Diagnosability for Patterns in Distributed Discrete Event Systems

Yuhong Yan¹, Lina Ye², and Philippe Dague²

¹ CSE, Concordia University, Montreal, Canada yuhong@cse.concordia.ca
² LRI, Univ. Paris-Sud 11, France {lina.ye, philippe.dague}@lri.fr

ABSTRACT

A pattern is a Finite State Machine that can describe rich faulty scenarios, such as the occurrence of single faults, multiple faults, multiple occurrences of a fault, or the repair of a system. In distributed systems, the events in the pattern, as well as in the system trajectories, are emitted from different components. Our approach is based on distributed simulation and communication to check the recognition of the pattern from the conclusion of local recognition of local patterns. The components communicate observable events and shared communication events, as well as their local recognition results during the checking process without sharing their local models in any way.

1 INTRODUCTION

Discrete Event Systems (DESs) are widely used to describe system dynamics. Classically, a fault is associated with one non-observable faulty event. The fault diagnosis task in discrete event systems is to reason about the occurrence of a fault from a sequence of observed events emitted. If a finite sequence of observable events can non-ambiguously identify a fault from the other faulty or correct behaviors, this fault is called diagnosable.

A pattern is modeled as a Finite State Machine (FSM) with the events from the original system (Jéron et al., 2006). It can uniformly describe various faulty scenarios, such as single fault, multiple faults, multiple occurrences of a fault, and the repair of a system after the occurrence of a fault, etc. For example, a single fault can be modeled as a FSM that includes one transition that emits a faulty event; and a pattern for two faults can be modeled as an FSM that contains two branches from the initial state and each of the branches presents a scenario that one of the two faulty events precedes another respectively (Jéron et al., 2006). Indeed, a pattern is a unified description of rich faulty scenarios. Thus, studying pattern diagnosis and diagnosability is of great interest. A system can be diagnosed by deducing the recognition of the pattern from

a sequence of observed events. The purpose of pattern diagnosability is to answer whether this recognition is deterministic for all the trajectories with the same observations.

Our paper is to solve the pattern diagnosability problem in distributed discrete event systems. Our approach is based on distributed simulation, which is new in the research of distributed diagnosability analysis. We align local trajectories into partially observable global trajectories by distributed simulation. The global consistency is ensured via the simulation, instead of calculating a verifier (Pencolé, 2004). Further, we develop a process to compute globally observed traces and recognition of the pattern, and further compute the pattern diagnosability from the local recognition results. We have proved the correctness of our method. One advantage of our approach is that we avoid expensive operation to calculate local diagnosers (Pencolé, 2004; Ye *et al.*, 2009) and verifier (Pencolé, 2004). Another advantage of our approach is that the components do not share their local models in any direct or indirect way (e.g. sharing their local diagnosers (Pencolé, 2004)).

2 PRELIMINARIES

A distributed DES is composed of a finite set of components, $\{G_0, G_1, ..., G_n\}$, each of which is modeled as an FSM (Definition 1).

Definition 1 (Local Model). The local model of a component G_i is an FSM $G_i = (Q_i, \Sigma_i, \delta_i, q_i^0)$, where

- Q_i is a finite set of states
- Σ_i is a finite set of events
- $\delta_i \subseteq Q_i \times \Sigma_i \times Q_i$ is a set of transitions
- q_i^0 is the initial state

The set of events Σ_i is divided into three parts $\Sigma_i = \Sigma_{i_o} \biguplus \Sigma_{i_u} \biguplus \Sigma_{i_c}$, where Σ_{i_o} is the set of locally observable events which are disclosed to any other components; Σ_{i_u} is the set of locally unobservable events; and Σ_{i_c} is a set of communication events which are shared by G_i and at least one of the other components, and are observable only to their owners. As the locally observable events are disclosed to

the other components, they can be considered globally observable as well in this paper. We assume that an event can be labeled its owner's name. Therefore, for any pair of local components G_i and G_j , we have $\Sigma_{i_o} \cap \Sigma_{j_o} = \emptyset$ and $\Sigma_{i_u} \cap \Sigma_{j_u} = \emptyset$. Globally, we have $\Sigma_o = \biguplus_{i=1}^n \Sigma_{i_o}, \Sigma_u = \biguplus_{i=1}^n \Sigma_{i_u}$ and $\Sigma_c = \bigcup_{i=1}^n \Sigma_{i_c}$. For convenient, $\delta_i \subseteq Q_i \times \Sigma_i \times Q_i$ is also used to represent $\delta_i \subseteq Q_i \times \Sigma_i^* \times Q_i$ in the following way:

- $(q, \epsilon, q) \in \delta_i$, where ϵ is the null event
- $(q, se, q1) \in \delta_i$ if $\exists q' \in Q, (q, s, q') \in \delta_i$ and $(q', e, q1) \in \delta_i$, where $s \in \Sigma_i^*, e \in \Sigma_i$

We assume that the components do not share their local model with the other components and there is no central unit that knows all the local models. We further assume that the components do not synchronize their local clocks. That means the observed sequence of observable events from different components is not necessarily the same as their occurrence sequence. If one event from one component occurs before a communication event and another event from a second component occurs after the communication event, we can determine their sequences. Otherwise, we deduce the two events can occur concurrently.

Definition 2 (Synchronization). Given two FSMs $G_1 = (Q_1, \Sigma_1, \delta_1, q_1^0)$ and $G_2 = (Q_2, \Sigma_2, \delta_2, q_2^0)$, their synchronized product based on the communication events is $G_{1||2} = G_1||_{\Sigma_{1_c} \cap \Sigma_{2_c}} G_2 = (Q_1 \times Q_2, \Sigma_1 \cup \Sigma_2, \delta_{1||2}, (q_1^0, q_2^0))$, where $\delta_{1||2}$ is defined in the following:

- $\{((q_1, q_2), \sigma, (q_1', q_2'))\}$, if $\sigma \in \Sigma_{1_c} \cap \Sigma_{2_c}$ and $(q_1, \sigma, q_1') \in \delta_1, (q_2, \sigma, q_2') \in \delta_2$.
- {((q_1, q_2), σ , ($q_1\prime, q_2$))} if $\sigma \in \Sigma_1 \land \sigma \notin \Sigma_2$ and $(q_1, \sigma, q_1\prime) \in \delta_1$.
- $\{((q_1, q_2), \sigma, (q_1, q_2'))\}$ if $\sigma \notin \Sigma_1 \land \sigma \in \Sigma_2$ and $(q_2, \sigma, q_2') \in \delta_2$.
- otherwise $\delta_{1\parallel 2}$ is undefined.

Figure 1 depicts a distributed system with components $\{G_1, G_2\}$. For $G_1, \Sigma_1 = \{c1, c2, u1, u2, o1\}, \Sigma_{1_o} = \{o1\}, \Sigma_{1_u} = \{u1, u2\}, \Sigma_{1_c} = \{c1, c2\}. G_2$ has similar definitions. The global model of the system is $G = \{Q, \Sigma, \delta, q^0\} = G_1 || G_2$. We try to avoid to use this operation to get the global model due to its computational expensiveness and non-sharable local models.



Figure 1: A distributed system with two components $G_1(\text{left})$ and $G_2(\text{right})$.

The prefix-closed language L(G) generated by G describes the behavior of the system, which implies all the states in G are final states. We will use **trajectories** instead of words:

$$L(G) = \{ s \in \Sigma^* | \exists q \in Q, (q^0, s, q) \in \delta \}$$

In our paper, both the generated language and the recognized language are assumed to be *observable alive*, which means that there is no cycle containing only unobservable events. Given a trajectory $s \in L$, we denote L/s as the post-language of L after s and denote $P_{\Sigma_o}(s)$ as the sequence of observable events in s. $P_{\Sigma_o}(s)$ (P(s) when no ambiguity) is also called a **trace**. The inverse projection of a trajectory s is denoted as $P^{-1}(P(s)) = \{t \in L(G) | P(t) = P(s), s \in L(G)\}$.

3 PATTERN DIAGNOSABILITY

From domain knowledge, we can associate a pattern with the occurrence of single or multiple faults, or the repair of a system (Jéron *et al.*, 2006). A pattern can be described as an FSM (Definition 3).

Definition 3 (*Pattern*). A pattern is an FSM with final states set F_{Ω} , $\Omega = (Q_{\Omega}, \Sigma_{\Omega}, \delta_{\Omega}, q_{\Omega}^{0}, F_{\Omega})$, which satisfies the following conditions:

- $\forall q \in Q_{\Omega}, \forall \sigma \in \Sigma_{\Omega}, \text{ if } (q, \sigma, q_1) \in \delta_{\Omega} \text{ and } (q, \sigma, q_2) \in \delta_{\Omega}, \text{ then } q_1 = q_2.$
- $\forall q \in Q_{\Omega}, \Sigma_{\Omega}(q) = \Sigma_{\Omega} \text{ where } \Sigma_{\Omega}(q) = \{ \sigma \in \Sigma_{\Omega} | \exists q' \in Q_{\Omega}, (q, \sigma, q') \in \delta_{\Omega} \}.$
- $F_{\Omega} \subseteq Q_{\Omega}$ and $\delta_{\Omega}(F_{\Omega}, \Sigma_{\Omega}) \subseteq F_{\Omega}$ where $\delta_{\Omega}(F_{\Omega}, \Sigma_{\Omega}) = \bigcup_{q \in F_{\Omega}, \sigma \in \Sigma_{\Omega}} \{q' \in Q_{\Omega} | (q, \sigma, q') \in \delta_{\Omega} \}.$
- $q_{\Omega}^{0} \notin F_{\Omega}$

The first two conditions describe the pattern as a deterministic and complete finite state automaton. The third condition characterizes that the final state set F_{Ω} is stable. The set of all the trajectories that transfer the FSM to its final states is a recognized language of this FSM. Then it can be deduced that its recognized language $L_m(\Omega)$ is "extension-closed", formally described as

$$\forall s \in L_m(\Omega), \forall s' \in \Sigma^*_\Omega, ss' \in L(\Omega) \to ss' \in L_m(\Omega)$$

which means that once the system arrives in a final state, it will be always in a final state in the future. As Ω is complete, $ss' \in L(\Omega)$ is always true. Given a system FSM G and a pattern Ω , we assume: $\Sigma_{\Omega} = \Sigma$, $\Sigma_{\Omega_o} = \Sigma_o$. Notice that the FSM for pattern has final states, while the model for components does not have final states. It is because the components are assumed to work continuously, while a pattern has a final state to mark the recognition of a fault. Figure 2 presents a pattern Ω for the system displayed in Figure 1.



Figure 2: Pattern Ω .

A system can be diagnosed by deducing the recognition of the pattern from a sequence of observed events. Pattern diagnosability is to answer whether this recognition is deterministic for all the trajectories with the same observations. In addition, we want the recognition process to be bound, i.e. there must exist $n \in N$, where N is the set of nature numbers, that whenever $s \in L(G) \cap L_m(\Omega) \cap \Sigma^* \Sigma_o$, for all $t \in L(G)/s \cap \Sigma^* \Sigma_o$, the inverse projection of P(st)is also recognized by $L_m(\Omega)$. The term $\Sigma^* \Sigma_o$ constrains the trajectory to end with an observable event for the convenience of recognition. The following definition follows the bound diagnosability in (Jéron *et al.*, 2006):

Definition 4 (Pattern Diagnosability). A system FSM G is Ω -diagnosable if $\exists n \in N, \forall s \in L(G) \cap L_m(\Omega) \cap \Sigma^* \Sigma_o, \forall t \in L(G)/s \cap \Sigma^* \Sigma_o,$ if $|t| \ge n$, then $P^{-1}P(st) \subseteq L_m(\Omega)$.

Definition 4 says that if exists a nature number n, for any trajectory s satisfying $s \in L(G) \cap L_m(\Omega) \cap \Sigma^* \Sigma_o$, and any sequence of events t satisfying $t \in L(G)/s \cap$ $\Sigma^* \Sigma_o$, if t is longer than n, then the inverse projection of the projection of st is recognized by the pattern Ω . In another word, all the possible trajectories of what we can observe (which is the projection of st) is recongized by the pattern Ω , if we wait long enough. This means we can non-ambiguously decide that the pattern is recognized from what we have observed, if we waitn long enough. This is the exact meaning of Ω -diagnosability.

4 PATTERN DIAGNOSABILITY IN DISTRIBUTED DISCRETE EVENT SYSTEMS

4.1 The Principle

Following Definition 4, we need to find out whether all the trajectories with the same trace can be recognized by the pattern or not. If all the trajectories with the same trace can be recognized by the pattern (or if neither of the trajectories with the same trace can be recognized by the pattern), this means that from what we can observe (*i.e.* the trace), we can non-ambiguously tell the recognition of the pattern (or not). Therefore, the system is Ω -diagnosable. If some trajectories are recognized while the others are not, but they have the same trace, we could not tell wether the pattern is recognized or not. In this case, the system is not Ω diagnosable.

Theorem 1 If there exist two trajectories for a system which have the same trace, one of which is recognized by the pattern Ω and the other is not, the system is not Ω -diagnosable.

Proof: We use reductio ad absurdum. Assume two trajectories $s_1, s_2 \in L(G)$ have the same trace, *i.e.* $P(s_1) = P(s_2)$, and s_1 is recognized by Ω , and s_2 is not, *i.e.* $s_1 \in L_m(\Omega)$ and $s_2 \notin L_m(\Omega)$. If we compute all the trajectories whose trace is $P(s_1)$ (or $P(s_2)$), we have at least two trajectories s_1 and s_2 , *i.e.* $s_1, s_2 \in P^{-1}(P(s_1))$. If the system is Ω -diagnosable, we should have both *i.e.* $s_1 \in L_m(\Omega)$ and $s_2 \in L_m(\Omega)$ from Definition 4, which violates the

assumption that $s_2 \notin L_m(\Omega)$. Therefore, the system is not diagnosable.

Based on Theorem 1, if we want to prove a system is not Ω -diagnosable, we need to find out at least two trajectories with the same trace, one of which is recognizable by the pattern, and the other is not. If we want to prove a system is Ω -diagnosable, we verify all the trajectories such that their traces are the same, they should be uniformly recognized or not recognized by the pattern. In a distributed system, we use distributed simulation to identify globally consistent trajectories, and then determine if the trajectories with the same trace are uniformly recognized by the pattern or not. The two definitions below are used in the rest of the paper.

Definition 5 The critical transitions of a pattern are the transitions in the pattern whose starting and ending states are not the same. And the critical events are the events emitted by the critical transitions.

The critical transitions in Figure 2 are (p_0, u_1, p_1) , (p_0, o_3, p_3) , and (p_1, o_2, p_2) . The critical transitions are critical because they progress the process of recognizing a pattern. The critical events are $\{u_1, o_2, o_3\}$.

Definition 6 A local pattern Ω_i for a component G_i is an FSM modified from Ω by keeping only the critical transitions and renaming the local non-observable events which do not belong to G_i to a silent event ϵ .

For Ω in Figure 2, the local pattern Ω_1 for component G1 is the same as Ω except the looped transitions are removed (see also Fig. 3 left). The local pattern Ω_2 for component G2 renames u_1 to ϵ and it is without the looped transitions (see also Fig. 3 right).



Figure 3: The local patterns for G_1 (left) and G_2 (right).

4.2 Distributed Simulation

We assume that the components can simulate their local models and can send and receive messages to communicate their local observable events and communication events with one another. Via simulation, the components identify whether their local patterns are recognized, and then exchange the local recognition results. Each component can conclude the same result about the global diagnosability by synthesizing the local recognition results it receives. This diagnoability analysis is conducted off line, i.e. no observations at the run time are used. We first present some data structures and then the simulation algorithm.

An **Execution Tree** is a tree T with labeled edges to record the execution paths in a component. A tree node is labeled by a state of the local model. An edge is labeled by the event emitted from a transition t and a unique id. The unique id is $id_1 \in N$ at the first level and in the format of $id_{n-1}.k$ ($k \in N$) at the *n*-th level, where N is the set of nature numbers. With an id, we can easily retrieve the execution path that leads to the current transition. A method id = T.grows(t)grows the tree T with the given transition t and returns the generated *id*. A method T.getPath(id) returns an array of transitions on the execution path that leads to the transition labeled *id*. G_i .*id* is to tell the *id* belongs to a component G_i .

Communications and messages. A message m has a message type $m.type \in \{$ "communicate", "confirm", "loop"}. The message content for "com-municate" and "confirm" messages is a tuple $\langle e, G_i.id, G_j.id \rangle$, in which e is an event, G_i is the sender component, G_j is the receiver com-ponent. The message content for "loop" messages is $\langle G_i.id1, G_i.id2, G_j.id \rangle$, where the path between $G_i.id1$ and $G_i.id2$ is a loop. If any fields in the messages are unknown, null is $m = sends(type, \langle e, G_i.id, G_j.id \rangle)$ and used. $m = sends(type, \langle G_i.id1, G_i.id2, G_j.id \rangle)$ compose the message m and send it to G_i . G_i . id is the last transition *id* in the matched trajectory in G_i (cf. below). If no matching trajectory exists, G_j is used. The communication protocol is as following:

- 1. When a communication event is emitted, the component sends a "communicate" message to all components which share this communication events (possibly more than one), and waits for "confirm" messages from all its correspondents before resuming the execution on the branch. "Time-out" can remove any unsynchronizable path from the execution paths pool. A method b = m.matches(m'.e, id) matches a message m with a message m'. It returns 1 iff $m \cdot e == m' \cdot e$ $\wedge m.G_i.id \geq id$, otherwise $b = 0. m.G_i.id \geq id$ holds if the current *id* in the receiver (aka $\overline{G_i}$) is a subbranch of G_j .id. G_j is considered as a root. A path in a component can match multiple paths in another component. To reuse the already built paths, we keep all the confirmed messages in memory.
- 2. When an observable event is emitted, the component sends a "communicate" message to all the components, and does not wait for an answer before continuing its execution.
- 3. When a loop is detected, the component sends a "loop" message to all the components.

Matrix to record trajectories. A component records its execution trajectories and their matched observed traces from the other components in a matrix. A cell contains an event and *component.id*. For convenience, the rows are aligned by the correspondent communication events (ref. Table 1 and Table 2).

Distributed Simulation. In Algorithm 1, a local model $G_i = (Q_i, \Sigma_i, \delta_i, q_i^0)$ is simulated and the execution paths are recorded. The main While loop alternatively operates two methods *simulate()* and processMessages(), until all the paths are explored or some time-out conditions satisfied. The method *simulate()* simulates the local model by exploring all the possible execution paths. At each invocation, *simulate()* stops only when all the execution paths pause at the communication transitions waiting for

confirmations, or all the execution paths go into stable loops. *processMessages()* is to process received messages.

As the local model has finite states, the whole simulation process runs into a loop eventually. We can detect a loop by comparing the states on the execution path. If a loop has no communication events, we just need to inform all the components that this loop is detected. There is no need to repeat the execution of the loop. If a loop contains communication events, we need to execute the loop again until all the components are in a stable and synchronized loop. An execution path is removed if its communication event is not confirmed after the "time-out" threshold. Therefore, Algorithm 1 terminates. We neglect some details in Algorithm 1 for simplicity.

Algorithm 1 Distributed Simulation

 $MUP, MP = \emptyset$ - the set of unprocessed/processed received messages.

 $NUP, NP = \emptyset$ - the set of unprocessed/processed sent messages.

- 1: $A = \{q_i^0\}, \delta' = \emptyset$
- 2: while $A \neq \emptyset$ or !(time-out) do
- simulate() 3:
- 4: processMessages()

5:
$$A = \{\delta_i(q, e) | \delta_i \in \delta', \delta_i.pause = false\}$$

6:
$$\delta' = \delta' \setminus \{\delta_i | \delta_i \in \delta', \delta_i.pause = false\}$$

7: end while

1: simulate()

2: while $A \neq \emptyset$ do

= $\bigcup_{q\in A, e\in\Sigma} \delta(q, e, q'),$ A'3: δ _ $\bigcup_{q \in A, e \in \Sigma} \delta(q, e)$ 4: for each transition $\delta_i(q, e, q') \in \delta$ do 5: $id = T.grows(\delta_i), Matrix \leftarrow (e, G_i.id)$ if $e \in \Sigma_c$ then 6: 7: $id.pause = true, A' = A' \setminus q', \delta' \leftarrow \delta_i$ $\begin{array}{l} \begin{array}{l} \underset{m}{ = \ sends("communicate", \langle e, G_i.id, \\ G_j.id\rangle) \text{ to } \forall G_j \neq G_i, e \in G_j, \\ NUP \leftarrow m \end{array}$ 8: 9: 10: else 11: if $e \in \Sigma_o$ then 12: $\forall G_j$ \neq G_i , = msends("communicate", $\langle e, G_i.id,$ $G_j.id\rangle)$ end if 13: end if 14: {Detect loops} if $\exists \delta_i(q', e, q'') \in$ 15: $T.getPath(id) \wedge$ $\delta_i.loop = false$ then $id' = \delta_j(q', e, q'').id$ 16: $\forall G_j \neq G_i, m = sends(``loop", \langle G_i.id',$ 17: $G_i.id, G_j.id\rangle$) $E = \{e | e \in \delta_k, \delta_k \in (T.getPath)\}$ 18: $G_{i}.id$) – $T.getPath(G_{i}.id'))$ } mark the loop from id' to id in Matrix 19: if $E \cap \Sigma_c = \emptyset \lor$ all aligned trajectories in 20: loop then $\hat{A}' = A' \backslash q'$ 21: end if 22: 23: end if

24: end for

25: A = A'

26: end while

- 1: processMessages()
- for $\forall m \in MUP$ do 2:
- 3: if m.type == "communication" $\land m.e \in \Sigma_o$ then
- record $(m.e, m.G_i.id)$ at the end of a line for 4: G_j in Matrix where $G_j == G_i \land G_i.id \le G_j.id$

5:
$$MUP = MUP \setminus m, MP \leftarrow m^1$$

- 6: end if
- if m.type == "confirm" $\land \exists m' \in NUP$ that $m.match(m'.e,m'.G_i.id) == true$ then 7:
- align $(m.e, m.G_i.id)$ with $(m'.e, m'.G_i.id)$ 8: in Matrix, $MUP = MUP \setminus m, MP \leftarrow m$
- if all correspondents aligned to m' then 9: 10:

$$m'.id.pause = false$$

$$NUP = MUP \backslash m', NP \leftarrow m'$$

12: end if

11:

- end if 13:
- if m.type == "communication" $\land m.e \in \Sigma_c$ 14: then

15: for
$$\forall m' \in NUP$$
 that $m.match(m'.e, m'.G_{i.id}) == true^2$ do

16: align
$$(m.e, m.G_i.id)$$
 with $(m'.e, m'.G_i.id)$ in Matrix

 $m'' = send("confirm", \langle m.e, m'.G_i.id,$ 17: $m.G_i.id\rangle$)

18:
$$MUP = MUP \setminus m, MP \leftarrow m$$

19: if all correspondents aligned to m' then m'.id.pause = false20:

$$NUP = MUP \backslash m', NP \leftarrow m'$$

- 21: 22: end if
- 23: end for
- end if 24:
- 25: end for

Example 1 Given two local models G_1 and G_2 as in Figure 1, the execution matrix are those displayed in Table 1 and Table 2. The execution trees are shown in Figure 4 (a) and (b). Please notice that the trajectory $G_2.3$ has no matching trajectories in G_1 . Thus the last row in Table 1 contains only the observable events of $G_2.3.$ So does the last row in Table 2. \Box

Simulation Complexity. A local model G_i has maximally $|Q_i|^2 * |\Sigma_i|$ transitions. From the code, the first for loop on line 4 in simulate() is executed maximally $|Q_i|^2 * |\Sigma_i|$ times. We assume that the system is well synchronized and that, after n limited execution steps, the system is in a stable synchronized loop. This means the while loop on line 2 in simulate() is executed n times. Therefore, the time complexity of Algorithm 1 is $O(|Q_i|^2 * |\Sigma_i| * n)$. We have bounds for n in some simple cases. For example, when the model has only one loop at the "end" state (like in our example), n is the number of the transitions in the longest path between the initial state and the "end" state. Assume all the components are executed *n* times. The time complexity to simulate the whole system is $O(\sum_i |Q_i|^2 * |\Sigma_i| * n)$.



Figure 4: (a) The execution tree for component G_1 ; (b) The execution tree for component G_2 .

4.3 Checking Diagnosability

A component can use the following reasoning process to decide the diagnosability of the system. The conclusion by any component should be the same.

1. The aligned local trajectories are globally consistent. This is ensured by the design of the simulation process which follows the only constraint of the distributed system - synchronization by communication events. The aligned local trajectories can have sequential parts and concurrent parts. For example, Path $G_{1.1}$ and Path $G_2.1$ in Table 1 are paired and aligned by c_1 . The pair can be represented by $u1c1((u_2o_1^*)||(o_2o_3^*))$. The concurrent part joined by || indeed means that two FSMs synchronize without common events. The second trajectory in G_1 formed by Path $G_1.2$ and Path $G_2.2$ can be represented by $o_2c_2((u_1u_2o_1^*)||o_3^*)$. And third trajectory in G_1 is formed by Path G_2 .3 can be represented by $a_3o_3^*$. For G_2 , the trajectory made by $G_2.1$ and $G_1.1$ is $c_1((o_2o_3^*)||o_1^*)$; the trajectory made by $G_2.2$ and $G_1.2$ is $o_2c_2(o_3^*||o_1^*)$; and the trajectory made by $G_2.2$ and $G_1.2$ is $o_2c_2(o_3^*||o_1^*)$; and the trajectory made by $G_3.3$ is $o_3.o_3^*$.

2. Each component can compute the global traces. By projecting the aligned trajectories on observable events, a component can obtain global traces. For example, the group of traces for Path $G_{1.1}$ and Path $G_2.1$ is $o_1^*||(o_2o_3^*)$; for Path $G_1.2$ and Path $G_2.2$ is $o_2(o_1^*||o_3^*)$, and for Path $G_2.3$ is $o_3o_3^*$. G2 gets the same results.

3. Compute the recognizability of the patterns. First step: each component decides the recognizability of the local observed trajectories against its local pattern. We project each pair of trajectories on critical events to simplify the computation, because only the sequences of critical events decide recognizability. For G_1 , the three projected sequences for the three pairs of aligned trajectories are: $u_1 o_2 o_3^*$, $o_2(u_1 || o_3^*)$, and $o_3 o_3^*$. It is easy to see that the first and the third are recognized against local pattern Ω_1 , and the second is rejected. For G_2 , we have three sequences: $o_2 o_3^*$, $o_2 o_3^*$ and $o_3 o_3^*$. All are recognized against local pattern Ω_2 . Please notice that, if a trajectory has concurrent terms, if and only if all the possible paths are recognized by the pattern, can we consider the trajectory is recognized. For example, assume the pattern is $o_1 o_2$ and the trajectory is $o_1 || o_2$. This trajectory is considered non-recognizable, because one possible path $o_2 o_1$ is not recognizable by the pattern. The complexity to

¹reuse of *MP* and *NP* is eliminated

² if no match, a new path from the initial state may start.

Path $G_1.1$	$ \{u_1, G_1.1\}$	$\{c_1, G_1.1.1\}, \otimes$	$\{u_2, G_1.1.1.1\}$	$\{o_1, G_1, 1.1.1.1\}^*$	-	
Path $G_2.1$	-	$\{c_1, G_2.1\}, \otimes$	$\{o_2, G_2.1.1\},\$	$\{o_3, G_2.1.1.1\}^*$	-	\checkmark by G_1, G_2
Path $G_1.2$	-	$\{c_2, G_1.2\}, \otimes$	$\{u_1, G_1.2.1\}$	$\{u_2, G_1.2.1.1\}$	$\{o_1, G_1.2.1.1.1\}^*$	\times by G_1
Path $G_2.2$	$\{o_2, G_2.2\}$	$\{c_2, G_2.2.1\}, \otimes$	$\{o_3, G_2.2.1.1\}^*$	-	-	\checkmark by G_2
Path $G_2.3$	$\{o_3, G_2.3\}$	$\{o_3, G_2.3.1\}^*$	-	-	-	\checkmark by G_1, G_2

Table 1: Matrix of execution trajectories of G1. \otimes marks communication events, \checkmark marks recognized trajectories.

Path $G_2.1$	$\{c_1, G_2.1\}, \otimes$	$\{o_2, G_2.1.1\}$	$\{o_3, G_2.1.1.1\}^*$	\checkmark by G_1, G_2
Path $G_1.1$	$\{c_1, G_1.1.1\}, \otimes$	${o_1, G_1.1.1.1.1}^*$	-	
Path $G_2.2$	$\{o_2, G_2.2\}$	$\{c_2, G_2.2.1\}, \otimes$	$\{o_3, G_2.2.1.1\}^*$	$ imes$ by G_1
Path $G_1.2$	-	$\{c_2, G_1.2\}, \otimes$	$\{o_1, G_1.2.1.1.1\}^*$	\checkmark by G_2
Path $G_2.3$	$\{o_3, G_2.3\}$	$\{o_3, G_2.3.1\}^*$	-	\checkmark by G_1, G_2

Table 2: Matrix of execution trajectories of G2. \otimes marks communication events, \checkmark marks recognized trajectories.

compute local recognition is discussed at the end of this section.

Second step: if and only if all the components unanimously vote for recognizing a pair of aligned trajectories using their local patterns, this pair of trajectories is recognized by the global pattern. That local recognition implies global recognition is ensured by Theorem 2. That a local rejection can cause a global rejection is easy to understand. If a global trajectory is recognized by the global pattern, its projection on one component's events can also satisfy this component's local pattern. Therefore, if the projection of a global trajectory on one component's events does not satisfy the component's local pattern, this global trajectory does not satisfy the global pattern either.

For our example, the pair of Path G_1 .1 and Path G_2 .1 is recognized globally; the pair of Path G_1 .2 and Path G_2 .2 is not recognized globally; and Path G_2 .3 is recognized globally.

4. Decide diagnosability. From Theorem 1, if there exist both recognized and unrecognized trajectories of the same trace, the system is not diagnosable, otherwise it is diagnosable. For our example, we find that o_2 is a possible trace for the pair composed by Path $G_1.1$ and Path $G_2.1$ and the pair composed by and Path $G_2.1$ and Path $G_2.2$. As the pair of Path $G_1.1$ and Path $G_2.1$ are recognized by the global pattern and the pair Path $G_1.2$ and Path $G_2.2$ are not, the system is not diagnosable. As we check all the possible trajectories, our decision is sound. The complexity to check the existence of a common trace for trajectories is discussed at the end of this section.

Our reasoning process relies on a proposition that the local recognition of the local patterns implies global recognition of the global pattern. It is easy to tell the inverse implication stands. As a local pattern is less constrained than the global pattern, if a local pattern is not recognized, the global pattern cannot be recognized. But there are cases that local recognition does not imply global recognition. Theorem 2 gives a sufficient condition that except the case that two consecutive critical transitions whose events are locally unobservable events and belong to two different components, local recognition implies global recognition.

Theorem 2 If there exist no two consecutive critical transitions in the pattern whose events are locally unobservable events and belong to different components,

the locally recognized trajectories against the local patterns are globally recognized against the global pattern.

Proof: Without losing generality, assume t_1 and t_2 are two consecutive critical transitions in the pattern and whose events are α and β respectively. $t_1 \prec t_2$ denotes that t_1 precedes t_2 .

If α and β belong to the same component, it is a trivial case, because one component can fully determine their sequence from the simulation algorithm.

If $\alpha \in G_1, \beta \in G_2$ belong to components G_1 and G_2 respectively. By the condition of the theorem, α and β cannot be both locally unobservable events. Therefore, α and β can be both locally observable, or one is locally unobservable and the other is locally observable. Thus, at least one of G_1 and G_2 has both α and β in its local pattern. Therefore, if this component can determine their sequences locally through the simulation algorithm, this means globally their sequence is determined. Thus, the local recognition leads to the global recognition.

Theorem 2 tells an exceptional case that if there are two consecutive critical transitions whose events are locally non-observable and belong to different components in the pattern, local recognition does not always imply global recognition. In this case, we cannot apply the methods developed in this paper. Figure 5 illustrates this exceptional case. In Figure 5(a), the pattern has two consecutive critical transitions whose events are u_1 and u_2 which belong to G_1 and G_2 respectively. Since one component can only simulate either u_1 or u_2 , there is no way to determine their sequence by either of the components. For example, Figure 5(c) and (d) illustrate two components G_1 and G_2 . Their local patterns are $(p_0, u_1, p_1) \rightarrow (p_1, o_1, p_3)$ and $(p_0, u_2, p_2) \rightarrow (p_2, o_1, p_3)$ respectively.

By distributed simulation, G_1 gets one trajectory $u_1c_1(o_1||o_2)$ and G_2 gets one trajectory $u_2c_1(o_2||o_1)$. It is easy to see that both G_1 and G_2 recognize their local patterns. However, globally, we can only deduce u_1 and u_2 are parallel, *i.e.* $u_1||u_2$, because they both precede the communication point c_1 . There is no way to distinguish their sequential order. This can also be proved by synchronizing the local models of G_1 and G_2 to get the global system model. In this case, the local recognition.

If we change the pattern to Figure 5(b), where

 u_1 and u_2 are separated by an observable event o_1 , the local patterns to G_1 and G_2 are $(p_0, u_1, p_1) \rightarrow (p_1, o_1, p_3)$ and $(p_0, o_1, p_2) \rightarrow (p_2, u_2, p_3)$ respectively. G_1 can get $u_1c_1(o_1||o_2)$ by simulation, and it recognizes its local pattern. G_2 can get $u_2c_1(o_1||o_2)$ by simulation, and it does not recognize its local pattern. Therefore, globally, the pattern is not recognized. This is the case that satisfies the condition of Theorem 2. We can see that the sequence orders of $u_1 \prec o_1$ and $o_1 \prec u_2$ in the pattern can be determined by one component. Thus local recognition implies global recognition. However, our example is a case that local non-recognition implies global non-recognition.



Figure 5: Two consecutive local events in a pattern.

Briefly, we discuss when a local component can determine the occurrence sequence of two critical events α and β emitted by two consecutive critical transitions in a pattern. Here we do not constrain the observability of α and β . If α and β belong to the same component, then this component can determine their occurrence sequence by simulation. If α and β belong to two different components G_1 and G_2 respectively, their occurrence sequence cannot be determined by their observed sequence. However, if there exists at least a communication event c shared by G_1 and G_2 , and $\alpha \prec c$, $c \prec \beta$, we can deduce that $\alpha \prec \beta$. This is illustrated as in Table 3. This helps us to check local recognition easily.

Path $G_{1.1}$	α	\rightsquigarrow	$c \otimes$		
Path $G_2.1$			$c \otimes$	$\sim \rightarrow$	β

Table 3: Matrix of execution trajectories of G1 and G2.

Complexity of Checking Diagnosability. The steps one and two in the diagnosability check reasoning process are trivial, which involve trajectories alignment and projection.

The third step needs to compute whether a pair of trajectories can be recognized by the local pattern. We first project the trajectories on critical events. In a general case, the projected trajectories have concurrent segments. Each projected trajectory can be represented as a regular expression. (Thompson, 1968) converts a regular expression E with n letters in length into a Nondeterministic Finite Automaton (NFA) with n states in linear time 1(n). The synchronized NFA II has $m = n1 \cdot n2$ states. Therefore, we need to consider whether $L(\Pi) \subseteq L(\Omega_i)$, where Ω_i is a local pattern which is a Deterministic Finite Automaton (DFA). It is known that it takes $O(n1 \cdot n2)$ time to check $L_1 \subseteq L_2$, for L_1 and L_2 are DFA with n_1 and n_2 states (Gelade and Neven, 2008). However, to convert a NFA II to a

DFA takes an exponential time 2^m . Therefore, the total complexity is $O(2^m \cdot p)$, where p is the number of states in the pattern. Normally a pattern Ω is relatively simple and critical events are not many. We can think that m is much smaller than the length of trajectories.

The fourth step needs to compute the existence of common traces for two pairs of trajectories when one pair is recognized globally and the other is not. We first project the trajectories on observable events. Then we can get two synchronized NFAs Π_1 and Π_2 as above. We need to tell whether $L(\Pi_1) \cap L(\Pi_2) = \emptyset$. This needs $O(m_1 \cdot m_2)$ time complexity (Gelade and Neven, 2008).

5 RELATED WORK

(Sampath et al., 1995) introduced the diagnosability problem for discrete event systems and proposed to solve it by detecting some transition cycles of ambiguous states in a global diagnoser. The main disadvantage of their approach is its exponential space complexity in the number of system states. (Jiang et al., 2001) proposed a classical twin plant approach to improve the algorithm complexity, which is only polynomial in the number of system states. (Pencolé, 2004) studied the diagnosability problem in distributed systems and provided a non scalable method of synchronizing local non reduced twin plants until a global critical path is detected. Then (Schumann and Pencolé, 2007) proposed a scalable approach for diagnosability verification in a distributed way through checking the existence of a set of local reduced twin plants, where at least one of them contains an observable possibly nondiagnosable cycle. On the other hand, (Jéron et al., 2006) extended the diagnosability problem for fault events to that for supervision patterns, which can be used to describe more general objectives. They verified pattern diagnosability by employing a global twin plant method. (Ye et al., 2009) is the first paper to discuss pattern diagnosability in distributed systems based on the model in (Jéron et al., 2006). Their study limited to simple patterns which contain only one linear branch leading to the final state. Compared to (Ye et al., 2009), our paper starts with the same model as in (Jéron et al., 2006), but without any constraints on patterns. Thus, the result is more general than (Ye et al., 2009). (Guillou et al., 2008) studies diagnosis of discrete event system with a formalism called chronicle (Dousson et al., 1993). A chronicle is a set of events and temporary constraints between those events. It is more abstract than pattern represented by FSM. (Guillou et al., 2008) extends the classic chronicle to include variables for modeling data flow and to include synchronization points for using in distributed systems. In (Guillou et al., 2008), faulty scenarios are modelled in the variables, while in our paper, the faults are presented by the patterns (or the FSM). In contrast to our paper, (Guillou et al., 2008) studies the problem of diagnosis, instead of diagnosability. In addition, (Guillou et al., 2008) uses local diagnosers and global diagnoser, while in our system we do not use any central node to synthesize local results.

6 CONCLUSIONS

We study pattern diagnosability in distributed discrete event systems. We present the principle of pattern diagnosability (Theorem 1). For a distributed system, we use distributed simulation to identify globally consistent trajectories, and then determine if the trajectories with the same trace are uniformly recognized by the pattern or not. We present the correctness proof (Theorem 2) and complexity analysis.

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