#### A-002

# A Fixed-Parameter Algorithm for Detecting a Singleton Attractor in an AND/OR Boolean Network with Bounded Treewidth

Tatsuya Akutsu<sup>†</sup>

Takeyuki Tamura<sup>†</sup>

### 1 Introduction

The Boolean network (BN) is known as a discrete mathematical model of gene regulatory networks [3]. In a BN, each vertex corresponds to a gene and takes one of two values 0 and 1, where 0 (resp., 1) means that the corresponding gene is inactive (resp., active). The value of a vertex at a given time step is determined according to a regulation rule, which is a Boolean function of the values of the predecessors of the vertex at the previous time instant. The values of vertices are updated synchronously, and the (global) state of a network at a given time step is the vector of its vertex values. Beginning from any initial state, the system eventually falls into an *attractor*, which is classified into two types: a singleton attractor corresponding to a stable state, and a *periodic attractor* corresponding to a sequence of states that repeats periodically.

It is known that the problem of finding an attractor of the shortest period is NP-hard even for BNs with maximum in-degree 2 consisting of AND/OR of literals [5]. Due to this hardness and the fact that there exist  $2^n$  global states for a BN with *n* vertices, previous theoretical studies focused on the development of  $o(2^n)$  time algorithms.

Since it is quite hard to develop such algorithms for general BNs, some restrictions were assumed on types of Boolean functions in all studies. For example, an  $O(1.587^n)$  time algorithm and an  $O(1.985^n)$ time algorithm were developed for detection of a singleton attractor [4] and an attractor of period 2 [1], respectively, both for AND/OR BNs which are BNs consisting of Boolean functions restricted to conjunctions and disjunctions of literals.

An  $O(n^{2p(w+1)}poly(n))$  time algorithm was also developed for finding an attractor of period p of a BN having bounded treewidth w and consisting of nested canalyzing functions, where p and w are constants, and nested canalyzing functions are a super class of AND/OR functions [1]. They also presented a fixedparameter algorithm (precisely, an algorithm working in O(q(p, w, d)poly(n)) time where q(p, w, d) depends only on p, w, d for a general BN with bounded degree and bounded treewidth [1]. However, it is unknown whether there exists a fixed-parameter algorithm even for an AND/OR BN with bounded treewidth but without degree constraint. In this short report, we present a fixed-parameter algorithm for detection of a singleton attractor in an AND/OR BN with bounded treewidth.

#### 2 Preliminaries

A BN N(V, F) consists of a set V of n vertices and a corresponding set  $F = \{f_v : v \in V\}$  of n Boolean functions. Let  $v(t) \in \{0,1\}$  represent the value of a vertex v at time t, and denote by  $\mathbf{v}(t) = \langle v(t) : v \in V \rangle$ the state of the network at time t. The values of all vertices are updated simultaneously according to the corresponding Boolean functions,  $v(t+1) = f_v(\mathbf{v}(t))$ . A directed graph G(V, E) can be associated with the network, with a directed edge  $(u, v) \in E$  if and only if  $f_v$  depends on u, where edges may be self-loops. The initial assignment of values  $\mathbf{v}(1) = \langle v(1) : v \in V \rangle$ uniquely determines the state of the network at all t > 1. An initial state is called a *periodic attractor* with period p if  $\mathbf{v}(1) = \mathbf{v}(p+1)$  and  $\mathbf{v}(1) \neq \mathbf{v}(q)$  holds for all 1 < q < p + 1. An attractor with period p = 1is called a *singleton attractor*.

A tree decomposition of a graph G(V, E) is a pair  $\langle T(V_T, E_T), (B_t)_{t \in V_T} \rangle$ , where  $T(V_T, E_T)$  is a rooted tree and  $(B_t)_{t \in V_T}$  is a family of subsets of V such that

- for every  $v_i \in V$ ,  $B^{-1}(v_i) = \{t \in V_T | v_i \in B_t\}$  is nonempty and connected in T, and
- for every edge  $\{v_i, v_j\} \in E$ , there exists  $t \in V_T$ such that  $v_i, v_j \in B_t$ .

The width of the decomposition is defined as  $\max_{t \in V_T} (|B_t| - 1)$  and the treewidth of G is the minimum of the widths among all the tree decompositions of G. Graphs with treewidth at most k are also known as partial k-trees [2].

## 3 Algorithm

Let  $\langle T(V_T, E_T), (B_t)_{t \in V_T} \rangle$  be a tree decomposition of G(V, E) associated to a given BN N(V, F). Let des(t) denote the set of descendant of  $t \in V_T$  with including t. Let  $V_t = \bigcup_{t' \in des(t)} B_{t'}$ . For each  $t \in V_T$ , p(t) denotes the parent node of t in  $V_T$ .<sup>‡</sup>

For each  $t \in V_T$ ,  $\phi_t$  denotes a function from  $B_t$  to  $\{0, 1, \omega\}$ , where we call such  $\phi_t$  an *assignment*. We define  $\hat{\phi}_t$  by

$$\hat{\phi}_t(v) = \begin{cases} \phi_t(v), & \text{if } \phi_t(v) \in \{0, 1\}, \\ 0, & \text{if } \phi_t(v) = \omega \text{ and } f_v \text{ is AND}, \\ 1, & \text{if } \phi_t(v) = \omega \text{ and } f_v \text{ is OR}, \end{cases}$$

It is to be noted that if  $\hat{\phi}_t(v) = 1$  and v is an AND vertex (resp.,  $\hat{\phi}_t(v) = 0$  and v is an OR vertex), Boolean

<sup>&</sup>lt;sup>†</sup>Institute for Chemical Research, Kyoto University

<sup>&</sup>lt;sup>‡</sup>We use nodes and vertices for  $\langle T(V_T, E_T), (B_t)_{t \in V_T} \rangle$ and N(V, F), respectively.

values of its input vertices are uniquely determined (in a singleton attractor). Therefore,  $\phi_t(v) = \omega$  holds only if v is an AND vertex and  $\hat{\phi}_t(v) = 0$ , or v is an OR vertex and  $\hat{\phi}_t(v) = 1$ .

For a 0-1 assignment  $\alpha$  to  $V' \subseteq V$ ,  $\phi_t(v) \in \{0, 1\}$ is called *validated* (by  $\alpha$  on V') if the Boolean value v is uniquely determined as  $\alpha(v)$  from  $\alpha$  regardless of an assignment to V - V'. We say that  $\alpha$  *violates* a Boolean function  $f_v$  (assigned to vertex v) if the value  $(b_v)$  of  $f_v$  is uniquely determined by  $\alpha$  but  $b_v \neq \alpha(v)$ . It is to be noted that we need not validate if  $\hat{\phi}_t(v) = 1$ and v is an AND vertex, or,  $\hat{\phi}_t(v) = 0$  and v is an OR vertex since it is enough for such a vertex to examine whether  $\alpha$  does not violate  $f_v$ .  $\phi_t$  is called *consistent* if the following is satisfied

- (1)  $\phi_t(v) \in \{0, 1\}$  holds for all  $v \in B_t B_{p(t)}$ , where we let  $B_{p(t)} = \emptyset$  if t is the root,
- (2) there exists a 0-1 assignment  $\alpha$  to  $V_t$  such that  $\hat{\phi}_t(v) = \alpha(v)$  holds for all  $v \in B_t$ ,  $\alpha$  does not violate any Boolean function assigned to  $V_t$ , and  $\phi_t(v)$  is validated for all v such that  $\phi_t(v) \in \{0, 1\}$  by  $\alpha$  on  $V_t$ .

Let  $A_t$  be the set of consistent assignments to  $B_t$ . We describe below how to compute  $A_t$  by dynamic programming from leaves to the root in  $V_T$ . For each leaf t,  $A_t$  is determined by  $A_t = \{\phi_t | \phi_t \text{ satisfies conditions (1) and (2) for } \alpha = \hat{\phi}_t.\}$ .

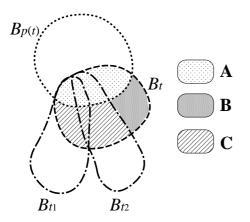


Figure 1: Part A  $(B_t \cap B_{p(t)})$  can be validated later, Part B  $(B_t - B_{p(t)} - \bigcup_j B_{t_j})$  must be validated by  $\phi_t$ , and Part C  $((B_t \cap (\bigcup_j B_{t_j})) - B_{p(t)})$ must be validated by  $\phi_t$ ,  $\phi_{t_1}$ ,  $\phi_{t_2}$ .

Let  $t_1, \ldots, t_d$  be the children of t. For all possible  $\phi_t$ , we examine whether it is consistent as follows. We call  $\phi_t$  and  $\phi_{t_i}$  are *compatible* if, for each  $v \in B_t \cap B_{t_i}$ ,  $\phi_t(v) = \phi_{t_i}(v)$  holds, or  $\phi_t(v) \in \{0, 1\}$ ,  $\phi_{t_i}(v) = \omega$  and  $\hat{\phi}_t(v) = \hat{\phi}_{t_i}(v)$  hold. We maintain a 0-1 table  $X(x_1, \ldots, x_h)$  where  $\{v_{j_1}, \ldots, v_{j_h}\} = B_t$  (i.e., X is a 0-1 table having  $2^{|B_t|}$  entries). We say that  $\phi_t$  is a *candidate* assignment if  $\phi_t(v) \in \{0, 1\}$  holds for any

vertex in  $B_t - B_{t(p)}$ ,  $\phi_t$  does not violate any function assigned to  $B_t$ , and  $\phi_t(v)$  is validated by  $\phi_t$  for any vertex  $B_t - B_{t(p)} - \bigcup_{j=1,\dots,d} B_{t_j}$ . The following is a pseudocode of the proposed algorithm (see also Fig. 1).

Procedure FpFindAttfor all leaves in  $V_T$  do compute  $A_t$ ; for all internal nodes t in  $V_T$  do  $A_t \leftarrow \emptyset;$ for all candidate assignments  $\phi_t$  do for all  $(x_1, \ldots, x_h) \in \{0, 1\}^h$  do  $X(x_1,\ldots,x_h) \leftarrow 0;$  $X(x_1,\ldots,x_h) \leftarrow 1$  where  $x_g = 1$  iff  $\phi_t(v_{j_g})$ has already been validated, or  $\phi_t(v_{j_q}) = \omega$ ; for i = 1 to d do  $Y \leftarrow X;$ for all  $\phi_{t_i} \in A_{t_i}$  compatible with  $\phi_t$  do let  $z_g = 1$  if  $\phi_{t_i}(v_{j_g}) \in \{0, 1\},\$ otherwise  $z_g = 0;$ for all  $(x_1, ..., x_h)$  with  $X(x_1, ..., x_h) = 1$ do  $Y(\max(x_1, z_1), \ldots, \max(x_h, z_h)) \leftarrow 1;$  $X \leftarrow Y;$ if  $X(1,1,\ldots,1) = 1$  then  $A_t \leftarrow A_t \cup \{\phi_t\}$ 

There exists a singleton attractor iff  $A_r \neq \emptyset$  for the root r of T. Furthermore, such a singleton attractor can be retrieved by using the standard traceback technique. Since the number of possible  $\phi_t$  is bounded by  $3^{k+1}$  per t and the size of table X is bounded by  $2^{k+1}$ for partial k-trees, we have:

**Theorem 1** The singleton attractor detection problem for an AND/OR BN with bounded treewidth kcan be solved in O(f(k)poly(n)) time, where f(k) is a function depending only on k.

### References

- T. Akutsu, S. Kosub, A. A. Melkman, and T. Tamura, Finding a periodic attractor of a Boolean Network, *IEEE/ACM Trans. Computational Biology and Bioinformatics* 9:151–160, 2012.
- [2] J. Flum and M. Grohe, *Parameterized Complexity Theory*, Springer, 2006.
- [3] S. A. Kauffman, The Origins of Order: Selforganization and Selection in Evolution, Oxford Univ. Press, 1993.
- [4] A. A. Melkman, T. Tamura, and T. Akutsu, Determining a singleton attractor of an AND/OR Boolean network in O(1.587<sup>n</sup>) time, Information Processing Letters, 110:565–569, 2010.
- [5] T. Tamura and T. Akutsu, Detecting a singleton attractor in a Boolean network utilizing SAT algorithms, *IEICE Trans. Fundamentals*, E92-A:493-501, 2009.