

• Supplementary File •

# Aggregation Method to Reachability and Optimal Control of Large-Size Boolean Control Networks

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## Appendix A Notations used in the body of this paper

- $\dot{\cup}$  denotes the disjoint union of sets.
- $\Delta_p := \{\delta_p^j := Col_j(I_p) : j = 1, \dots, p\}$ . For example,  $\delta_3^1 := [1 \ 0 \ 0]^\top$ ,  $\delta_3^2 := [0 \ 1 \ 0]^\top$ ,  $\delta_3^3 := [0 \ 0 \ 1]^\top$ ,  $\Delta_3 := \{\delta_3^j : j = 1, 2, 3\}$ .
- $\mathcal{L}_{k \times h} := \{[\delta_k^{i_1} \ \dots \ \delta_k^{i_h}] := \delta_k[i_1 \ \dots \ i_h] : i_j \in \{1, \dots, k\}, j = 1, \dots, h\}$ .
- $N = \{x_1, x_2, \dots, x_n\}$  denotes a set of  $n$  nodes, and  $M = \{x_{i_1}, x_{i_2}, \dots, x_{i_m}\} \subseteq N$ ,  $1 \leq m < n$ . Denote the state of node  $x_i$  by  $\varsigma_i$ , where  $\varsigma_i \in \Delta_2$ ,  $i = 1, \dots, n$ . Then, the natural projection from  $N$  to  $M$ , denoted by  $\sigma_{N,M} : \Delta_{2^n} \rightarrow \Delta_{2^m}$ , is defined as  $\sigma_{N,M}(\times_{i=1}^n \varsigma_i) = \times_{j=1}^m \varsigma_{i_j}$ . In addition, for a nonempty ordered set  $N' = \{a_1, \dots, a_s\} \in \Delta_{2^n}^s$ ,  $\sigma_{N,M}(N') := \{\sigma_{N,M}(a_1), \dots, \sigma_{N,M}(a_s)\}$ ; for a set group  $N'' = \{N'_i \in \Delta_{2^n}^s : N'_i \neq \emptyset, i = 1, \dots, h\}$ ,  $\sigma_{N,M}(N'') := \{\sigma_{N,M}(N'_i) : i = 1, \dots, h\}$ .

## Appendix B Proofs in the body of the letter

### Appendix B.1 Proof of Corollary 1

When  $Z_i = \emptyset$ , this theorem becomes Theorem 3.3 in [1]. Thus, we only consider  $Z_i \neq \emptyset$  in the following proof.

When  $\kappa = 1$ , by Definition 1,  $\alpha_i^d = \delta_{2^{n_i}}^{\theta_i}$  is reachable from  $\alpha_i^0 = \delta_{2^{n_i}}^{\lambda_i}$  at the first step, if and only if there exists  $\beta_i(0) := \delta_{2^{m_i}}^\mu \in \Delta_{2^{m_i}}$  such that

$$\delta_{2^{n_i}}^{\theta_i} = F_i \delta_{2^{q_i}}^{\xi_i} \delta_{2^{m_i}}^\mu \delta_{2^{n_i}}^{\lambda_i}.$$

Noticing that  $F_i \delta_{2^{q_i}}^{\xi_i} \delta_{2^{m_i}}^\mu = F_i \delta_{2^{m_i+q_i}}^{2^{m_i}(\xi_i-1)+\mu}$ , that is,  $\delta_{2^{n_i}}^{\theta_i} = F_i \delta_{2^{m_i+q_i}}^{2^{m_i}(\xi_i-1)+\mu} \delta_{2^{n_i}}^{\lambda_i}$ , we have  $[F_i \delta_{2^{m_i+q_i}}^{2^{m_i}(\xi_i-1)+\mu}]_{\theta_i, \lambda_i} = 1 > 0$ . Then, it is easy to see that

$$\begin{aligned} [R_i(1)]_{\theta_i, \lambda_i} &= [M_i(0)]_{\theta_i, \lambda_i} \\ &= \left[ \sum_{j=1}^{2^{m_i}} F_i \delta_{2^{m_i+q_i}}^{2^{m_i}(\xi_i-1)+j} \right]_{\theta_i, \lambda_i} \\ &\geq [F_i \delta_{2^{m_i+q_i}}^{2^{m_i}(\xi_i-1)+\mu}]_{\theta_i, \lambda_i} > 0. \end{aligned}$$

Hence, the conclusion is true for  $\kappa = 1$ .

Assuming the truth of the conclusion for  $\kappa = s > 1$ , we prove the truth of the conclusion for  $\kappa = s + 1$ . When  $\kappa = s + 1$ , we divide the proof into two steps. Firstly, there exists  $\delta_{2^{n_i}}^{\zeta_i} \in \Delta_{2^{n_i}}$  such that  $\delta_{2^{n_i}}^{\zeta_i}$  is reachable from  $\delta_{2^{n_i}}^{\lambda_i}$  at the  $s$ -th step. Secondly,  $\delta_{2^{n_i}}^{\theta_i}$  is reachable from  $\delta_{2^{n_i}}^{\zeta_i}$  at the first step. Then, we have  $[R_i(s)]_{\zeta_i, \lambda_i} > 0$  and  $[M_i]_{\theta_i, \zeta_i} > 0$ . Therefore,

$$\begin{aligned} [R_i(s+1)]_{\theta_i, \lambda_i} &= [M_i^s M_i(0)]_{\theta_i, \lambda_i} \\ &= \sum_{j=1}^{2^{n_i}} [M_i]_{\theta_i, j} [M_i^{s-1} M_i(0)]_{j, \lambda_i} \\ &\geq [M_i]_{\theta_i, \zeta_i} [R_i(s)]_{\zeta_i, \lambda_i} > 0. \end{aligned}$$

Therefore, the conclusion is true for  $\kappa = s + 1$ .

By induction, the conclusion is true for any positive integer  $\kappa$ .

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## Appendix B.2 Proof of Theorem 1

(Sufficiency) From condition (i) and Corollary 1, subnetwork  $\Sigma_i$  is reachable from  $\alpha_i^0 = \delta_{2n_i}^{\lambda_i}$  to  $\alpha_i^d = \delta_{2n_i}^{\theta_i}$  at the  $\kappa$ -th step,  $i = 1, 2, \dots, \rho$ . Then, for any subnetwork  $\Sigma_i$ ,  $i \in \Phi_2$ , one can obtain  $\Omega_i$ .

By condition (ii) and Definition 2, there exists at least one  $\kappa$ -matchable control sequence, denoted by  $\{w_i^{a_i, c_i} : i \in \Phi_2\}$ , where  $w_i^{a_i, c_i} \in \Omega_i$ ,  $a_i \in \{1, \dots, b_i\}$  and  $c_i \in \{1, \dots, |\Omega_i^{a_i}|\}$ .

Then, for this  $\kappa$ -matchable control sequence, one can obtain a control sequence driving BCN (1) from  $x^0$  to  $x^d$  at the  $\kappa$ -th step, denoted by  $u = \{u(t) : t = 0, \dots, \kappa - 1\}$ , where  $u(t) = \times_{j=1}^n u_j(t)$ . In fact, if  $u_j \in U_i$ , then  $\{u_j(t) : t = 0, \dots, \kappa - 1\}$  can be constructed as  $\{u_j(t) : t = 0, \dots, \kappa - 1\} = \sigma_{Z_i \cup U_i, \{u_j\}}(w_i^{a_i, c_i})$ . Therefore, under the control  $u = \{u(t) : t = 0, \dots, \kappa - 1\}$ , BCN (1) is reachable from  $x^0$  to  $x^d$  at the  $\kappa$ -th step.

(Necessity) If BCN (1) is reachable from  $x^0$  to  $x^d$  at the  $\kappa$ -th step, then there exists at least one control sequence  $\{u(t) : t = 0, \dots, \kappa - 1\}$  satisfying  $x(\kappa; x^0, u) = x^d$ . Denote the corresponding state trajectory by  $\{x(t) : t = 0, \dots, \kappa\}$ , where  $x(0) = \delta_{2n}^{\lambda}$ ,  $x(\kappa) = \delta_{2n}^{\theta}$ . Obviously, subnetwork  $\Sigma_i$ ,  $i \in \Phi_1$  is reachable from  $\alpha_i^0 = \delta_{2n_i}^{\lambda_i}$  to  $\alpha_i^d = \delta_{2n_i}^{\theta_i}$  at the  $\kappa$ -th step, and the state trajectory is  $T_i^1 = \{\delta_{2n_i}^{t_0} \rightarrow \delta_{2n_i}^{t_1} \rightarrow \dots \rightarrow \delta_{2n_i}^{t_\kappa}\}$ , where  $\delta_{2n_i}^{t_\kappa} = \sigma_{X, X_i}(x(t))$ ,  $t = 0, 1, \dots, \kappa$ . In addition, under the following control sequence, subnetwork  $\Sigma_i$ ,  $i \in \Phi_2$  is reachable from  $\alpha_i^0 = \delta_{2n_i}^{\lambda_i}$  to  $\alpha_i^d = \delta_{2n_i}^{\theta_i}$  at the  $\kappa$ -th step:

$$w_i^{a_i, c_i} := \{\tilde{u}_i(t) : t = 0, \dots, \kappa - 1\}, \quad (\text{B1})$$

where  $\tilde{u}_i(t) = \gamma_i(t) \times \beta_i(t)$ ,  $\gamma_i(t) = \sigma_{X, Z_i}(x(t))$ , and  $\beta_i(t) = \sigma_{U, U_i}(u(t))$ . Then, by Corollary 1,  $[R_i(\kappa)]_{\theta_i, \lambda_i} > 0$ ,  $\forall i = 1, 2, \dots, \rho$ . Therefore, condition (i) holds.

Denote the state trajectory corresponding to  $w_i^{a_i, c_i}$  by  $T_i^{a_i} = \{\delta_{2n_i}^{t_0^{a_i}} \rightarrow \delta_{2n_i}^{t_1^{a_i}} \rightarrow \dots \rightarrow \delta_{2n_i}^{t_\kappa^{a_i}}\}$ . Then

$$\delta_{2n_i}^{t_\kappa^{a_i}} = \sigma_{X, X_i}(x(t)), \quad (\text{B2})$$

$t = 0, 1, \dots, \kappa$ . Set  $\tilde{T}_i^{a_i} := \{\delta_{2n_i}^{t_0^{a_i}}, \delta_{2n_i}^{t_1^{a_i}}, \dots, \delta_{2n_i}^{t_{\kappa-1}^{a_i}}\}$ ,  $i = 1, 2, \dots, \rho$ , and for any  $i \in \Phi_1$ ,  $a_i = 1$ . For any  $i \in \{1, 2, \dots, \rho\}$ ,  $j \in \Phi_2$ ,  $i \neq j$ , denote  $Y_i^j = \{y_{i, j_r} : j_r \in \{1, \dots, p_i\}, r = 1, \dots, p_i^j\}$ , whose elements keep the order in  $Y_i$ . Then, for any  $i \in \{1, 2, \dots, \rho\}$ ,  $j \in \Phi_2$ ,  $i \neq j$ , on one hand, by (B2),  $\sigma_{X_i, Y_i^j}(\tilde{T}_i^{a_i}) = \{\times_{r=1}^{p_i^j} y_{i, j_r}(t) : t = 0, \dots, \kappa - 1\}$ ; on the other hand, by (B1),  $\sigma_{Z_j \cup U_j, Z_j^i}(w_j^{a_j, c_j}) = \{\times_{r=1}^{p_j^i} y_{i, j_r}(t) : t = 0, \dots, \kappa - 1\}$ . Therefore,  $\sigma_{X_i, Y_i^j}(\tilde{T}_i^{a_i}) = \sigma_{Z_j \cup U_j, Z_j^i}(w_j^{a_j, c_j})$  holds for any  $i \in \{1, 2, \dots, \rho\}$ ,  $j \in \Phi_2$ ,  $i \neq j$ . By Definition 2,  $\{w_i^{a_i, c_i} : i \in \Phi_2\}$  is a  $\kappa$ -matchable control sequence, that is,  $\mathcal{M} \neq \emptyset$ , which implies that condition (ii) holds.

## Appendix B.3 Proof of Proposition 2

On one hand, assume that large-size BCN (2) is reachable from  $x^0 = \delta_{2n}^{\lambda}$  to  $x^d = \delta_{2n}^{\theta}$  at the  $\kappa$ -th step. Then, at least one control sequence can be obtained, denoted by  $\{u(t) : t = 0, \dots, \kappa - 1\}$ , and the corresponding state trajectory is  $\{x(t) : t = 0, \dots, \kappa\}$ , where  $x(0) = x^0$ ,  $x(\kappa) = x^d$ . For any partition  $\Xi$  satisfying Assumption 1, assume that  $\Xi$  contains  $\hat{\rho}$  subnetworks, denoted by  $\hat{\Sigma}_i$ ,  $i = 1, 2, \dots, \hat{\rho}$ . In addition, for each subnetwork  $\hat{\Sigma}_i$ , denote the parameters by  $\hat{X}_i$ ,  $\hat{U}_i$ ,  $\hat{Z}_i$ ,  $\hat{Y}_i$ ,  $\hat{n}_i$ ,  $\hat{m}_i$ ,  $\hat{q}_i$ ,  $\hat{p}_i$ ,  $\hat{\alpha}_i^0 = \sigma_{X, \hat{X}_i}(x^0) = \delta_{2n_i}^{\lambda_i}$ , and  $\hat{\alpha}_i^d = \sigma_{X, \hat{X}_i}(x^d) = \delta_{2n_i}^{\theta_i}$ . Denote  $\hat{\Phi}_1 := \{i \in \{1, 2, \dots, \hat{\rho}\} : \hat{Z}_i \cup \hat{U}_i = \emptyset\}$  and  $\hat{\Phi}_2 := \{1, 2, \dots, \hat{\rho}\} \setminus \hat{\Phi}_1$ . Then, by virtue of the necessity part of Theorem 1, for each subnetwork  $\hat{\Sigma}_i$ ,  $i \in \hat{\Phi}_1$ , it is obvious that  $\hat{\Sigma}_i$  is reachable from  $\hat{\alpha}_i^0 = \delta_{2n_i}^{\lambda_i}$  to  $\hat{\alpha}_i^d = \delta_{2n_i}^{\theta_i}$  at the  $\kappa$ -th step; for each subnetwork  $\hat{\Sigma}_i$ ,  $i \in \hat{\Phi}_2$ , one can obtain that  $\hat{\Sigma}_i$  is reachable from  $\hat{\alpha}_i^0 = \delta_{2n_i}^{\lambda_i}$  to  $\hat{\alpha}_i^d = \delta_{2n_i}^{\theta_i}$  at the  $\kappa$ -th step under control sequence  $\hat{w}_i^{a_i, c_i} := \{\tilde{u}_i(t) : t = 0, \dots, \kappa - 1\}$ . Thus, by Corollary 1, one can conclude that condition (i) in Theorem 1 holds. In addition, one can verify that  $\{\hat{w}_i^{a_i, c_i} : i \in \hat{\Phi}_2\}$  is a  $\kappa$ -matchable control sequence. Thus, condition (ii) in Theorem 1 holds.

On the other hand, given a partition  $\Xi$  (here, we assume that the parameters of  $\Xi$  is the same as the above  $\Xi$ ) satisfying Assumption 1, and suppose that the condition (i) and condition (ii) of Theorem 1 are satisfied. Then, according to Definition 2, one can obtain a  $\kappa$ -matchable control sequence, denoted by  $\{\hat{w}_i^{a_i, c_i} : i \in \hat{\Phi}_2\}$ . In addition, based on the sufficiency part of Theorem 1, from the above  $\kappa$ -matchable control sequence, a control sequence  $u = \{u(t) : t = 0, \dots, \kappa - 1\}$  driving BCN (2) from  $x^0$  to  $x^d$  at the  $\kappa$ -th step can be obtained. Therefore, by resorting to the first part of this proof, it is easy to see that for any other partition  $\Xi'$  satisfying Assumption 1, the conditions in Theorem 1 are still satisfied.

## Appendix C Algorithm of obtaining an acyclic aggregated graph which satisfies Assumption 1

Consider large-size BCN (1) and denote the network graph of (1) by  $G = (N, E)$ .

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**Algorithm C1** The algorithm of obtaining an acyclic aggregated graph which satisfies Assumption 1.

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- Step 1: Obtain all strongly connected components of  $G$ , and consider each one as a super node;
  - Step 2: If there exists at least one state node in every super node, then an acyclic aggregated graph which satisfies Assumption 1 is obtained and stop. Otherwise, arbitrarily choose a super node  $N_i$  contains no state node, and go to Step 3;
  - Step 3: Choose another super node  $N_j$  satisfying the condition that there exists an edge from some  $v_s \in N_i$  to some  $v_t \in N_j$ . Then, combine  $N_i$  and  $N_j$  to form a super node and go back to Step 2.
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## Appendix D Computational complexity analysis of Theorem 1

Given an aggregation of large-size BCN (2) which contains  $\rho$  subnetworks. For subnetwork  $\Sigma_i$ ,  $i = 1, 2, \dots, \rho$ , there exist at most  $2^{\kappa(m_i+q_i)}$  control sequences which can drive  $\Sigma_i$  from  $\alpha_i^0$  to  $\alpha_i^d$  at the  $\kappa$ -th step. Let  $\zeta := \max_{i \in \{1, 2, \dots, \rho\}} \{m_i + q_i\}$ . Then, in order to verify the reachability of BCN (2) via Theorem 1, one needs to handle matrices of sizes  $2^{n_i} \times 2^{m_i+q_i}$ ,  $i = 1, 2, \dots, \rho$  and enumerate at most  $2^{\rho\zeta}$  combinations of control sequences. Therefore, the time complexity of Theorem 1 is exponential in the number of nodes. However, the establishment Theorem 1 makes it possible to verify the reachability of large-size BCNs in the following two special cases: (i) Note that it is feasible to verify the reachability of each subnetwork. When there exists a subnetwork which is not reachable, the original large-size BCN is not reachable. (ii) When  $|\Omega_i|$  is very small, say  $|\Omega_i| \ll 2^{\zeta\kappa}$ , it is possible to verify the  $\kappa$ -matchable condition. In the future, we devote to reducing the computational complexity of Theorem 1 for the application to general large-size BCNs.

## Appendix E Examples

### Appendix E.1 An example used to illustrate how Corollary 1 and Proposition 1 work

Consider the following BCN:

$$\begin{cases} x_1(t+1) = x_1(t) \vee (x_2(t) \wedge x_3(t)), \\ x_2(t+1) = x_2(t) \vee x_3(t), \\ x_3(t+1) = \neg x_3(t), \\ x_4(t+1) = x_5(t) \wedge u(t), \\ x_5(t+1) = x_4(t) \vee u(t) \vee x_2(t), \\ x_6(t+1) = x_8(t) \vee x_5(t), \\ x_7(t+1) = x_6(t) \vee x_3(t), \\ x_8(t+1) = x_7(t), \end{cases} \quad (\text{E1})$$

where  $x_i$ ,  $i = 1, \dots, 8$  and  $u$  denote states and control input, respectively. Fig. 1 shows an aggregation of BCN (E1). Denote the subnetwork corresponding to  $N_i$  by  $\Sigma_i$ ,  $i = 1, 2, 3$ . Letting  $x^0 = \delta_{256}^{25}$ ,  $x^d = \delta_{256}^{163}$  and  $\kappa = 3$ , we consider the reachability of subnetworks  $\Sigma_1$ ,  $\Sigma_2$  and  $\Sigma_3$ , respectively.

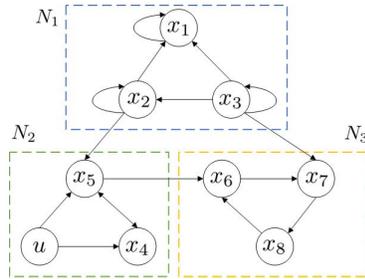


Fig. 1: Aggregation of BCN (E1).

For subnetwork  $\Sigma_1$ , the state trajectory from  $\alpha_1^0 = \delta_8^1$  to  $\alpha_1^d = \delta_8^6$  is  $T_1^1 = \{\delta_8^1 \rightarrow \delta_8^8 \rightarrow \delta_8^7 \rightarrow \delta_8^6\}$ .

Consider subnetwork  $\Sigma_2$ , we have  $Z_2 = \{x_2\}$ ,  $Y_2 = \{x_5\}$ . Calculating  $F_2$  and splitting  $F_2$  into 4 equal blocks, we have  $[M_2^2 M_2(0)]_{1,4} = 4 > 0$ . By Corollary 1, subnetwork  $\Sigma_2$  is reachable from  $\alpha_2^0$  to  $\alpha_2^d$  at the third step. By calculation, all possible state trajectories of  $\Sigma_2$  are  $T_2^1 = \{\delta_4^4 \rightarrow \delta_4^3 \rightarrow \delta_4^1 \rightarrow \delta_4^1\}$ ,  $T_2^2 = \{\delta_4^4 \rightarrow \delta_4^3 \rightarrow \delta_4^3 \rightarrow \delta_4^1\}$  and  $T_2^3 = \{\delta_4^4 \rightarrow \delta_4^4 \rightarrow \delta_4^3 \rightarrow \delta_4^1\}$ . In addition, by resorting to Proposition 1, the set of control sequences is

$$\Omega_2 = \left\{ \{\delta_4^1, \delta_4^3, \delta_4^1\}, \{\delta_4^1, \delta_4^2, \delta_4^3\}, \{\delta_4^2, \delta_4^2, \delta_4^3\}, \{\delta_4^2, \delta_4^3, \delta_4^3\} \right\}.$$

Subnetwork  $\Sigma_3$  is reachable from  $\alpha_3^0 = \delta_8^1$  to  $\alpha_3^d = \delta_8^3$  at the third step, and the corresponding state trajectories are  $T_3^1 = \{\delta_8^1 \rightarrow \delta_8^3 \rightarrow \delta_8^6 \rightarrow \delta_8^3\}$ ,  $T_3^2 = \{\delta_8^1 \rightarrow \delta_8^3 \rightarrow \delta_8^2 \rightarrow \delta_8^3\}$ . In addition, the set of control sequences is  $\Omega_3 = \left\{ \{\delta_4^2, \delta_4^3, \delta_4^3\}, \{\delta_4^2, \delta_4^4, \delta_4^1\} \right\}$ .

### Appendix E.2 An example used to show how Theorem 1 works

Consider the BCN model of *Pseudomonas aeruginosa* QS system [2]. Given an aggregation shown in Fig. 2. Verify whether or not the *Pseudomonas aeruginosa* QS system is reachable from  $x^0 = \delta_{16777216}^{68870173}$  to  $x^d = \delta_{16777216}^{8706044}$  at the second step.

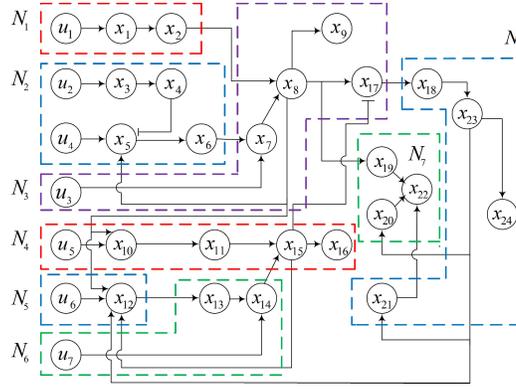


Fig. 2: Aggregation of *Pseudomonas aeruginosa* QS system.

Since  $[M_1 M_1(0)]_{3,2} = 1 > 0$ , subnetwork  $\Sigma_1$  is reachable from  $\delta_4^2$  to  $\delta_4^3$  at the second step. The corresponding state trajectory and the set of control sequences are  $T_1 = \{\delta_4^2 \rightarrow \delta_4^1 \rightarrow \delta_4^3\}$  and  $\Omega_1 = \{\{\delta_2^1, \delta_2^2\}\}$ , respectively. By calculation, subnetwork  $\Sigma_i$  is reachable from  $\alpha_i^0$  to  $\alpha_i^d$  at the second step,  $i = 2, 3, \dots, 8$ .

By Definition 2, one can obtain 64 different 2-matchable control sequences. Therefore, the *Pseudomonas aeruginosa* QS system is reachable from  $x^0$  to  $x^d$  at the second step. By Remark 1, the set of control sequences is

$$U = \left\{ \{\delta_{128}^i, \delta_{128}^j\} : i = 5, \dots, 8, 13, \dots, 16, 21, \dots, 24, 29, \dots, 32; j = 72, 80 \right\}.$$

### Appendix E.3 An example used to show the necessity of verifying $\kappa$ -matchable condition

Consider the Boolean network model of colitis-associated colon cancer with 70 nodes and 153 edges [3]. The network graph of colitis-associated colon cancer network is shown in Fig. 3.

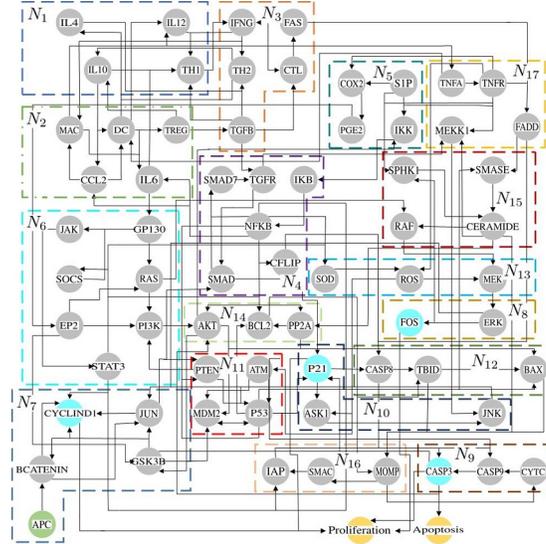


Fig. 3: Network graph of colitis-associated colon cancer network.

Colorectal cancer is one of the most common malignancies. It is shown that colorectal cancer is closely correlated with inflammation. The modeling and analysis of colitis-associated colon cancer network establish a framework for the study of inflammation-associated cancer [3]. Note that the existing results on colitis-associated colon cancer network are mainly based on experiments, and it is meaningful to develop a mathematical tool for the study of colitis-associated colon cancer network.

In the colitis-associated colon cancer network, “APC” denotes an input node, “Proliferation”, “Apoptosis” denote two output nodes, and the remaining ones are state nodes. According to [3], the output dynamics of colitis-associated colon cancer network is

$$\begin{cases} \text{Proliferation}(t) = (\text{FOS}(t) \wedge \text{CYCLIND1}(t)) \wedge \neg(\text{P21}(t) \vee \text{CASP3}(t)), \\ \text{Apoptosis}(t) = \text{CASP3}(t), \end{cases} \quad (\text{E2})$$

Given  $X_0 = (1011010111111011111111111111111011111111101110111111111111111011110)$  and  $X_d = (1111110111110111010111101001111011110111011101011010000111111111101)$

1 1 0 1 1). According to (E2), it is easy to verify that state  $X_0$  corresponds to “Proliferation” being “ON” and “Apoptosis” being “OFF”, while state  $X_d$  corresponds to “Proliferation” being “OFF” and “Apoptosis” being “ON”. In the following, we investigate the 3-step reachability of colitis-associated colon cancer network from state  $X_0$  to state  $X_d$  based the aggregation method. For full names and Boolean logical rules of nodes in colitis-associated colon cancer network, please refer to [3].

Choose the aggregation shown in Fig. 3, where the whole BCN is partitioned into 17 small-size subnetworks (Table 1), denoted by  $\Sigma_i$ ,  $i = 1, 2, \dots, 17$ .

**Table 1:** Notations of each subnetwork in aggregation.

	$X_i$	$Z_i \cup U_i$
$\Sigma_1$	TH1, IL4, IL12, IL10	TREG, TH2, TGFB, MAC, IFNG, CTL
$\Sigma_2$	TREG, MAC, IL6, DC, CCL2	NFKB, TNFA, IL10, IFNG
$\Sigma_3$	FAS, TH2, TGFB, IFNG, CTL	TREG, TH1, IL4
$\Sigma_4$	CFLIP, IKK, NFKB, SMAD, SMAD7, TGFR	IKK, JUN, TGFB
$\Sigma_5$	COX2, IKK, PGE2, S1P	AKT, SPHK1, TNFR
$\Sigma_6$	EP2, GP130, JAK, PI3K, RAS, STAT3	PGE2, PTEN, IL6
$\Sigma_7$	BCATENIN, CYCLIND1, GSK3B, JUN	AKT, EP2, ERK, JNK, STAT3
$\Sigma_8$	ERK, FOS	MEK
$\Sigma_9$	CASP3, CASP9, CYTC	CASP8, IAP, MOMP, P21
$\Sigma_{10}$	ASK1, JNK, P21	CASP3, GSK3B, MEKK1, P53, ROS, SMAD
$\Sigma_{11}$	ATM, MDM2, P53, PTEN	AKT, GSK3B, JNK, JUN, NFKB, ROS
$\Sigma_{12}$	BAX, CASP8, TBID	AKT, BCL2, CFLIP, FADD, P21, P53, PP2A
$\Sigma_{13}$	MEK, ROS, SOD	NFKB, RAF, STAT3, TNFR
$\Sigma_{14}$	AKT, BCL2, PP2A	CASP3, CERAMIDE, NFKB, P53, PI3K, STAT3
$\Sigma_{15}$	CERAMIDE, RAF, SMASE, SPHK1	ERK, FADD, P53, RAS, TNFR
$\Sigma_{16}$	IAP, MOMP, SMAC	BAX, BCL2, CERAMIDE, NFKB, STAT3, TBID
$\Sigma_{17}$	FADD, MEKK1, TNFR, TNFA	CERAMIDE, FAS, TGFR, MAC

Firstly, we consider the reachability of subnetworks. Take subnetwork  $\Sigma_3$  as an example. We have  $\alpha_3^0 = \delta_{32}^1$ ,  $\alpha_3^d = \delta_{32}^{18}$  and  $\gamma_3(0) = \delta_8^1$ . Since  $[M_3^2 M_3(0)]_{18,1} = 2 > 0$ , by Corollary 1,  $\alpha_3^d$  is reachable from  $\alpha_3^0$  at the third step. Similar to subnetwork  $\Sigma_3$ , one can verify that subnetwork  $\Sigma_i$  is reachable from  $\alpha_i^0$  to  $\alpha_i^d$  at the third step,  $i \in \{1, 2, \dots, 16\} \setminus \{3\}$ .

Next, we check the 3-matchable condition. For subnetwork  $\Sigma_4$  with  $\alpha_4^0 = \delta_{32}^1$ ,  $\alpha_4^d = \delta_{32}^{18}$  and  $\gamma_4(0) = \delta_8^1$ , there exists one state trajectory from  $\alpha_4^0$  to  $\alpha_4^d$  at the third step as  $T_4^1 = \{\delta_{64}^1 \rightarrow \delta_{64}^{30} \rightarrow \delta_{64}^{56} \rightarrow \delta_{64}^{21}\}$ . In addition, the set of control sequences is

$$\Omega_4 = \left\{ \{\delta_8^1, \delta_8^1, \delta_8^1\}, \{\delta_8^1, \delta_8^2, \delta_8^1\}, \{\delta_8^1, \delta_8^3, \delta_8^1\}, \{\delta_8^1, \delta_8^4, \delta_8^1\}, \{\delta_8^1, \delta_8^1, \delta_8^3\}, \{\delta_8^1, \delta_8^2, \delta_8^3\}, \{\delta_8^1, \delta_8^3, \delta_8^3\}, \{\delta_8^1, \delta_8^4, \delta_8^3\} \right\}.$$

Considering subnetwork  $\Sigma_3$ , there exists one state trajectory from  $\alpha_3^0$  to  $\alpha_3^d$  at the third step, that is,  $T_3^1 = \{\delta_{32}^1 \rightarrow \delta_{32}^{10} \rightarrow \delta_{32}^{32} \rightarrow \delta_{32}^{18}\}$ . Then, we have  $\tilde{T}_3^1 = \{\delta_{32}^1, \delta_{32}^{10}, \delta_{32}^{32}\}$ .

It is easy to see from Table 1 that  $Y_3^4 = Z_4^3 = \{\text{TGFB}\}$ . By a simple calculation, for  $\alpha_3(2) = \delta_{32}^{32}$ , one can obtain that  $\sigma_{X_3, Y_3^4}(\{\alpha_3(2)\}) = \{\delta_8^2\}$ . However, for  $\gamma_4(2) = \delta_8^3$  and  $\gamma_4(2) = \delta_8^3$ , it holds that  $\sigma_{Z_4, Z_3^4}(\{\gamma_4(2)\}) = \{\delta_8^1\}$ . Then, for each  $w_4 \in \Omega_4$ , we have  $\sigma_{X_3, Y_3^4}(\tilde{T}_3^1) \neq \sigma_{Z_4, Z_3^4}(w_4)$ . Therefore, according to Definition 2, the colitis-associated colon cancer network is not 3-matchable. Thus, the CACC network is not reachable from  $X_0$  to  $X_d$  at the third step.

From this example, one can see that for a given aggregation, although all subnetworks are reachable, the whole large-size network maybe not reachable, which supports the necessity of verifying  $\kappa$ -matchable condition.

## Appendix E.4 An example used to illustrate Remark 2

Recall the example in Appendix E.1. We check whether or not BCN (E1) is 3-matchable.

It is obvious that the aggregation given in Fig. 1 is an acyclic aggregation, and it holds that  $Y_i^j = Z_j^i = \emptyset$ ,  $i > j$ ,  $i, j = 1, 2, 3$ .

Since subnetwork  $\Sigma_1$  has no input, according to the unique state trajectory from  $\alpha_1^0$  to  $\alpha_1^d$ , that is,  $T_1^1 = \{\delta_8^1 \rightarrow \delta_8^8 \rightarrow \delta_8^7 \rightarrow \delta_8^6\}$ , we have  $\sigma_{X_1, Y_1^2}(\tilde{T}_1^1) = \{\delta_2^1, \delta_2^2, \delta_2^2\}$ ,  $\sigma_{X_1, Y_1^3}(\tilde{T}_1^1) = \{\delta_2^1, \delta_2^2, \delta_2^2\}$ , where  $\tilde{T}_1^1 = \{\delta_8^1, \delta_8^8, \delta_8^6\}$ .

By enumerating control sequences in  $\Omega_2$ , one can obtain that only control sequence  $w_2^{3,2} = \{\delta_4^2, \delta_4^3, \delta_4^3\}$  satisfies  $\sigma_{X_1, Y_1^2}(\tilde{T}_1^1) = \sigma_{Z_2 \cup U_2, Z_2^1}(w_2^{3,2})$ . Then, the corresponding state trajectory is  $T_2^3 = \{\delta_4^4 \rightarrow \delta_4^4 \rightarrow \delta_4^3 \rightarrow \delta_4^1\}$ . In addition, similar to subnetwork  $\Sigma_2$ , there exists a unique control sequence  $w_3^{2,1} = \{\delta_4^2, \delta_4^4, \delta_4^1\} \in \Omega_3$  satisfying  $\sigma_{X_1, Y_1^3}(\tilde{T}_1^1) = \sigma_{Z_3 \cup U_3, Z_3^1}(w_3^{2,1})$ .

Then, we just need to consider the matchability between subnetworks  $\Sigma_2$  and  $\Sigma_3$ . Since  $\sigma_{X_2, Y_2^3}(\tilde{T}_2^3) = \sigma_{Z_3 \cup U_3, Z_3^2}(w_3^{2,1})$ , one can conclude that  $\{w_2^{3,2}, w_3^{2,1}\}$  is a 3-matchable control sequence. Therefore, BCN (E1) is 3-matchable.

In order to check the 3-matchable condition of BCN (E1), according to Definition 2, one needs to verify whether or not (9) holds for any  $i \in \{1, 2, 3\}$ ,  $j \in \{2, 3\}$ ,  $i \neq j$ . However, in this example, by virtue of the acyclic aggregation, one just needs to verify the cases of  $i = 1$ ,  $j = 2$ ,  $i = 1$ ,  $j = 3$ , and  $i = 2$ ,  $j = 3$ . Thus, acyclic aggregation can reduce the number of times for matchability when verifying the  $\kappa$ -matchable condition.

## Appendix E.5 An example used to show how Algorithm 1 works

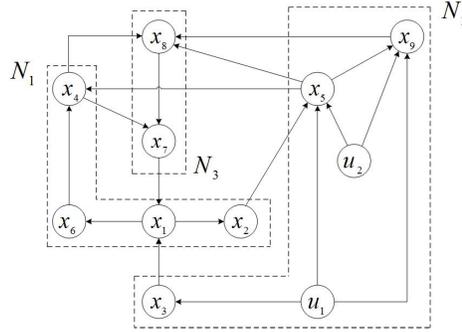
Consider the following Boolean model for the lac operon in Escherichia coli [4]:

$$\begin{cases} x_1(t+1) = \neg x_7(t) \wedge x_3(t), \\ x_2(t+1) = x_1(t), \\ x_3(t+1) = \neg u_1(t), \\ x_4(t+1) = x_5(t) \wedge x_6(t), \\ x_5(t+1) = \neg u_1(t) \wedge x_2(t) \wedge u_2(t), \\ x_6(t+1) = x_1(t), \\ x_7(t+1) = \neg x_4(t) \wedge \neg x_8(t), \\ x_8(t+1) = x_4(t) \vee x_5(t) \vee x_9(t), \\ x_9(t+1) = \neg u_1(t) \wedge (x_5(t) \wedge u_2(t)). \end{cases} \quad (\text{E3})$$

Fig. 4 shows an aggregation of BCN (E3). Denote the subnetwork corresponding to  $N_i$  by  $\Sigma_i$ ,  $i = 1, 2, 3$ . Given  $x^0 = \delta_{512}^{53}$ ,  $\kappa = 2$  and  $\mathbf{r} = (r_1, r_2, \dots, r_{512})^\top \in \mathbb{R}^{512}$ , where the element  $r_j$ ,  $j = 1, 2, \dots, 512$  in  $\mathbf{r}$  is given by the following function:

$$r_j = -j^2 + 9.8j - 14.$$

We solve this Mayer-type optimal control problem according to Algorithm 1.



**Fig. 4:** Aggregation of Boolean model (E3) for the lac operon in Escherichia coli.

Firstly, by Algorithm 1, setting  $i = 1$  and calculating  $x_1^d$  satisfying  $\mathbf{r}^\top x_1^d = \max\{r_j : j = 1, 2, \dots, 512\}$ , we can obtain  $x_1^d = \delta_{512}^{53}$ , where the lac operon is “on”.

Secondly, verify whether or not BCN (E3) is reachable from  $\delta_{512}^{53}$  to  $\delta_{512}^{53}$  at the second step by Theorem 1. On one hand, by a simple calculation, we can obtain that  $[M_i M_i(0)]_{\theta_i, \lambda_i} > 0$ ,  $i = 1, 2, 3$ . On the other hand, we can respectively find all the control sequences driving  $\alpha_i^0$  to  $\alpha_i^d$  at the second step,  $i = 1, 2, 3$  as  $\Omega_1 = \{\{\delta_8^4, \delta_8^2\}\}$ ,  $\Omega_2 = \{\{\delta_8^1, \delta_8^3\}, \{\delta_8^2, \delta_8^3\}, \{\delta_8^3, \delta_8^3\}, \{\delta_8^4, \delta_8^3\}\}$ , and  $\Omega_3 = \{\{\delta_8^7, \delta_8^1\}, \{\delta_8^7, \delta_8^2\}, \{\delta_8^7, \delta_8^3\}, \{\delta_8^7, \delta_8^4\}, \{\delta_8^7, \delta_8^5\}, \{\delta_8^7, \delta_8^6\}, \{\delta_8^7, \delta_8^7\}, \{\delta_8^7, \delta_8^8\}\}$ . Then, it can be verified by Definition 2 that the control sequence  $\{\{\delta_8^4, \delta_8^2\}, \{\delta_8^3, \delta_8^3\}, \{\delta_8^7, \delta_8^5\}\}$  is a 2-matchable control sequence, that is, BCN (E3) is 2-matchable. Therefore, condition (ii) of Theorem 1 holds. Therefore, BCN (E3) is reachable from  $x^0$  to  $x_1^d$  at the second step. In addition, the control sequence  $u_1^*$  steering BCN (E3) from  $x^0$  to  $x_1^d$  at the second step is  $\{u(0), u(1)\} = \{\delta_4^3, \delta_4^3\}$ .

By Theorem 2, the optimal control sequence is  $\{u(0), u(1)\} = \{\delta_4^3, \delta_4^3\}$ .

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