

Approximability and Nonapproximability by Binary Decision Diagrams (Extended Abstract)

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Abstract. Many BDD (binary decision diagram) models are motivated by CAD applications and have led to complexity theoretical problems and results. Motivated by applications in genetic programming Krause, Savický, and Wegener (1999) have shown that for the inner product function IP_n and the direct storage access function DSA_n all functions which approximate them on considerably more than half of the inputs need exponential π -OBDD size for most variable orderings π . In this paper, the results of Krause, Savický, and Wegener are generalized to more general BDD models like k-IBDDs and BP1s (read-once branching programs). Furthermore, the size of OBDDs for functions which approximate a function f_n is compared with the size of randomized OBDDs with two sided error for f_n. An exponential gap is presented.

Keywords: Computational complexity, binary decision diagrams, approximations.

1 Introduction and Definitions

Branching programs (BPs) or Binary Decision Diagrams (BDDs) are a well established representation type or computation model for Boolean functions $f \in B_n$, i.e., $f : \{0, 1\}^n \to \{0, 1\}$.

Definition 1. A branching program (BP) or binary decision diagram (BDD) on the variable set $X_n = \{x_1, \ldots, x_n\}$ is a directed acyclic graph with one source and two sinks labelled by the constants 0 or 1, resp. Each non-sink node (or inner node) is labelled by a Boolean variable and has two outgoing edges one labelled by 0 and the other by 1. At each node v a Boolean function $f_v : \{0,1\}^n \to \{0,1\}$ is represented. For the evaluation of $f_v(a)$ start at v. At x_i -nodes, the outgoing a_i -edge is chosen and $f_v(a)$ equals the label of the sink which is finally reached on the computation path.

The size of a branching program G is equal to the number of its nodes and is denoted by |G|. BP(f) denotes the size of the smallest BP for a function f. The

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depth of a branching program is the maximum length of a path from the source to one of the sinks.

Probabilistic methods have turned out to be useful in almost all areas of computer science. Ablayev and Karpinski (1996) have introduced randomized BPs defined analogously to probabilistic circuits. In the following we consider an alternative approach.

Definition 2. A randomized BP (or BDD) G on the variable set $X_n = \{x_1, \ldots, x_n\}$ is a directed acylic graph with decision nodes for Boolean variables and randomized nodes. A randomized node is an unlabelled node with two outgoing edges. The random computation path for a is defined as follows. At decision nodes labelled by x_i , the outgoing a_i -edge is chosen. At randomized nodes, each outgoing edge is chosen independently from all other random decisions with probability 1/2. The acceptance probability $acc_G(a)$ or Prob(G(a) = 1) of G on a is the probability that the random computation path reaches a 1-sink. The rejection probability $rej_G(a)$ equals Prob(G(a) = 0) = 1 - Prob(G(a) = 1).

G represents $f \in B_n$ with two-sided ϵ -bounded error, $0 \leq \epsilon < 1/2$, if $Prob(G(a) \neq f(a)) \leq \epsilon$ for all inputs a.

In order to develop and strengthen lower bound techniques and for applications restricted computation models are considered.

Definition 3. i) A branching program is called read k times (BPk) if each variable is tested on each path at most k times.

ii) A BP is called oblivious if the node set can be partitioned into levels such that edges lead from lower to higher levels and all inner nodes of one level are labelled by the same variable.

In his seminal paper Bryant (1986) has introduced ordered binary decision diagrams (OBDDs) which are up to now the most popular representation not only for formal circuit verification but also in genetic programming.

Definition 4. A permutation π on $\{1, \ldots, n\}$ describes the variable ordering $x_{\pi(1)}, \ldots, x_{\pi(n)}$. A π -OBDD for a variable ordering π is a BP where the sequence of tests on a path is restricted by the variable ordering π , i.e., if an edge leads from an x_i -node to an x_j -node, the condition $\pi(i) < \pi(j)$ has to be fulfilled. An OBDD is a π -OBDD for some variable ordering π .

Unfortunately, several important and also quite simple functions have exponential OBDD size. Therefore, more general representations with good algorithmic behaviour are necessary. Jain, Abadir, Bitner, Fussell, and Abraham (1994) have introduced k-IBDDs.

Definition 5. A k-IBDD is a branching program which can be partitioned into k layers such that the *i*th layer is an OBDD (with possible many sources) respecting the ordering π_i and such that the edges leaving the *i*th layer reach only nodes of the layers j > i and the sinks.

One main problem in learning theory is the search for a small size representation of a function $f \in B_n$ which coincides with the partial function $g \in B_n$ defined by the set of labelled training examples $(a_1, g(a_1)), \ldots, (a_m, g(a_m))$ where $a_i \in \{0,1\}^n$ and $m = n^{O(1)}$. The idea to look for a small size representation is based on the well-known Occam's razor principle. See Krause, Savický, and Wegener (1999) for a discussion of this background. They also have described how such a small representation can be obtained by a genetic programming approach. In particular, they have underlined the importance of the choice of a good variable ordering. Simple functions like the direct storage access function DSA_n and the inner product function IP_n have small OBDD size but almost all variable orderings lead to exponential size representations for the considered functions and even all reasonably good approximations. In learning theory we cannot expect to learn the unknown function exactly. It is good enough to obtain a representation of a function which almost looks like the unknown function. This motivates the investigation of a BDD theory for the approximate representation of Boolean functions.

Definition 6. i) A function $g \in B_n$ is a c-approximation of $f \in B_n$ if $Pr(f(\tilde{x}) = g(\tilde{x})) \ge c$ for a random input \tilde{x} .

ii) A function $g \in B_n$ is a strong c-approximation of $f \in B_n$ if, for $a \in \{0, 1\}$, $Pr(g(\tilde{x}) = a) \ge c$ for a random input $\tilde{x} \in f^{-1}(a)$.

In applications, strong approximations are necessary. However, our lower bounds even hold for c-approximations. Since one of the two constant functions 0 and 1 is always a 1/2-approximation we consider $(1/2 + \epsilon_n)$ -approximations. Using methods from information theory and communication complexity Krause, Savický, and Wegener (1999) have proved that besides DSA_n the inner product function IP_n has only $(1/2 + \epsilon_n)$ -approximations of exponential π -OBDD size for almost all variable orderings π . Futhermore, they have introduced the permuted inner product function PIP_n for which all $(1/2 + \epsilon_n)$ -approximations have exponential OBDD size. The shifted inner product function SIP_n is related to IP_n and has appeared in several disguises in the literature on BDDs (see, e.g., Krause (1992)).

- **Definition 7.** *i)* For even n, $IP_n(x_1, \ldots, x_n) = x_1x_2 \oplus x_3x_4 \oplus \ldots \oplus x_{n-1}x_n$ is the inner product function.
- ii) For even n, let $\operatorname{IP}_n^{\pi}(x_1, \ldots, x_n) = \operatorname{IP}_n(x_{\pi(1)}, \ldots, x_{\pi(n)})$. For $l = \lceil \log(n!) \rceil$, $\operatorname{PIP}_n(a_0, \ldots, a_{l-1}, x_1, \ldots, x_n) = \operatorname{IP}_{\pi_a}(x_1, \ldots, x_n)$ is the permuted inner product function, where π_a is the permutation on $\{1, \ldots, n\}$ coded by the variables a_0, \ldots, a_{l-1} .
- iii) Let $n = 2^l$. The function SIP_n is defined on 2n+l Boolean variables $a_{l-1}, \ldots, a_0, x_{n-1}, \ldots, x_0, y_{n-1}, \ldots, y_0$. Let

$$\mathrm{SIP}_n^i(x,y) = \bigoplus_{0 \le j \le n-1} x_j y_{(j+i) \mod n},$$

then

$$\operatorname{SIP}_n(a, x, y) = \bigvee_{0 \le i \le n-1} (|a| = i) \wedge \operatorname{SIP}_n^i(x, y).$$

Now, we give an overview on the rest of the paper. In Section 2, we present some tools from communication complexity and show how distributional communication complexity can be used to prove lower bounds on the BDD size for approximations of Boolean functions. The relation between BDDs for approximations and randomized BDDs with two-sided bounded error is investigated in Section 3. A function f_n is presented which cannot well be represented by oblivious randomized BDDs of linear length but for which there exists a function g_n with linear OBDD size, where g_n approximates f_n on almost all inputs. In Section 4, we prove that the permuted inner product function PIP_n has no small and good approximation by k-IBDDs or BP1s.

2 Communication Complexity and Approximations

We start with some notions from communication complexity needed later on. For a thorough introduction into communication complexity, we refer to the monographs of Hromkovič (1997) and Kushilevitz and Nisan (1997). The communication complexity of two-party protocols has been introduced by Yao (1979). It has turned out to be the strongest tool for proving lower bounds on the size of oblivious BDDs. The basic model is the following one. Alice and Bob want to cooperate for the evaluation of a Boolean function $f(x, y) \in B_{m+n}$, i.e., $f: \{0,1\}^m \times \{0,1\}^n \to \{0,1\}$. Alice knows the first part $X = (x_1, \dots, x_m)$ and Bob the second part $Y = (y_1, \ldots, y_n)$ of the input. Before Alice and Bob obtain their partial inputs, they reach an agreement on a communication protocol which decides who starts the communication and how the communication is organized. If Alice starts the communication, she sends the first message $m_1(x)$ depending on her partial input x, Bob answers with a message $m_2(y, m_1(x))$ depending on his partial input y and Alice's message. Then Alice anwers and so on until one player knows f(x, y). The cost of a protocol P for f on input (x, y) is the total length of all messages sent by Alice and Bob. The cost of the protocol P is the cost for a worst case input. The deterministic communication complexity C(f) of f is the minimum cost of P, over all protocols P that compute f. Many generalizations are possible, e.g., nondeterministic protocols and randomized protocols. Furthermore, the number of communication rounds may be restricted.

Distributional communication complexity provides lower bounds for randomized protocols that are allowed two-sided error.

Definition 8. Let μ be a probability distribution on $X \times Y$. The (μ, ϵ) -distributional communication complexity of f, $D^{\mu}_{\epsilon}(f)$, is the cost of the best deterministic protocol that gives the correct answer for f on at least a $(1 - \epsilon)$ -fraction of all inputs in $X \times Y$, weighted by μ .

For ease of description we rename the variables such that $IP_n(x, y) = x_1y_1 \oplus \cdots \oplus x_{n/2}y_{n/2}$. We give the x-variables to Alice and the y-variables to Bob. The distributional communication complexity of the inner product function has been studied by Chor and Goldreich (1988). They have proved the following fact which has turned out to be very useful for our investigations.

Fact 1: $D_{1/2-\epsilon}^{uniform}(\mathrm{IP}_n) \ge n/4 - \log(1/\epsilon).$

The following proposition shows how lower bounds on the distributional communcation complexity of a function f lead to lower bounds on the size of BDDs for approximations of f.

Proposition 1. Let $f \in B_{m+n}$ and g be a c-approximation of f and G be a k-IBDD representing g with respect to the variable orderings π_1, \ldots, π_k where for all $\pi_i, 1 \leq i \leq k$, it holds that the x-variables are tested before the y-variables. Then it holds that

$$(2k-1)\left[\log|G|\right] \ge D_{1-c}^{uniform}(f).$$

Proof. This result follows by the fact that $D_{1-c}^{uniform}(f) \leq C(g)$. The upper bound $(2k-1)\lceil \log |G| \rceil$ for C(g) is well-known (Wegener (2000)).

Communication complexity cannot be used directly for lower bound proofs on the size of BPks which are not oblivious. The only remaining known lower bound technique is the method of generalized rectangles based on so-called (k, a)rectangles (a name introduced by Jukna (1995)) which first has been described by Borodin, Razborov, and Smolensky (1993) and Okol'nishnikova (1993). Using results from randomized communication complexity Sauerhoff (1998) has generalized this technique to the randomized case to prove lower bounds on the size of randomized BPks. Finally, Thathachar (1998) has applied Sauerhoff's technique to separate the read-k-times branching program hierarchy.

Definition 9. A Boolean function $g \in B_n$ is called a (k, a)-rectangle if it can be represented as a conjunction of functions g_i , $1 \le i \le ka$, such that g_i essentially depends only on variables from X(i) where $|X(i)| \le \lceil n/a \rceil$ and each variable x_j belongs to at most k of the X(i)-sets.

We identify a (k, a)-rectangle R, which is defined as Boolean function, with the set $R^{-1}(1)$. Such a rectangle R is called f-monochromatic if $R \subseteq f^{-1}(0)$ or $R \subseteq f^{-1}(1)$.

Definition 10. A function $f \in B_n$ is called an (s, k, a)-step function if $\{0, 1\}^n$ can be partitioned into $(2s)^{ka}$ f-monochromatic (k, a)-rectangles.

Functions $f \in B_n$ whose BPk size equals s are (s, k, a)-step functions. Applying a method due to Yao (1983), Sauerhoff (1998) has proved that functions with small randomized BPk size can be approximated with small error by (s, k, a)-step functions.

Theorem 1. Let μ be a probability measure on $\{0,1\}^n$. If $f \in B_n$ can be represented by a randomized BPk of size s which has two-sided ϵ -bounded error, there exists an (s, k, a)-step function g such that g μ -approximates f with ϵ -bounded error, i.e., the set of all a where $f(a) \neq g(a)$ has a μ -measure bounded above by ϵ .

Therefore, the randomized BPk size of a function f is an upper bound for the BPk size of an $(1 - \epsilon)$ -approximation of f if μ is the uniform distribution. How far can the best upper bound obtained by this method be away from the BPk size of an $(1 - \epsilon)$ -approximation of f? In the next section we answer this question for oblivious BDDs of linear length.

3 Randomized BDDs and Approximations by BDDs

Theorem 2. Let $n = 2^l$. There exists a function f_n on 3n + l Boolean variables which needs exponential size for oblivious randomized BDDs of linear length with two-sided ϵ -bounded error, $0 \leq \epsilon < 1/2$, but has a strong $(1 - 1/2^n)$ approximation g_n which can be represented by π -OBDDs of size 4n + 2l + 1for arbitrary π .

Proof. The function $f_n : \{0,1\}^{3n+l} \to \{0,1\}$ is defined on the variable set $A_n \cup X_n \cup Y_n \cup Z_n$ where $A_n = \{a_0, \ldots, a_{l-1}\}, X_n = \{x_0, \ldots, x_{n-1}\}, Y_n = \{y_0, \ldots, y_{n-1}\}, \text{ and } Z_n = \{z_0, \ldots, z_{n-1}\}$. Let $h_n(a, x, y) := \text{SIP}_n(a, x, y)$. Then the function f_n is given as

$$f_n(a, x, y, z) := \begin{cases} a_0 \oplus \dots \oplus a_{l-1} \oplus x_0 \oplus \dots \oplus x_{n-1} \oplus y_0 \oplus \dots \oplus y_{n-1}, & \text{if } z \neq \mathbf{0}, \\ h_n(a, x, y), & \text{if } z = \mathbf{0}. \end{cases}$$

For the lower bound it is sufficient to show that the subfunction h_n of f_n cannot well be represented by oblivious randomized BDDs of linear length with twosided ϵ -bounded error. It is known that deterministic oblivious BDDs of linear length representing the function SIP_n have exponential size (Krause (1992)). Let $s = (s_1, \ldots, s_l)$ be a sequence of variables from X_n and Y_n such that no variable appears more than k times. There are two sets $X'_n \subseteq X_n$ and $Y'_n \subseteq Y_n$ such that $|X'_n| \ge |X_n|/2^{2k-1}$, $|Y'_n| \ge |Y_n|/2^{2k-1}$, and the number of layers in s with respect to X'_n and Y'_n is bounded by 2k+1 (Alon and Maass (1988)). There are at least $(n/2^{2k-1})^2$ pairs (x_{j_1}, y_{j_2}) with $x_{j_1} \in X'_n$ and $y_{j_2} \in Y'_n$. By the pigeonhole principle there exists an index $i \in \{0, \ldots, n-1\}$ such that there are at least $n/2^{4k-2}$ pairs $(x_j, y_{(j+i) \mod n})$. The x-variables of these pairs are contained in the set X''_n and the y-variables in Y''_n respectively. We fix the a-variables in such a way that |a| = i and we set all variables of $X_n \cup Y_n$ which are not in $X''_n \cup Y''_n$ to 0.

Using the fact that the inner product function IP_n has a communication complexity of $\Omega(n)$ for randomized communication protocols with two-sided ϵ bounded error (where Alice holds the *x*- and Bob the *y*-variables and the protocols may use an arbitrary number of rounds) (see e.g. Babai, Frankl, and Simon (1986)) we obtain an exponential lower bound of $\Omega(2^{n/2^{4k-2}})$ for the size of oblivious randomized BDDs of linear length with two-sided ϵ -bounded error, $0 \leq \epsilon < 1/2$, representing the function f_n .

Now, we prove the upper bound. The function $g_n(a, x, y, z) = a_0 \oplus \cdots \oplus a_{l-1} \oplus x_0 \oplus \cdots \oplus x_{n-1} \oplus y_0 \oplus \cdots \oplus y_{n-1}$ is obviously a strong $(1 - 1/2^n)$ -approximation of f_n . The π -OBDD size of g_n is 4n + 2l + 1 for arbitrary π .

The result shows that for proving lower bounds on the size of oblivious randomized BDDs for a function f_n it is sufficient to find a hard subfunction which is difficult to represent. Therefore, it is also possible to apply reducibility concepts like the so-called rectangular reductions (see, e.g., Kushilevitz and Nisan (1997)). The situation is different for the representations of strong approximations for f_n . If the error probability is close to 1/2, we have to prove that all subfunctions of f_n are hard to represent.

4 The Permuted Inner Product Function

The permuted inner product function PIP_n defined on $l := \lceil \log(n!) \rceil$ selection variables and n data variables x_1, \ldots, x_n has been introduced by Krause, Savický, and Wegener (1999) showing that there exists a function which cannot be approximated well by OBDDs. Now, we prove that PIP_n cannot be approximated well by k-IBDDs and BP1s either. The l selection variables describe a permutation π on $\{1, \ldots, n\}$, more precisely, for each permutation the number of codewords is one or two. The function PIP_n realizes IP_n on the data variables which are permuted according to π .

For $\operatorname{IP}_n^{\pi}$ we call $x_{\pi(2i-1)}$ the partner of $x_{\pi(2i)}$, $1 \leq i \leq n/2$, and vice versa. A variable is called singleton with respect to a variable ordering π' , or a π' -singleton for short if it is in the first half of the variables according to π' but its partner is in the second half. Similar we define singletons according to a variable set $A \subseteq \{x_1, \ldots, x_n\}$, or a A-singletons for short if a variable is in A and its partner is not.

First, we determine the number of different functions IP_n . We consider the following experiment. In the beginning x_1, \ldots, x_n are free. Then we choose the free variable with the smallest index and choose another arbitrary free variable as its partner. The chosen variables are no longer free. The process is repeated n/2 times. Hence, the number of different functions IP_n^{π} equals

$$d(n) = \prod_{1 \le i \le n/2} (2i - 1) = 2^{\Theta(n \log n)}.$$

Fact 2 (Krause, Savický, and Wegener (1999)): With probability at least $1-e^{-4\delta^2 n}$ a random variable ordering π' leads to at least $(1-\delta)n/8 \pi'$ -singletons for $\operatorname{IP}_n^{\pi}$.

Lemma 1. Let k be a constant and let π_1, \ldots, π_k be independent random variable orderings. With probability at least $1-e^{-\Omega(n)}$ there is a set $I \subseteq \{x_{\pi(2i-1)} | 1 \le i \le n/2\}$ of at least $((1-\delta)/8)^k n = \Omega(n)$ elements with the following property. It is possible to partition each variable ordering π_j , $1 \le j \le k$, into a top part and a bottom part such that for each $x_{\pi(2i-1)} \in I$ the top part contains exactly one of the variables $x_{\pi(2i-1)}$ and $x_{\pi(2i)}$.

Proof. Let $I_0 = \{x_{\pi(2i-1)} | 1 \le i \le n/2\}$. The set I_1 is constructed in the way described by Krause, Savický, and Wegener (1999). The set I_j is constructed

in the same way based on I_{j-1} and the variable ordering π_j restricted to the variables of I_{j-1} and their partners. Hence, $|I_j| \ge ((1 - \delta)/8)|I_{j-1}| \ge ((1 - \delta)/8)^j n$ proving the claim on the size of $I = I_k$. The error probability is bounded above by $e^{-4\delta^2|I_{j-1}|}$ implying that the total error probability is bounded by $ke^{-4\delta^2|I|} = e^{-\Omega(n)}$.

Now, we are ready to prove the following theorem.

Theorem 3. Let k be a constant. Each function which is a c-approximation of PIP_n with $c \geq 1/2 + ke^{-4\delta^2((1-\delta)/8)^k n} + e^{-n^{1-\epsilon'}}$, $0 < \epsilon' < 1$, has a k-IBDD size which is bounded below by $2^{\Omega(n)}$.

Proof. Let π_1, \ldots, π_k be arbitrary variable orderings and G be a k-IBDD according to π_1, \ldots, π_k representing a c-approximation g_n of PIP_n. By the definition of PIP_n we have $N := 2^{\lceil \log(n!) \rceil}$ codewords. Therefore, some π have a probability of 1/N to be chosen and others have a probability of $2/N \leq 2/(n!)$. Hence, the error probability of Lemma 1 is at most doubled and we get the result that for at least $(1 - 2ke^{-4\delta^2((1-\delta)/8)^k n})2^{\Theta(n \log n)}$ subfunctions IP_n^{π} of PIP_n obtained by replacing the selection variables with constants there are at least $((1 - \delta)/8)^k n$ variables in a subset I such that π_1, \ldots, π_k can be partitioned into a top part and a bottom part such that for each $x_{\pi(2i-1)} \in I$ the top part contains exactly one of the variables, we obtain the function $IP_{((1-\delta)/8)^k n}^{\pi}$ or its negation as subfunction. Let $\gamma := 2ke^{-4\delta^2((1-\delta)/8)^k n}$.

By averaging and using Proposition 1

$$|G| \ge 2^{1/(2k-1)(D_{1/(1-\gamma)(1-c)}^{uniform}(\operatorname{IP}_{((1-\delta)/8)^{k_n}}))}.$$

Using Fact 1 it follows that

$$|G| > 2^{1/(2k-1)(((1-\delta)/8)^k n/2 - \log((1/2 - (1/(1-\gamma))(1-c))^{-1}))}$$

It holds that

$$-\log((1/2 - (1/(1-\gamma))(1-c))^{-1}) = \log((-1-\gamma+2c)/(2(1-\gamma)))$$

$$\geq \log(-1-\gamma+2c) - 1.$$

Since $c \ge 1/2 + (1/2)\gamma + e^{-n^{1-\epsilon'}}$ it follows that

$$|G| \ge 2^{1/(2k-1)(((1-\delta)/8)^k n/2 - (\log e)n^{1-\epsilon'})} = 2^{\Omega(n)}.$$

Now, we consider *c*-approximations of PIP_n represented by BP1s. First, we consider the subproblem to represent the function IP_n by restricted BP1s. Often it turns out to be easier to prove lower bounds on the size of BP1s if they have the following special property.

Definition 11. A BP1 on the variables x_1, \ldots, x_n is called regular if for each node v the same set of variables is tested on all paths from the source to v. Furthermore, it is required that on each path from the source to the sinks all n variables are tested.

It is not difficult to see that an arbitrary BP1 G can be converted into a regular BP1 G' with size $|G'| \leq (n+1)|G|$ by inserting dummy tests.

Since the function IP_n can be represented by OBDDs of linear size, it is not possible to apply directly the method of generalized rectangles to prove exponential lower bounds. Our aim is to adapt the known proof technique to prove a lower bound on the size of restricted BP1s for IP_n . Therefore, we combine an adapted version of the rectangle method with results from distributional communication complexity.

The following theorem is well-known (see e.g. Okol'nishnikova (1993)).

Theorem 4. If a function $f \in B_n$ can be represented by a BP1 G of size s, $\{0,1\}^n$ can be partitioned into 2s f-monochromatic rectangles.

The proof idea of Theorem 4 is to decompose G into 2-dimensional rectangles. Therefore, we define a cut through G consisting of all nodes v such that on each path from the source of G to v exactly n/2 variables have been tested. By our assumption on G on all paths to v the same set A(v) of variables has been tested before reaching v. In this way paths using the same set of variables reaching the same node v on the cut are "bundled" together. A 2-dimensional rectangle hence represents the computation paths running through a sequence (v_0, v_1, v_2) of nodes, where v_0 is the source of G, v_2 is one of the sinks, and v_1 is a node on the cut of G. From this construction it follows immediately that properties of the paths to the nodes on the cut of G correspond to properties of the 2-dimensional rectangles.

A 2-dimensional rectangle R corresponds to a balanced partition of the variables into two sets X_1 and X_2 . R is a 2-dimensional rectangle with at least $(1 - \delta)n/8$ singletons if there are at least $(1 - \delta)n/8$ variables in X_1 whose partners belong to X_2 .

Proposition 2. Let G be a BP1 for IP_n of size s. If for all nodes v of G which can be reached after testing exactly n/2 variables it holds that there exist at least $(1-\delta)n/8 A(v)$ -singletons, $\{0,1\}^n$ can be partitioned into 2s IP_n-monochromatic 2-dimensional rectangles with at least $(1-\delta)n/8$ singletons.

The discrepancy technique (see, e.g., Kushilevitz and Nisan (1997)) is known as a method for proving lower bounds for the distributional communication complexity D_{ϵ}^{μ} by giving upper bounds on the size of rectangles that are "almost" monochromatic.

Definition 12. Let $f : X \times Y \to \{0,1\}$ be a Boolean function, R be any rectangle, and μ be a probability distribution on $X \times Y$. Denote

$$\begin{split} Disc_{\mu}(R, \ f) &= \\ & \left| Pr_{\mu}(f(x,y) = 0 \ and \ (x,y) \in R) - Pr_{\mu}(f(x,y) = 1 \ and \ (x,y) \in R) \right|. \end{split}$$

The discrepancy of f according to μ is $Disc_{\mu}(f) = \max_{R} Disc_{\mu}(R, f)$ where the maximum is taken over all rectangles R.

Sauerhoff (1998) has generalized this method by introducing the rectangle balance property. Afterwards, he has shown how this property can be used for proving lower bounds on the size of randomized BPks. Here, we present the method for proving lower bounds on the size of BP1s for $(1 - \epsilon)$ -approximations of functions $f \in B_n$.

Definition 13. Let $f \in B_n$, μ be a probability measure on $\{0,1\}^n$, and $\gamma(n)$ be a real-valued function. The function f has the rectangle balance property with respect to $(\mu, \gamma(n))$ if

$$\mu(R \cap f^{-1}(0)) \ge \mu(R \cap f^{-1}(1)) - \gamma(n)$$

holds for each 2-dimensional rectangle R.

Theorem 5 (Sauerhoff(1998)). If $f \in B_n$ has the rectangle balance property with respect to $(\mu, \gamma(n))$, the size of each BP1 representing a $(1 - \epsilon)$ approximation of f is bounded below by

$$1/2[(\mu(f^{-1}(1)) - \epsilon)/\gamma(n)].$$

Now, we adapt this method to our case. In the rest of the section we set μ to the uniform distribution.

Corollary 1. Let G be a regular BP1 representing a $(1 - \epsilon)$ -approximation of IP_n with the property that for all nodes v of G which can be reached after testing exactly n/2 variables it holds that there exist at least $(1 - \delta)n/8 A(v)$ -singletons according to IP_n . Let μ be the uniform distribution and $\gamma(n)$ be a real-valued function. If for each 2-dimensional rectangle R with $(1 - \delta)n/8$ singletons it holds that $\mu(R \cap \operatorname{IP}_n^{-1}(0)) \geq \mu(R \cap \operatorname{IP}_n^{-1}(1)) - \gamma(n)$, the size of G is bounded below by

$$1/2[(\mu(\mathrm{IP}^{-1}(1)) - \epsilon)/\gamma(n)].$$

Fact 3: $\mu(\operatorname{IP}_n^{-1}(1)) = 1/2 - 1/2^{n/2+1}$ where μ is the uniform distribution on $\{0,1\}^n$.

Now, we prove that $\mu(R \cap \operatorname{IP}_n^{-1}(0)) \ge \mu(R \cap \operatorname{IP}_n^{-1}(1)) - 2^{-(1-\delta)n/16}$ for the uniform distribution μ and 2-dimensional rectangles R with $(1-\delta)n/8$ singeltons. Since R corresponds to a balanced partition of the input variables into two sets X_1 and X_2 , R can be described as a rectangle $A \times B$ where A is a set of assignments to X_1 and B a set of assignments to X_2 . We consider the set $X'_1 \subseteq X_1$ of all singletons in X_1 and the set $X'_2 \subseteq X_2$ of all singletons in X_2 . Each partial assignment to the variables outside $X'_1 \cup X'_2$ restricts R to some rectangle $A' \times B'$ where A' contains assignments to the variables in X'_1 and B' contains assignments to the variables in X'_2 . Moreover, IP_n is restricted to the subfunction $\operatorname{IP}_{(1-\delta)n/8}$ or its negation. Finally, the uniform distribution μ is restricted to the uniform distribution μ' on $\{0, 1\}^{(1-\delta)n/4}$, the input set for the variables in $X'_1 \cup X'_2$. Now, we use the fact that $Disc_{uniform}(\mathrm{IP}_{(1-\delta)n/8}) \leq 2^{-1/2(1-\delta)n/8}$ (Chor and Goldreich (1988)). We easily obtain the restricted balance property for IP_n and R by averaging over all partial assignments. Altogether, we have proved the following result.

Corollary 2. Let μ be the uniform distribution. Let G be a regular BP1 representing a c-approximation of IP_n with $c \geq (1/2 + 1/2^{n/2+1} + 2^{-\Omega(n^{1-\epsilon'})}), 0 < \epsilon' \leq 1$. If for all nodes v of G which can be reached after testing exactly n/2 variables it holds that there exist at least $(1-\delta)n/8 A(v)$ -singletons according to IP_n , then the size of G is bounded below by $2^{\Omega(n)}$.

Now, we are ready to prove the following theorem.

Theorem 6. There is no c-approximation of PIP_n with $c \ge (1/2 + 1/2^{n/2+1} + 1/2^{n^{1-\epsilon'}})$, $0 < \epsilon' \le 1$, which can be represented by BP1s of polynomial size.

Proof. We assume that there is a *c*-approximation g_n of PIP_n with $c \ge (1/2 + 1/2^{n/2+1} + 1/2^{n^{1-\epsilon'}})$ which can be represented by a BP1 *G* of polynomial size. First, we transform *G* to a regular BP1 *G'* representing g_n . By our assumption *G'* is of polynomial size *s*. We define a cut through *G'* consisting of all nodes *v* such that on each path from the source of *G'* to *v* exactly n/2 data variables have been tested before reaching *v*. By our assumption on *G'* on all paths to *v* the same set A(v) of variables has been tested. By the definition of PIP_n some permutations π are described by two codewords. Therefore, we double the error probability of Fact 2. Since the number of nodes in *G'* is bounded above by *s* it follows that for at most $2se^{-4\delta^2n}2^{\Theta(n\log n)}$ subfunctions IP_n^{π} of PIP_n there exists a node *v* on the cut of *G'* such that there are at most $(1-\delta)n/8 A(v)$ -singletons according to π . Furthermore, $Pr(IP_n^{\pi'}(\tilde{x}) = g'_n(\tilde{x})) \ge (1-2se^{-4\delta^2n})c =: c'$ for a random input \tilde{x} where g'_n is the subfunction of g_n choosing a setting to the selection variables which corresponds to a codeword for π' . Since $s \ll e^{2\delta^2n}$ if *n* is large enough, it holds that $2se^{-4\delta^2n} < e^{-2\delta^2n}$. Using Corollary 2 it follows

$$\begin{aligned} |G'| &\geq 1/2(1/2 - 1/2^{n/2+1} - (1 - c'))2^{(1-\delta)n/16} \\ &= 1/2(-1/2 - 1/2^{n/2+1} + c')2^{(1-\delta)n/16} \\ &\geq 1/2(-1/2 - 1/2^{n/2+1} + (1 - e^{-2\delta^2 n})(1/2 + 1/2^{n/2+1} + 1/2^{n^{1-\epsilon'}}))2^{(1-\delta)n/16} \\ &\geq (1/2)2^{(1-\delta)n/16 - n^{1-\epsilon'} - 1} \\ &= 2^{\Omega(n)}. \end{aligned}$$

Since $|G| \ge |G'|/(n+1)$, there is a contradiction of our assumption that G is of polynomial size.

Conclusion

Problems from learning theory and genetic programming motivate the investigation of the approximative representation of Boolean functions by BDDs. Lower bounds for the approximative representation are shown to be harder to obtain as similar bounds for randomized representations. Exponential lower bounds were known for OBDDs and are proved here for the more general models k-IBDDs and read-once branching programs.

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