nodes that represent causes to nodes that represent immediate effects [3]. Thus, directed links in the graph denote an immediate

cause-and-effect relationship. In a combinational circuit, the immediate causes of switching at a line are the switchings at the input lines of the corresponding gate.

IV. FORMATION OF THE LOGIC-INDUCED DIRECTED-ACYCLIC-GRAPH STRUCTURED BAYESIAN NETWORK (LIDAG-BN)

We first illustrate with an example how switching in a combinational circuit at circuit level can be represented by an LIDAG-BN. We then show how the conditional probabilities that quantify the links of LIDAG-BN are specified.

Let us consider the circuit with fives gates shown in Fig. 1. We are interested in the switching at each of the nine numbered lines in the circuit. Each line can take four values corresponding to the four possible transitions $\{x_{00}, x_{01}, x_{10}, x_{11}\}$. Note that this way of formulating the random variable effectively models temporal correlation since only first-order temporal correlation is exhibited in combinational circuit under zero-delay scenario [14]. To capture all higher order spatial correlations, we form the interconnection (through edges) and quantify them by the conditional probabilities for the child-parent group in the LIDAG. The probability of switching at a line would be given by $P(X_i =$ $(x_{01}) + P(X_i = x_{10})^2$ The LIDAG structure for the circuit is shown in Fig. 2. Dependence among the nodes that are not connected directly is implicit in the network structures. For example, nodes X_1 and X_2 are independent of each other, however, they are conditionally dependent given the value of, say, node X_9 , or the transition at line 5, i.e., X_5 , is dependent on the transitions at lines 1 and 2, represented by the random variables X_1 and X_2 , respectively. Thus, the transitions of line 5 are conditionally independent of all transitions at other lines given the transition states of lines 1 and 2.

The joint probability function that is modeled by a BN can be expressed as the product of the conditional probabilities. We also say that joint probability function p factorizes in the BN. For the BN structure in Fig. 2, the corresponding joint probability density is given by the following factored form. It has to be noted that this factored form can only be obtained for circuits without a feedback as follows:

$$P(x_1, ..., x_N) = \prod_{v} P(x_v | x_{\text{parent}(v)})$$

$$P(x_1, ..., x_9) = P(x_9 | x_7, x_8) P(x_8 | x_4) P(x_7 | x_5, x_6)$$

$$\times P(x_6 | x_3, x_4) P(x_5 | x_1, x_2) P(x_4)$$

$$\times P(x_3) P(x_2) P(x_1). \tag{6}$$

The conditional probabilities of the lines that are directly connected by a gate can be obtained knowing the type of the gate. For example, $P(X_5=x_{01}|X_1=x_{01},X_2=x_{00})$ will always be 1 because if one of the inputs of an OR gate makes a transition from 0 to 1 and the other stays at 0, then the output always makes a transition from 0 to 1. A complete specification of the conditional probability of $P(x_5|x_1,x_2)$ will have 4^3 entries since each variable has four states. These conditional probability specifications are determined by the gate type. Thus, for an AND gate, if one input switches from

²Probability of the event $X_i = x_i$ will be denoted simply by $P(x_i)$ or by $P(X_i = x_i)$.

0 to 1 and the other from 1 to 0, the output remains at 0. By specifying a detailed conditional probability, we ensure that the spatio-temporal effect (first-order temporal and higher order spatial) of any node are effectively modeled.

The last four terms in the right-hand side of (7) represent the statistics of the input lines. Given the statistics of the input lines, we would like to infer the probabilities of all the other nodes. A brute-force way of achieving this would be to compute the marginal probabilities by summing over possible states, thus, $P(x_9,x_1)=\sum_{x_2,\dots,x_8}P(x_1,\dots,x_9)$. This, obviously, is computationally very expensive and, in addition, does not scale well. In Section V, we show how the structure of the BN can be used to efficiently compute the required probabilities.

V. COMPUTATIONS USING A BN

It would be computationally convenient if we could compute probabilities in a BN by local message passing. Unfortunately, we cannot directly update a BN by local message passing if the underlying undirected graph structure has cycles. Thus, the first step of the computation strategy is to transform the original DAG structure into a undirected tree structure whose nodes are subsets of the original random variables [2]. Such a tree is referred to as the junction tree of cliques of random variables and the process is referred to as the BN compilation process. After the compilation process, the required probabilities can be easily computed by local message passing in a simple propagation algorithm. In this section, we will first outline the steps involved in the compilation process and then sketch the propagation algorithm. We should point out to the reader that there are many BN softwares that will automatically do these steps. For this study, we used the HUGIN BN software for the implementation of inference engine, which performs the steps discussed throughout this section.3

A. Network Compilation

The compilation process ensures the following two aspects in that: 1) it builds a tree of cliques, which implies that there is no loop in the tree and 2) the tree is built in a special fashion to help a local message passing scheme described in Section V-B. The first step of the compilation process is the formation of the corresponding moral graph. This is followed by triangularization of the moral graph. The clique set is then identified and a junction tree of cliques is formed. These steps are shown as a flowchart in Fig. 3.

1) Moral Graph: The first step of the compilation process is to create an undirected graph structure called the moral graph (denoted by D^m) given the BN DAG structure (denoted here by D). The moral graph represents the Markov structure of the underlying joint function [2]. If probability function P factorizes recursively like that shown in (7) in a DAG D, it also can be expressed in a factorized form in the moral graph of D [2]. From a DAG, which is the structure of a BN, a moral graph is obtained by adding undirected edges between the parents of a common child node and dropping the directions of the links (shown via the dashed line in Fig. 4). This step ensures that every parent

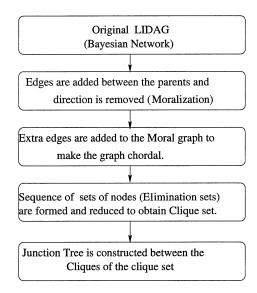


Fig. 3. Network compilation steps.

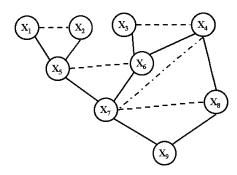


Fig. 4. Triangulated undirected graph structure that encodes that same dependencies as the DAG structure shown in Fig. 2. The solid links are from the BN. The dashed links are added during moral-graph construction. The dotted–dashed link is added during the final triangulation step.

child set is a complete subgraph. A moral graph is undirected and, due to the added links, some of the independencies displayed in a DAG will not be graphically visible in a moral graph. Some of the independencies that are lost in the transformation contributes to the increased computational efficiency but does not affect the accuracy [2]. The independencies that are graphically seen in the moral graph are used in the inference process to ensure local message passing.

2) Chordal Graph: This step is the triangulation of the moral graph to make it chordal (often referred to as triangulated). We add additional links to the moral graph so that all cycles longer than three nodes are broken into cycles of three nodes. In our example, the dashed–dotted line between X_4 and X_7 in Fig. 4 is added to the moral graph to triangulate it. One can always add all possible links between nodes to make the graph chordal, transforming the moral graph into a complete graph; however, the challenge is to make a graph chordal using a minimum number of additional links. This problem is, in general, NP hard. Hence, heuristics are used. A sub-optimal solution to find the minimum number of links may hamper computational efficiency, but the resultant graph will be a chordal one, thus maintaining the accuracy of the inference process. In HUGIN, a minimum fill-in edges heuristics is used [2].

³[Online]. Available: http://www.hugin.com/

3) Junction Tree:

Definition 9: A junction tree T of cliques is a tree with nodes representing cliques (collection of completely connected nodes) and between any two cliques C_i and C_j in the tree T there is a unique path. Also, the elements in the intersection set $C_i \cap C_j$ are present in all the cliques in that unique path between C_i and C_j . This property in a junction tree is also referred to as the running intersection property.

The existence of at least one junction tree in a chordal graph is guaranteed [2]. The running intersection property of a junction tree is an important one and is utilized in the local message-passing-based updating algorithm described later. It is worth noting that it is possible to construct a tree of node cliques from the moral graph itself, but it will not have the running intersection property and, hence, it will not be suitable for local message-passing-based updating.

4) From Moral Graph to Junction Tree: In this section, we outline the details of how a moral graph is transformed into a junction tree. The process essentially consists of the three following steps.

Step 1: Moral Graph to Elimination Set

All the vertices of the moral graph are first unnumbered. An unnumbered vertex that needs the minimum number of additional edges between its neighbors (considering both numbered or unnumbered neighbors) is chosen. This vertex is then labeled with the highest available node number, say, i, starting from a number equal to the total number of nodes. A set C_i is then formed consisting of the selected vertex and its still unnumbered neighbors. Edges are filled in between any two unlinked nodes in this set C_i . The maximum available node number i is decremented by one. This process is repeated until there is no unnumbered nodes. The resultant graph is now a chordal one. The generated sequence of cliques C_i 's is termed the *elimination set* of cliques of the graph, a subset of which is used to construct the junction tree.

Due to the way the cliques in the elimination set are derived, they will possess the running intersection property [2]. It can also be shown that the set of cliques in the final junction tree of a graph is subset of the elimination set [2].

In our example moral graph, node X_9 is first selected since no fill-in edge is needed because all the neighbors (remember the moral graph is undirected) are already linked. This node X_9 is assigned the number 9—the total number of nodes in the graph. The set C_9 is then formed by nodes $\{X_9, X_8, X_7\}$. The nodes X_8 and X_7 are not yet numbered. For the second cycle, the nodes X_8 , X_7 , X_6 , and X_4 cannot be selected, as they each would require one fill-in edges amongst its neighbors, whereas the neighbors of X_3 does not require any fill-in edges. Hence, X_3 is numbered 8 in our example and C_8 is formed by $\{X_3, X_4, X_6\}$. For the third cycle, we then select X_2 , numbering it as 7 and forming $C_7 = \{X_2, X_1, X_5\}$. In the fourth cycle, node X_1 is assigned 6 and $C_6 = \{X_1\}$ is formed. We then select X_5 , assign a number 5, and form $C_5 = \{X_5, X_6, X_7\}$. Node X_8 is assigned number 4, and $C_4 = \{X_8, X_7, X_4\}$ is formed. In this step, a fill-in edge between X_4 and X_7 is added. We then assign the number 3 to X_7 , the number 2 to X_6 , and the number 1 to X_4 . Thus, we form $C_3 = \{X_7, X_6, X_4\}, C_2 =$ $\{X_6\}$ and $C_1 = \{X_4\}$.

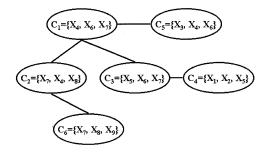


Fig. 5. Junction tree of cliques.

Step 2: Reducing the Elimination Set

It is shown in [2] that if C_1,\ldots,C_k is a sequence of sets having running intersection property and $C_t\subseteq C_p$ for some $t\neq p$, then the ordered set $C'=\{C_1,\ldots,C_{t-1},C_p,C_{t+1},\ldots,C_{p-1},C_{p+1},C_k\}$ also has running intersection property. By this property, C_t can be eliminated for all $C_t\subseteq C_p, p\neq t$. Hence, the elimination set can be reduced to obtain the minimal ordered set of cliques called the *clique set* representing the chordal graph completely. This ordering of cliques is guaranteed to possess the running intersection property.

Step 3: Elimination Set to Junction Tree

For each clique C_i of the clique set ordered to have running intersection property, we have to add a link to clique C_j , where j is any one of the set $\{1,2,\ldots,i-1\}$ such that $C_i\cap (C_1\cup C_2\cup\cdots\cup C_{i-1})\subseteq C_j$.

Fig. 5 shows the junction tree for our running example. It can be observed that the cliques in our example are $C_1 = \{X_4, X_6, X_7\}$, $C_2 = \{X_7, X_4, X_8\}$, $C_3 = \{X_5, X_6, X_7\}$, $C_4 = \{X_1, X_2, X_5\}$, $C_5 = \{X_3, X_4, X_6\}$, and $C_6 = \{X_7, X_8, X_9\}$. This cliques are obtained by reducing the elimination set. The edges between them are formed in the algorithm in step 3. For our example in Fig. 5, for the clique C_6 , we have to find the set that contains the following intersection set:

$$C_6 \cap (C_1 \cup C_2 \cup \dots, \cup C_6)$$
= $\{ \{X_7, X_8, X_9\} \cap \{X_4, X_6, X_7, X_8, X_5, X_1, X_2, X_3\} \}$
= $\{X_7, X_8\}$.

It is obvious that $\{X_7, X_8\} \subseteq C_2$, hence, we add a link between C_6 and C_2 . Every clique with an edge between them will have a nonnull set of nodes common between them, which is referred to as the *separator set*. These common variables play key role in evidence propagation. If there were two cliques that are not connected by an edge and still have common variables, then these variables must be present in all the cliques in between the unique path between the two cliques to guarantee correct probability values for the random variables of the entire network while probabilities are updated by local propagation. As it can be seen, C_3 and C_6 both contain X_7 , and X_7 is also present in all the cliques in the path from C_3 to C_6 , namely, in C_1 and C_2 . This helps to preserve probabilities of the random variables of the entire network while updating an evidence by local message passing.

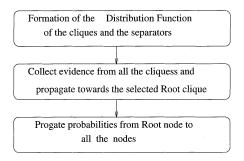


Fig. 6. Steps involved in propagation of evidence in the junction tree.

B. Propagation in Junction Trees

Once the junction tree of cliques is formed in the compilation process, we need to form the distribution function for the cliques. We then follow a two-phase algorithm for message propagation. All the messages propagated are between neighboring cliques. These steps are shown in Fig. 6.

It can be proven that the dependency properties of a DAG, which carry over to moral graphs, are also preserved in the triangulated moral graph [2]. The joint probability function that factorizes on the moral graph will also do so on the triangulated moral graph since each clique in the moral graph is either a clique in this graph or subset of a clique. Let $\{\mathbf{x}_c\}$ be the set of nodes in clique c in the junction tree. The joint probability function over these variables is denoted by $p(\mathbf{x}_c)$. Let $\{\mathbf{x}_s\}$ be the set of nodes in a separator set s between two cliques in the junction tree. The joint probability function over these variables is denoted by $p(\mathbf{x}_s)$. Let CS denote the set of all cliques and SS denote the set of separators in the junction tree. The joint probability function factorizes over the junction tree in the following form [2]:

$$p(x_1, \dots, x_N) = \frac{\prod_{c \in CS} p(\mathbf{x_c})}{\prod_{s \in SS} p(\mathbf{x_s})}.$$
 (7)

A separator set s is contained in the two neighboring cliques c_1 and c_2 . If we associate each of the product terms over the separators in the denominator with one its two neighboring cliques, say, c_1 , then we can write the joint probability function in a pure product form as follows. Let $\phi_{c_1}(\mathbf{x_{c1}}) = p(\mathbf{x_{c_1}})/p(\mathbf{x_s})$ and $\phi_{c_2}(\mathbf{x_{c_2}}) = p(\mathbf{x_{c_2}})$, then the joint probability function is expressed as

$$P(x_1, \dots, x_N) = \prod_{c \in CS} \phi_c(\mathbf{x}_c)$$
 (8)

where the factors $\phi_c(\mathbf{x_c})$ are also commonly referred to as the potential function over the nodes $\{x_c\}$ in clique c and CS is the set of cliques. These functions $\phi_c(\mathbf{x_c})$ s can be formed by multiplying the conditional probabilities from the input BN specification of nodes in the clique c.

Let us now focus on the information flow in the neighboring cliques to understand the key feature of the Bayesian updating scheme. As shown in Fig. 7, let two cliques A and B have probability potentials ϕ_A and ϕ_B , respectively. Let S be the set of nodes that separates cliques A and B. Let us say that there is new evidence for some nodes. This will change the probabilities of all the other nodes such that two neighboring cliques

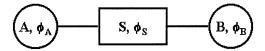


Fig. 7. Two cliques of the junction tree along with the separator set.

agree on probabilities on the node set S, which is their separator. To achieve this, we first compute the marginal probability of S from the probability potential of clique A and then use that to scale the probability potential of B, as captured by (10). To achieve this, we need to transmit the scaling factor along the link, and this process is referred to as message passing. We have to repeat this process in the reverse direction by computing the marginal probability of S from probability potential of clique B and then use that to scale the probability potential of A. This will ensure that evidence at both the cliques are taken into account. New evidence is absorbed into the network by passing such local messages. The pattern of the message is such that the process is multithreadable and partially parallelizable. Since the junction tree has no cycles, messages along each branch can be treated independently of the others as follows:

$$\phi_S^* = \sum_{X \in A, X \notin S} \phi_A \tag{9}$$

$$\phi_B^* = \phi_B \frac{\phi_S^*}{\phi_S}. (10)$$

It would appear that we will need to initiate a message passing over the whole network for each new evidence, however, this need not be the case. In fact, there is a two-phase message passing scheme that can integrate all new evidence in two passes. In this scheme, a clique is selected from the junction tree to be the root node. In the first phase, i.e., the collection phase, all the leaf nodes of the tree send messages toward the root node, which are recomputed according to (9) and (10) at each node along the way. Once the messages from all the leaf nodes have been received, the second phase is initiated when messages are passed back from the root clique toward the leaves. Note that this ensures that, along any one link, we have messages along both directions, thus ensuring all nodes have been updated based on information from all the new evidence.

Thus, junction-tree-based probabilistic propagation serve the following two purposes in that: 1) they solve the problem of updating probabilities in a loop and 2) the cliques have a running intersection property, thus updating is performed locally from one clique to the neighboring clique just once. In Section VI, we discuss the experiments and results obtained for benchmark circuits.

VI. EXPERIMENTAL RESULTS

We mapped 14 ISCAS and five MCNC circuits to their corresponding LIDAG structured BNs. The conditional probabilities are predetermined by the type of gate connecting the parents and child. We have already discussed inSection IV that each node in BN represents switching at a line in the circuit and can be in one of the four states $(x_{00}, x_{01}, x_{10}, x_{11})$. Conditional probabilities

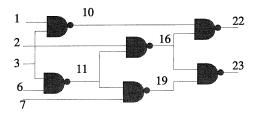


Fig. 8. Combinational circuit c17.

are assigned by the logical relationship of the gate. In combinational circuits, it is fairly common to encounter gates with a large number of inputs. This may be a problem in terms of the conditional probability table size, which grows exponentially with the number of inputs. We solve this problem by adding a few two-input dummy nodes still preserving the logic structure. The switching activity on the original nodes are modeled exactly by this method. We do not consider the switching activity on dummy nodes for the error estimates. For example, a four-input NAND gate is modeled as two AND gates followed by a NAND gate. We ignore switching activity on the outputs of the first two AND gate and the inputs of the last NAND gate. Thus, the switching estimates of the terminals of the original NAND is captured accurately without adding switching activity on dummy nodes. We used HUGIN's BN tool for compiling the junction tree and propagating the probabilities. We also performed logic simulation providing "ground truth" estimates of switching.

A. Small Example

Let us tabulate the switching activity estimation of each node of the c17 ISCAS benchmark. As can be seen in c17 (see Fig. 8), nodes 10 and 11 and 16 and 19 are highly correlated to each other. Since nodes 10 and 16 both are dependent on node 3, they are also correlated. Table I shows all the circuit nodes under random input sequences and under biased sequences. As can be observed in both cases, we have extremely accurate estimates on all nodes of the circuit.

B. Random Inputs

We tabulate the results of switching activity estimation at circuit level by the LIDAG structured BNs in Tables II and III. We also performed simulation with pseudorandom inputs for comparison of the switching estimates. In Table II, we present the circuits where we model the circuit as a single BN without segmentation.

It is evident that if the circuit is considerably large or if the circuit has high connectivity between nodes, then the size of the cliques formed becomes large. This results in large probability potential tables and might make handling large circuits in one stage impossible with limited memory resources. To handle large circuits, we adopt a divide-and-conquer strategy. We first segment the large circuit into smaller ones and then estimate switching activity in them by ensuring consistency of singleton probability of the nodes at the boundary. In Table III, we present the circuits where we model the circuit as multiple BNs. In both Tables II and III, columns 2 and 3 provide the mean and standard deviation of the error of switching activity, which is the error

TABLE I
COMPARISON OF ESTIMATED SWITCHING ACTIVITY BY BN MODELING AND
SIMULATED SWITCHING ACTIVITY OF EACH NODE OF BENCHMARK c17

nodes of	Random	Inputs	Biased Inputs	
c17	Est. Sw_{BN}	Sim. Sw	Est. Sw_{BN}	Sim. Sw
1	0.5	0.499	0.4	0.401
2	0.5	0.500	0.4	0.402
3	0.5	0.499	0.4	0.4
6	0.5	0.5	0.4	0.4
7	0.5	0.5	0.4	0.4
8	0.5	0.499	0.4	0.4
9	0.5	0.499	0.4	0.4
10	0.375	0.375	0.160	0.160
11	0.375	0.374	0.160	0.160
14	0.375	0.374	0.160	0.160
15	0.375	0.374	0.160	0.160
16	0.469	0.469	0.380	0.382
19	0.469	0.469	0.380	0.379
20	0.469	0.469	0.380	0.382
21	0.469	0.469	0.380	0.382
22	0.492	0.493	0.435	0.436
23	0.492	0.492	0.480	0.481

TABLE II
EXPERIMENTAL RESULTS ON SWITCHING ACTIVITY ESTIMATION BY SINGLE
BN MODELS FOR ISCAS'85 AND MCNC'89 BENCHMARK CIRCUITS FOR
RANDOM INPUT SEQUENCES

	Error statistics		Overall % Error and time			
	over each node		for all nodes			
Circuits	μ_{Err}	σ_{Err}	% Error	Elapsed Time (s)		
				Total	Update	
c17	0.0002	0.0004	0.02%	<.001	<.001	
max_flat	0.0004	0.0005	0.10%	< 0.001	<.001	
voter	0.0002	0.0005	0.04%	0.11	<.001	
count	0.0001	0.0006	0.03%	0.33	<.001	
comp	0.0000	0.0003	0.00%	0.22	<.001	
pcler8	0.0002	0.0007	₅₇ Q.Q3%	0.11	<.001	

TABLE III

EXPERIMENTAL RESULTS ON SWITCHING ACTIVITY ESTIMATION BY SINGLE
BN MODELS FOR ISCAS'85 AND MCNC'89 BENCHMARK CIRCUITS FOR
RANDOM INPUT SEQUENCES

	Error statistics		Overall % Error and time			
	over each node		for all nodes			
Circuits	μ_{Err}	σ_{Err}	% Error	Elapsed Time (s)		
				Total	Update	
b9	0.0004	0.0020	0.10%	0.33	<.001	
c8	0.0002	0.0017	0.05%	0.93	<.001	
alu4	0.0001	0.0198	0.86%	0.54	<.001	
malu4	0.0011	0.0204	0.23%	0.21	<.001	
c432	0.0111	0.0300	2.08%	1.33	<.001	
c499	0.0002	0.0038	0.09%	0.33	<.001	
c880	0.0012	0.0088	0.57%	1.24	<.001	
c1355	0.0015	0.0189	0.41%	2.29	<.001	
c1908	0.0009	0.0090	0.22%	6.09	<.001	
c3540	0.0029	0.0400	0.79%	8.62	<.001	
c5315	0.0035	0.0269	0.80%	15.08	<.001	
c6288	0.0140	0.0465	3.34%	18.70	<.001	
c7552	0.0031	0.0460	0.70%	18.26	<.001	

between the switching activity estimated and the switching activity obtained through simulation on each node, respectively. Moreover, we present the percentage error between the average switching activity estimated over all the nodes with average switching activity through simulation in column 4 (This column is presented separately from column 2 to indicate the percentage

error). The elapsed total time for BN-based estimation, which is time to compile and time to propagate, is shown in column 5. We separately show the time for propagation of evidence in column 6. As can be observed, the propagation time is extremely low, irrespective of the size of the circuits.

The platform used here is a single-processor DELL PC with 250-MB RAM and 450-MHz clock speed. We want to emphasize that the time estimates are obtained from a Windows-based timing feature (function ftime in VISUAL C++) that provides the total time spent in a function module, which includes memory access and I/O time along with CPU time. In fact, while handling a larger circuit, CPU time is an inadequate indicator in terms of the actual time due to high memory accesses by the estimation algorithms.

As can be observed from Table II, the circuits that are small (namely, c17, comp, count, pcler8, etc.) and, thus, do not require any segmentation. The BN is extremely accurate and the standard deviation of error calculated on each node is extremely low, indicating that switching at every node is estimated accurately. The estimation time is low. The circuits that are segmented, shown in Table III, have a standard deviation of the order of 0.02. The maximum error that is encountered is around 3.5% for c6288. The minimum error is zero. Out of the 19 benchmark circuits, 17 of them had less than 1% error and two of them have an error between 1%-3.5%. It can also be observed that updating and propagating all the input evidences in almost all the circuits were performed in the order of 1 ms. This is advantageous particularly if the switching activity has to be estimated for different input signal statistics. The circuits can be pre-compiled; only propagation has to be done for different input statistics. We found that the BN is an accurate model of switching activity. We analyzed the circuits that showed a standard deviation above 0.03 (namely, c6288 and c7552), and we observed that errors occurred for some nodes in the segment boundaries. We are currently investigating an accurate way to handle segmentation so that errors do not propagate through the segments. For all the large circuits, including c5315, c6288, c7552, we have very low mean error, which is encouraging considering the size of the circuit. With accurate segmentation, we can achieve very high accuracy uniformly throughout the circuit with considerably low elapsed time.

VII. CONCLUSIONS AND FUTURE DIRECTIONS

This paper has introduced a switching activity estimation tool that encapsulates all the dependencies, both in the internal nodes and inputs, in a reasonable time and with high accuracies. We have shown results of the estimated switching activity for pseudorandom inputs and biased inputs. The results are very competitive in terms of accuracy and the elapsed time for estimation. The model handles spatio-temporal dependencies between the nodes. It also models conditional dependencies of the nodes. The proposed model is accurate and captures higher order correlation among lines and is not restricted to pair-wise correlations. Our future effort focuses on the segmentation techniques and input modeling for capturing spatial correlation at the primary inputs using the same BN model.

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Sanjukta Bhanja received the B.E. degree in electrical engineering from Jadavpur University, Calcutta, India, the M.Sc. (Engg.) degree in electrical engineering from the Indian Institute of Science, Bangalore, India, and the Ph.D. degree in computer science and engineering from the University of South Florida, Tampa.

She is currently an Assistant Professor with the Department of Electrical Engineering, University of South Florida. Her research interests includes design automation, low-power VLSI design, power estima-

tion, optimization, system-on-a-chip, probabilistic modeling, VLSI testing, and expert systems-based modeling.



N. Ranganathan (S'81–M'83–SM'92–F'02) received the B.E. degree (with honors) in electrical and electronics engineering from Regional Engineering College, Tiruchirapalli, University of Madras, Madras, India, in 1983, and the Ph.D. degree in computer science from the University of Central Florida, Orlando, in 1988.

He is currently a Professor with the Department of Computer Science and Engineering and the Nanomaterials and Nanoelectronics Research Center, University of South Florida, Tampa. He has developed

many special-purpose VLSI chips for computer vision, image processing, pattern recognition, data compression, and signal-processing applications. He has authored or coauthored over 175 papers in reputed journals and conferences. He co-holds five U.S. patents. His research interests include VLSI system design, design automation, power estimation and optimization, high-level synthesis, embedded systems, and computer architecture.

Dr. Ranganathan has served on the Program Committees of several conferences and on the Editorial Boards of the IEEE TRANSACTIONS ON VERY LARGE SCALE INTEGRATION (VLSI) SYSTEMS, the IEEE TRANSACTIONS ON CIRCUITS AND SYSTEMS—PART II: ANALOG AND DIGITAL SIGNAL PROCESSING, and the IEEE TRANSACTIONS ON CIRCUITS AND SYSTEMS FOR VIDEO TECHNOLOGY.