# Using Lower Bounds during Dynamic BDD Minimization

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### Abstract

Ordered Binary Decision Diagrams (BDDs) are a data structure for representation and manipulation of Boolean functions often applied in VLSI CAD. The choice of the variable ordering largely influences the size of the BDD; its size may vary from linear to exponential. The most successful methods for finding good orderings are based on dynamic variable reordering, i.e. exchanging of neighboring variables. This basic operation has been used in various variants, like sifting and window permutation.

In this paper we show that lower bounds computed during the minimization process can speed up the computation significantly. First, lower bounds are studied from a theoretical point of view. Then these techniques are incorporated in dynamic minimization algorithms. By the computation of good lower bounds large parts of the search space can be pruned resulting in very fast computations. Experimental results are given to demonstrate the efficiency of our approach.

# 1 Introduction

Decision Diagrams (DDs) are often used in VLSI CAD systems for efficient representation and manipulation of Boolean functions. The most popular data structure are ordered Binary Decision Diagrams (BDDs) [3, 5].

nary Decision Diagrams (BDDs) [3, 5]. But, as well known BDDs are very sensitive to the variable ordering, i.e. the size of a BDD (measured in the number of nodes) may vary from linear to exponential. Finding the optimal variable ordering is an NP-hard problem [2] and the best known algorithm has runtime  $O(n^2 \cdot 3^n)$  [7], where n denotes the number of variables.

This is the reason why many authors presented heuristics for finding good variable orderings from circuit descriptions in the last few years (see e.g. [8]). The most promising methods for BDD minimization are based on *Dynamic Variable Ordering* (DVO) [9], i.e. improving graph size using exchanges of neighboring variables. The best results measured in the number of nodes of the resulting BDD were obtained using sifting [14, 13], but unfortunately sifting is very time consuming for large functions. For this, recently in [12] an algorithm has been proposed how to partition the search space to improve sifting runtimes. But this algorithm is largely dependent on the initial variable ordering. Another approach based on "sampling" has been suggested in [15], but dependent on the chosen candidates the quality of the result varies widely, i.e. the results can be up to a factor of two worse than "classical" sifting. In this paper we investigate the use of lower bound tech-

In this paper we investigate the use of lower bound techniques during dynamic minimization. Lower bounds have so far mainly be used in exact BDD minimization [10, 11, 6]. We first theoretically analyze how lower bounds can be computed for the exchange of variables. Then it is shown how these theoretical results can be used by integrating the tech-

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niques in the sifting algorithm [14]. The method has been incorporated in the CUDD package [16]. We give a large set of experimental results to show the quality of our approach. Using lower bounds the number of exchanges can be reduced by more than 50% on average resulting in a speedup of nearly 70%. Furthermore, we study heuristic methods to "finetune" sifting by relaxation of lower bounds. It turns out that the use of lower bounds gives a very robust method to trade off runtime versus quality.

# 2 Preliminaries

### 2.1 Binary Decision Diagrams

As well-known, each Boolean function  $f : \mathbf{B}^n \to \mathbf{B}$  can be represented by a *Binary Decision Diagram* (BDD) [3], i.e. a directed acyclic graph where a Shannon decomposition is carried out in each node. In the following, only reduced, ordered BDDs are considered and for briefness these graphs are called BDDs. (For more details see [5].)

In the following we make use of *Complement Edges* (CEs) without mentioning it further. (Note that all results shown here directly transfer to BDDs without CEs.) We denote the number of nodes labelled with a variable  $x_i$  with label $(x_i)$ . For a subset  $A \subseteq \{x_1, \ldots, x_n\}$  of variables, this definition is extended to label $(A) = \sum_{x_i \in A} label(x_i)$ .

# 2.2 Exchange of Neighboring Variables

The basic operation of dynamic variable ordering is the exchange of adjacent variables [9, 14]. The exchange is performed very quickly since only edges must be redirected within these levels. Thus, the size is optimized without a complete reconstruction of the BDD. Only local transformations for the two levels are performed. This is due to the observation that BDDs are a canonical form. The exchange of two variable does not change the sub-BDDs of other levels.

## 2.3 Sifting

The sifting algorithm [14] successively considers all variables of a given BDD. When a variable is chosen, the goal is to find the best position of the variable, assuming that the relative order of all other variables remains the same. In a first step, the order in which the variables are considered is determined. This is done by sorting the levels according to their size with largest level first. To find the best position, the variable is moved across the whole BDD. In [14], this is done in three steps:

1. The variable is exchanged with its successor variable until it is the last variable in the ordering.

2. The variable is exchanged with its predecessor until it is the topmost variable.

3. The variable is moved back to the closest position which has led to the minimal size of the BDD.

Some improvements to the original sifting algorithm have already been proposed:

**Upper limit:** As the size of the BDD can grow much during the movement of one variable, it is possible to set an upper limit to the growth of the BDD. If this limit is exceeded, moving into this direction is aborted. This avoids large intermediate BDDs.

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**Closest end:** The considered variable is not always moved downwards first. Instead it is moved to the closest end and then to the opposite one.

Several other techniques have been proposed, like the use of symmetries and of interaction  $matrix^1$  [13]. We make use of these methods in the following without mentioning it further.

## 3 Lower Bounds on BDD Sizes

During sifting we want to stop moving a variable in a direction as early as possible if no further improvement can be obtained. In our approach we make use of an efficient lower bound computation that gives information that further size reduction is not possible. In this section we prove lower bounds and show by examples that they are tight. In the next section we explain how these bounds are incorporated in the sifting algorithm.

If the modification of the variable ordering is not restricted, it is well known that an exponential difference is possible [3]. However, if the kind of modification is restricted in some way (which is the case if used during sifting, since only one variable changes its location), stricter lower bounds can be given.

A very general lower bound technique has been proposed in [4]. The lower bound holds if the variables of the upper part of the BDD remain unchanged, but the variable order of the lower part can be modified arbitrarily. If used for exact BDD minimization, significant reductions in runtime are possible [6]. However, if used for sifting, these lower bounds are too general to give good estimates for the size of the resulting BDD. Futhermore, the computation of these lower bounds is much too time consuming, since a traversal of the whole graph is needed.

#### 3.1 Bounds for Level Exchanges

In the following, the node increase for the exchange of neighboring levels is examined. In the whole section it is assumed that the function essentially depends on all its variables.

Let  $f : \mathbf{B}^n \to \mathbf{B}^m$  be a Boolean function represented by a BDD. Then the following theorem holds (see [1]):

**Theorem 1** Let  $1 \leq i < n$ . If the levels *i* and *i* + 1 are exchanged, the resulting level sizes label'(·) are bounded by

$$1 \leq \text{label}'(x_i) \leq 2 \cdot \text{label}(x_i)$$

and

$$\frac{1}{2} \cdot \operatorname{label}(x_{i+1}) \leq \operatorname{label}'(x_{i+1}) \leq \operatorname{label}(x_i) + \operatorname{label}(x_{i+1}).$$

All other levels remain unchanged.

**Proof**: We consider the four cases:

Case 1:  $1 \leq \text{label}'(x_i)$ : By assumption, the function depends on  $x_i$ , thus there must be at least one node in that level.

Case 2: label' $(x_i) \leq 2$  · label $(x_i)$ : The inequation holds because each node in level *i* which depends on  $x_{i+1}$  can lead to at most two nodes, one for each possible value of  $x_{i+1}$ . If a node does not depend on  $x_{i+1}$ , it remains unchanged.

Case 3:  $\frac{1}{2} \cdot \operatorname{label}(x_{i+1}) \leq \operatorname{label}'(x_{i+1})$ : The inequation is simply the inversion of the previous inequation, i.e. exchanging two levels twice leads to the original representation.

Case 4: label' $(x_{i+1}) \leq label(x_i) + label(x_{i+1})$ : The inequation holds because the number of cofactors which have to be represented in level *i* cannot be larger than the number of cofactors which are already represented in levels *i* and *i* + 1 (see [7]). Since each node in the BDD represents a cofactor with respect to all variables in the upper part, this number cannot change by a local operation.

All results hold for BDDs with and without CEs. 
$$\Box$$



Figure 1: Result of one level exchange to the node count

The bounds proven in the theorem are tight. This can be seen as follows:

**Example 1** Consider the function with two outputs given in Figure 1. If  $x_i$  is the topmost variable the size of the first two levels is 4, while the number of nodes is reduced by a factor of two if the two topmost variables are exchanged.

**Remark 1** If neighboring levels do not interact, the resulting level sizes remain unchanged, i.e.  $label'(x_i) = label(x_i)$ and  $label'(x_{i+1}) = label(x_{i+1})$ . This can be checked efficiently using the interaction matrix.

#### 3.2 Lower Bounds for one Variable

In this section we give lower bounds for the BDD size if a single variable is moved in the variable ordering and the relative order of all other variables remains unchanged.

A trivial lower bound (also used in [16]) is the following: Assume variable  $x_i$  is in level *i*. Moving level *i* down to level *j*, all nodes which are above level *i* (i.e. in levels  $1, \ldots, i-1$ ) and all nodes below level *j* (i.e. in levels  $j + 1, \ldots, n$ ) do not change. Furthermore, nodes between these levels for which the variable does not interact with  $x_i$  do not change. Thus, the sum of these three numbers constitutes a lower bound. But since this technique only sums up the number of nodes which are not affected by moving variable it is often not effective.

To give improved lower bounds, the problem is further divided into two cases, one for the case the variable is moved upwards and one for the other direction.

upwards and one for the other direction. Let  $f : \mathbf{B}^n \to \mathbf{B}^m$  be a Boolean function represented by a BDD and assume that the initial variable ordering is  $(x_1, x_2, \ldots, x_n)$ . From iteratively applying Theorem 1 and making use of the results from [1] and [6] we get the following two corollaries:

**Corollary 1** Let  $1 \le i < j \le n$ . If level *i* is moved down to level *j*, resulting in the variable ordering

$$(x_1, x_2, \ldots, x_{i-1}, x_{i+1}, \ldots, x_j, x_i, x_{j+1}, \ldots, x_n),$$

for the size of the resulting BDD |G'| holds

$$|G'| \ge \operatorname{label}(A) + 1 + \frac{1}{2} \cdot \operatorname{label}(B) + \operatorname{label}(C)$$

and

$$|G'| \ge \text{label}(A) + \text{label}(x_i) + \text{label}(C)$$

with  $A := \{x_1, \ldots, x_{i-1}\}, B := \{x_{i+1}, \ldots, x_j\}$ , and  $C := \{x_{j+1}, \ldots, x_n\}.$ 

**Corollary 2** Let  $1 \leq j < i \leq n$ . If level *i* is moved up to level *j*, resulting in the variable ordering

$$(x_1, x_2, \ldots, x_{j-1}, x_i, x_j, \ldots, x_{i-1}, x_{i+1}, \ldots, x_n),$$

for the size |G'| of the resulting BDD holds

$$|G'| \ge \operatorname{label}(A) + (i-j) + \frac{1}{2^{i-j}} \cdot \operatorname{label}(x_i) + \operatorname{label}(C),$$

with  $A := \{x_1, \ldots, x_{j-1}\}, B := \{x_j, \ldots, x_{i-1}\}, \text{ and } C := \{x_{i+1}, \ldots, x_n\}.$ 

Furthermore, levels which do not interact with  $x_i$  can be added to both lower bounds.

 $<sup>^1{\</sup>rm For}$  two variables the interaction matrix gives information whether an output of the BDD depends on both variables.

Table 1: Lower bound sifting

				sift in		lb-sift	ting	difference (%)	
circuit	$_{ m in}$	initial	final	exch.	time	exch.	time	exch.	$\operatorname{time}$
bigtest	328	95224	93066	204803	332.4	101925	138.9	-50.2	-58.2
$c1\bar{3}55$	41	39648	30103	3108	17.1	1580	12.8	-49.2	-24.9
c1908	33	23158	7582	2015	6.3	1105	4.7	-45.2	-25.3
c2670	233	222404	16106	71002	216.0	26762	104.8	-62.3	-51.5
c499	41	39649	30459	3117	15.5	1579	11.2	-49.3	-28.0
c5315	178	4737	2377	52607	2.2	27875	1.2	-47.0	-46.4
c7552	207	331495	13934	79092	75.2	44911	53.4	-43.2	-28.9
c880	60	8116	4384	6462	1.6	3328	0.8	-48.5	-48.8
dalu	75	12355	1216	10520	2.4	8210	1.9	-22.0	-22.4
$\operatorname{des}$	256	9413	3065	103031	1.5	73370	1.2	-28.8	-19.8
i10	257	326484	97256	121280	447.3	63998	286.5	-47.2	-36.0
i2	201	334	204	69401	3.8	29133	1.8	-58.0	-51.5
i4	192	420	246	68985	1.1	13371	0.4	-80.6	-60.6
i8	133	3869	1665	28049	1.0	18868	0.8	-32.7	-20.1
pair	173	14432	6068	56820	3.7	33710	3.0	-40.7	-18.7
$\operatorname{rot}$	135	8147	6200	33690	2.4	15895	1.5	-52.8	-35.7
s13207.1	700	10432	3152	797041	7.6	426990	7.2	-46.4	-4.9
s1423	91	4117	1837	15425	1.2	9621	0.9	-37.6	-29.7
s15850.1	611	42944	14660	643009	39.3	350288	28.3	-45.5	-28.0
s38417	1664	608542	560835	5011193	3336.4	1442982	703.2	-71.2	-78.9
s38584.1	1464	62901	16245	3751859	52.6	2453578	41.5	-34.6	-21.0
s5378	199	5256	2391	71740	1.4	38904	1.2	-45.8	-18.5
s9234.1	247	25135	4267	112799	8.3	60300	6.2	-46.5	-25.6
$\operatorname{sum}$		1899212	917318	11317048	4576.3	5248283	1413.4	-53.6	-69.1

 $\begin{array}{l} \mbox{SiftingDown}(i:\mbox{level to sift}) \\ lb = \mbox{compute_lower_bound}(); \\ best = \mbox{size_of_BDD}(); \\ \mbox{while } (i < n \mbox{ and } lb \leq best) \\ \mbox{level_exchange}(i, i + 1); \\ lb = \mbox{compute_lower_bound}(); \\ \mbox{if } (\mbox{size_of_BDD}() < best) \mbox{ best } = \mbox{size_of_BDD}(); \\ i = i + 1; \end{array}$ 

Figure 2: Sketch for the sifting down-algorithm

# 4 Lower Bound Sifting

During sifting, moving a variable  $x_i$  can be aborted if it cannot improve the best already known BDD size. In other words, if it is known that the BDD size cannot be improved by further shifting the variable, this can be stopped. To decide whether an improvement is still possible, the lower bounds given in the previous section are used. For the case of moving a variable  $x_i$  down, this yields

$$\begin{split} \mathrm{lb}(x_i) &= \min_{j=1,\dots,i-1} |G'_j| \\ &\geq \mathrm{label}(A) + \mathrm{max}(\mathrm{label}(x_i), 1 + \tfrac{1}{2} \cdot \mathrm{label}(B)), \end{split}$$

with  $A := \{x_1, \ldots, x_{i-1}\}, B := \{x_{i+1}, \ldots, x_n\}$  and  $G'_j$  denoting the BDD after moving variable  $x_i$  to position j. If this lower bound is larger than the best previously found BDD size, a movement in that direction cannot lead to a better position for the variable. Analogously the lower bound is computed for moving a variable down. The computation of the lower bound can be done locally, and by this is much more efficient than the approach in [6], where a traversal of the whole BDD is needed. A sketch of the algorithm for sifting down is given in Figure 2. In the following this technique is denoted as *lower bound sifting* (lb-sifting). Notice that the quality (measured in number of nodes) of lb-sifting is the same as for "classical" sifting, but due to the lower bounds the computation is speeded up, since much less exchanges have to be performed.

The runtime of dynamic variable reordering can be further improved, if the lower bounds are relaxed: It is unlikely that during sifting half of the nodes are vanishing in each step. For this, we suggest to not only consider the bound  $\frac{1}{2}$ , but the straightforward generalization to  $\frac{1}{b}$  ( $b \ge 2$ ). Obviously, this is not a "real" lower bound, since it overestimates, but by this extension sifting only consideres parts of the search space where large improvements are possible. (In typical BDD applications, like verification, a small improvement in number of nodes is not of much interest.) The  $\frac{1}{b}$  bound is used for moving variables up and down, i.e. we do not distingiush between these cases.

It turned out in our experiments (see below) that using the  $\frac{1}{b}$  bound is a good choice for  $2 \le b \le 10$ . Within this range, the relaxed lower bound often tremendously speeds up the sifting algorithm without increasing the resulting BDD sizes too much. This heuristic approach of using lower bounds allows the user to trade off runtime versus quality.

#### 5 Experimental Results

In this section we describe experimental results that have been carried out on a SUN Ultra 1 with 256 MBytes. All times are given in CPU seconds. Our algorithm has been integrated in the CUDD package [16].

In a first series of experiments, we compare the original sifting algorithm [14] with the lower bound improvement. Results are given in Table 1. In column *circuit* the name of the benchmark circuit is given and column *in* refers to the number of inputs of the circuit. We build the BDD using an initial ordering (given in column *initial*). The BDD size after application of sifting is shown in column *final*. (Obviously, Ib-sifting results in the same sizes.) For both sifting methods we give the number of exchanges that are carried out during sifting and the runtime needed. The improvements are given in the last two columns. Ib-sifting needs on average 53.5% less exchanges resulting in a reduction of runtime by 69.1%. Especially for bigger examples large reductions are achieved (see e.g. s38417). Compared to the trivial lower bound (as it is implemented in [16]) improvements of up to 70% can be overserved using Ib-sifting.

In a second series of experiments, we study the influence of relaxing the lower bound. For increasing values of b, we analyze the influence of the relaxed lower bound on final node count and on runtime. (The bound is used in both directions, i.e. for moving a variable up and down.) The results for various values of b in comparison to sifting and

Table 2: Relaxed lower bound sifting

Table 2. Itelaxed lower bound shtting												
	$\operatorname{nodes}$		deviation/circuit			$\operatorname{time}$		deviation/circuit				
$\mathrm{met}\mathrm{hod}$	$\operatorname{sum}$	$\mathbf{rel}$	av.	max	$\min$	$\operatorname{sum}$	$\operatorname{rel}$	av.	$\max$	$\min$		
sifting	929078	0.0	0.0	0.0	0.0	4582.7	0.0	0.0	0.0	0.0		
lb-sifting	929081	0.0	0.0	0.0	0.0	1418.1	-69.1	-29.9	-4.9	-81.9		
b=2	931321	0.2	1.4	38.9	-7.4	1534.6	-66.5	-21.8	4.5	-76.9		
b = 3	931938	0.3	3.0	40.7	-7.4	1277.5	-72.1	-30.5	-3.0	-84.3		
b = 4	932552	0.4	3.8	67.1	-7.4	1073.0	-76.6	-35.6	3.0	-84.3		
b = 6	934314	0.6	5.1	67.1	-7.1	841.5	-81.6	-42.4	-9.1	-88.3		
b = 8	935381	0.7	5.6	67.1	-7.1	707.6	-84.6	-47.3	3.0	-90.6		
b = 10	934869	0.6	6.3	67.1	-7.1	628.7	-86.3	-49.8	-3.0	-92.0		
b = 12	939009	1.1	9.5	115.0	-7.1	565.8	-87.7	-53.0	-15.0	-93.0		
b = 14	940209	1.2	12.1	115.1	-7.1	509.4	-88.9	-55.4	-15.0	-94.0		
b = 16	939689	1.1	12.1	115.1	-7.1	467.8	-89.8	-57.5	-22.1	-94.7		
b = 18	941073	1.3	13.7	115.1	-7.1	440.1	-90.4	-57.6	-7.7	-95.1		
b = 20	973241	4.8	19.0	239.1	-7.1	409.6	-91.1	-58.8	-7.7	-95.6		
b = 22	973553	4.8	19.8	239.1	-7.1	391.7	-91.5	-60.7	-15.2	-95.8		
b = 24	974849	4.9	20.6	239.1	-7.1	374.8	-91.8	-62.2	-15.2	-96.0		



Figure 3: CPU time in seconds (left) and size in 1000 nodes (right) for relaxing bound

lb-sifting are given in Table 2. Columns 2-6 and 7-11 give informations about the behavior of the algorithms for resulting BDD size and runtime, respectively. For all experiments 43 circuits have been used and we computed the average (column av.), maximal (column max) and minimal (column min) deviation per circuit given in %. rel gives the relative improvement compared to "classical" sifting. To make this data easier readable, we show the overall behavior in a diagram: The runtimes and the final BDD sizes in 1000 nodes for various values of b are given in Figure 3. It can be seen that for small values of b (e.g. b = 10), the increase in number of nodes is very small, i.e. less than 1%, but the gain in runtime is large, i.e. more then a factor of 7 over "classical" sifting. Our results show that relaxed lb-sifting is a very robust method to trade off runtime versus BDD size.

#### 6 Conclusions

In this paper we studied the use of lower bounds during dynamic BDD minimization. It turned out that it is possible to speed up the original algorithm significantly without increasing the final size of the BDD. We proved tight bounds that can be computed efficiently during sifting. If a small increase is allowed, i.e. the lower bounds are relaxed, the algorithm can be further speeded up.

Experimental results have shown that an improvement of runtime by nearly 70% becomes possible without making the results worse. Using relaxed lb-sifting the runtime could be further improved by more than a factor of two while increasing the BDD size by less than 1% on average.

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