

# MODELING LARGE SCALE MANUFACTURING PROCESS FROM TIMED DATA

## *Using the TOM4L Approach and Sequence Alignment Information for Modeling STMicroelectronics' Production Processes*

Pamela Viale<sup>1,2</sup>, Nabil Benayadi<sup>1</sup>, Marc Le Goc<sup>1</sup> and Jacques Pinaton<sup>2</sup>

<sup>1</sup>LSIS, Laboratory for Information and System Sciences, University of Marseille, Marseille, France

<sup>2</sup>STMicroelectronics, Rousset, France

**Keywords:** Process model discovery, Temporal knowledge discovering, Markov processes, Sequence alignment.

**Abstract:** Modeling manufacturing process of complex products like electronic chips is crucial to maximize the quality of the production. The Process Mining methods developed since a decade aims at modeling such manufacturing process from the timed messages contained in the database of the supervision system of this process. Such process can be complex making difficult to apply the usual Process Mining algorithms. This paper proposes to apply the TOM4L Approach (Timed Observations Mined for Learning) to model large scale manufacturing processes. A series of timed messages is considered as a sequence of class occurrences and is represented with a Markov chain from which models are deduced with an abductive reasoning. Because sequences can be very long, a notion of process phase is introduced. Sequences are cut based on the concept of class of equivalence and information obtained from an appropriate alignment between the sequences considered. A model for each phase can then be locally produced. The model of the whole manufacturing process is obtained from the concatenation of the models of the different phases. This paper presents the application of this method to model STMicroelectronics' manufacturing processes. STMicroelectronics' interest in modeling its manufacturing processes is based on the necessity to detect the discrepancies between the real processes and experts' definitions of them.

## 1 INTRODUCTION

<sup>1</sup> Modeling manufacturing processes for the construction of complex objects like electronic chips is crucial for maximizing productivity. The methods and the algorithms developed in the Process Mining domain aims at modeling such manufacturing processes with graphs of manufacturing steps (treatments, operations or tasks) from the timed messages contained in the database (Cook and Wolf, 1998). The proposed algorithms generally generate complex models that are difficult to read and interpret when the process contains hundred of steps.

This paper proposes a modeling method and the corresponding algorithms to model manufacturing processes having hundreds of steps. The proposed method is based on cutting the sequences in subsequences called *process phases*. The way of cutting

the sequences is based on information obtained from an alignment between them. However, finding an appropriate sequence alignment is not a simple task.

The section 2 resumes the state of art on the process mining area and it introduces some ideas about sequence alignment. Section 3 defines the notions of *class of equivalence* and *process phase* that we propose and describes the algorithms that are required to this aim. Section 4 presents the application of the algorithms to model STMicroelectronics' manufacturing processes. These manufacturing processes are used for creating electronics chips, micro-controllers, etc. The resulting models will be useful to control that the implemented process correspond to the experts' ideas. Discrepancies found will be helpful to alarm experts' about possible problems in production. In section 5 a summary of the proposed method is presented. The paper concludes in section 6 with the introduction of our current works.

<sup>1</sup>This work has been made possible thanks to the financial support of STMicroelectronics, Convention 642/2008.

## 2 STATE OF ART

### 2.1 Process Mining Area

In the Process Mining framework, a series of messages is considered as an ordered set of events from where a process model can be inferred and represented in some formalism (workflows, state charts or Petri nets for examples) (van der Aalst and Weijters, 2004). One of the first algorithm was proposed in (Agrawal et al., 1998). The algorithm aims at finding workflow graphs from a set of series of events contained in a workflow log. An event represents the start time of a task. To avoid the problem of potential cycles (i.e. repeated events in a series), the algorithm first renames the repeated labels of task before enumerating the binary dependency relations between the tasks. This set of relations is then reduced with the use of the transitivity property of the binary relations. Labels are again renamed to merge the tasks, making possible the introduction of cycles in the model. Difficulties arise with this approach when (i) the tasks are statistically independent and (ii) the number of tasks is large (Agrawal et al., 1998). Nevertheless, Pinter (Pinter and Golani, 2004) extends this algorithm notably with the introduction of events marking the end of the tasks. Similar issues in the context of software engineering processes are investigated in (Cook and Wolf, 1998) where the aim is to build a finite state machine from the set of the most frequent event patterns mined in a given log. In particular, the *Markov* algorithm is based on a two order Markov chain that is converted in states and state transitions. Cook and Wolf (Cook and Wolf, 2004) extend this method to concurrent processes and uses a first order Markov chain for this aim. The difficulties come from the pruning of the finite state machine to obtain a minimal model and the sensibility of pruning metrics to the "noise" (van der Aalst and Weijters, 2004). Aalst (van der Aalst et al., 2004) defines the class of process that can be modeled with the  $\alpha$ -algorithm but this algorithm requires the series of events in the log to be noise-free and complete.

There is a consensus to consider that finite state machines are difficult to understand and to validate. And most of the proposed methods have difficulties when (i) the process contains a lot of steps, (ii) the series in the log induce potential cycles in the models and (iii) the sequences are not noise-free and complete. The *TOM4L Approach* (Timed Observations Mined for Learning, previously called Stochastic Approach Framework) (Le Goc et al., 2005) for discovering temporal knowledge from timed observations provides a general framework for modeling dynamic

processes that is based on a markovian representation but uses abstract chronicle models (Ghallab, 1996) instead of finite state machines. This framework considers that the timed messages of a series are written in a database by a program, called a monitoring cognitive agent *MCA*, that monitors a production process *Pr*. A timed message is represented with an occurrence of a discrete event class  $C^i = \{e_i\}$  that is an arbitrary set of discrete event  $e_i = (x_i, \delta_i)$ , where  $\delta_i$  is one of the discrete value of the variable  $x_i$ . When the variable  $x_i$  is not known, an abstract variable  $\phi_i$  is used to define the discrete event  $e_i = (\phi_i, \delta_i)$  corresponding to the constant  $\delta_i$ . A discrete event class is often a singleton because in that case, two discrete event classes  $C^i = \{(x_i, \delta_i)\}$  and  $C^j = \{(x_j, \delta_j)\}$  are only linked with the variables  $x_i$  and  $x_j$  when the constants  $\delta_i$  and  $\delta_j$  are independent (Le Goc, 2006). This condition is only concerned with the programs the *MCA* is made with. A sequence of discrete event class occurrences is then considered as the observable manifestation of a series of state transitions in a timed stochastic automaton representing the couple  $(Pr, MCA)$ . The *BJT4G* algorithm represents a set of sequences of discrete event class occurrences with a one order Markov chain and uses an abductive reasoning to identify the set of the most probable timed sequential binary relations between discrete event classes leading to a given class. A timed sequential binary relation  $R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+])$  is an oriented relation between two discrete event classes  $C^i$  and  $C^j$  that is timed constrained with the interval  $[\tau_{ij}^-, \tau_{ij}^+]$ .  $[\tau_{i,j}^-, \tau_{i,j}^+]$  is the time interval for observing an occurrence of the  $C^j$  class after an occurrence of the  $C^i$  class. The set of timed sequential binary relation is an abstract chronicles model where the nodes are discrete event classes and the links are timed sequential binary relations. This paper proposes to tackle the two main problems of the Process Mining approaches with the extension of the TOM4L Approach. The first ideas of this approach has been presented in (Benayadi et al., 2008).

### 2.2 Sequence Alignment

Before introducing the extension to the TOM4L Approach, it is necessary to introduce some ideas about sequence alignment. These ideas are necessary to understand the steps proposed in the algorithms for modeling large scale manufacturing processes.

A sequence alignment consists in a way of arranging sequences to identify regions of similarity between them. Aligned sequences are usually represented as rows within a matrix. Gaps ('-') are inserted between the residues so that identical or simi-

lar residues are aligned in successive columns. In our problem, the residues are the discret event class occurrences.

One of the first issue that arise when comparing sequences concerns the estimation of substitution costs. These substitution costs provide useful information that leads the algorithms to align or not the residues of the sequences considered. The scores for aligning the different characters in the alphabet  $\Sigma$  are obtained from a matrix  $S = [s_{x,y}]_{x \in \Sigma, y \in \Sigma}$ , called similarity matrix (it is also called substitution matrix). The value  $s_{x,y}$  represents the similarity between the elements of the pair  $(x,y), x \in \Sigma, y \in \Sigma$ . In biology, there exists some well-known substitution matrices. However, this problem has also been encountered in many other areas, for example, in social sciences where there are no well known similarity matrices precalculated. The problem in social science is to compare chronological sequences of various kind (e.g. professional or familial statuses) where there is no equivalent to such a strong theoretical and empirical framework like evolutionary theory. Thus the relationship between the symbols constituting sequences of social events remain at least partly uncertain. The ideas from social science field can guide us towards to find a strategy to build similarity matrices for any kind of sequences. The idea is to find a way to construct the similarity matrix  $S$  for any kind of sequence, deriving costs empirically. An adaptation of the algorithm for calculating substitution costs iteratively is necessary (Gauthier et al., 2008).

### 3 EXTENSION OF THE TOM4L APPROACH FOR PROCESS MINING

#### 3.1 Motivation

Let us take an example to illustrate the proposed extensions with a manufacturing process having a set  $S = \{A, B, C, D, E, F\}$  of 6 manufacturing steps. Suppose the supervision system records the execution of a step with a message  $X(t_k)$  denoting the time  $t_k$  of the beginning of the execution of the step  $X$ . The three series of messages of table 1 are represented with the abstract chronicle model of figure 1. In this model, if nodes labeled with  $A$  denote the same manufacturing step, then the nodes can be confused, introducing a cycle in the model. The same reasoning can be done over the other nodes, making the model difficult to read and to understand.

This repetition of manufacturing steps can be due

Table 1: Three series of event.

$A(t_1)B(t_2)F(t_3)B(t_4)E(t_5)D(t_6)C(t_7)A(t_8)E(t_9)B(t_{10})$
$A(t_{11})B(t_{12})E(t_{13})C(t_{14})D(t_{15})A(t_{16})E(t_{17})B(t_{18})$
$A(t_{19})B(t_{20})E(t_{21})C(t_{22})D(t_{23})A(t_{24})E(t_{25})B(t_{26})$

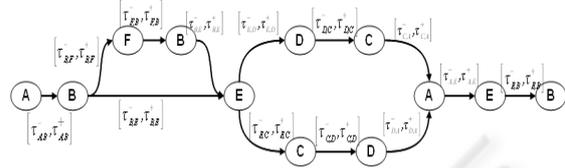


Figure 1: Model for the three Sequences.

to a situation that arrives frequently during the execution of a manufacturing process. When an object is being manufactured, the object goes from one machine to another for making all the necessary treatments on it. Nevertheless, usual controls in production sometimes show that certain objects did not achieved the expected characteristics. Some actions (special actions) has to be done to correct the problem. These actions usually consist on performing special treatments and/or the repetition of tasks.

This is the reason why we need to align the different process execution sequences to identify equivalent treatments over the different objects being produced in each execution.

Given a set  $\Omega = \{\omega_i\}_{i=\{1, \dots, n\}}$  of sequences obtained from the execution of the same manufacturing process, we need a 'good' sequence alignment between them,  $SA(\Omega)$ . By 'good' we mean an alignment that aligns similar treatments made on the different objects on the same column and special tasks (task performed to correct errors) with gaps ('-').

Two possible 'good' alignment for the sequences shown in Table 1 are proposed in Table 2.

Suppose now that each of the three sequences are cut when a label is aligned with exactly the same one in all sequences and this alignment appears two times. The model of the first part of the sequences will be similar to the one of figure 1 but without the path  $A - E - B$  at the end of the model. This fact motivates the notion of *process phase* proposed in this paper. But this notion is not sufficient to solve the cycles that are introduced with the steps  $C$  and  $D$ . We consider that this problem is due to the fact that the three series of events provide no information about the order of the steps  $C$  and  $D$ . Consequently, any solution of this problem must take into account some *a priori* knowledge about the process which we want to avoid it. The notion of *potential cycle* is then defined to detect this kind of situation to be able to make further investigations (i.e. finding new series or discussing

Table 2: Two Sequence Alignments proposed for sequences on Table 1.

Possible Sequence Alignments	
$SA_1 =$	$\left( \begin{array}{cccccccccccc} A(t_1) & B(t_2) & F(t_3) & B(t_4) & E(t_5) & D(t_6) & C(t_7) & - & A(t_8) & E(t_9) & B(t_{10}) \\ A(t_{11}) & B(t_{12}) & - & - & E(t_{13}) & - & C(t_{14}) & D(t_{15}) & A(t_{16}) & E(t_{17}) & B(t_{18}) \\ A(t_{19}) & B(t_{20}) & - & - & E(t_{21}) & - & C(t_{22}) & D(t_{23}) & A(t_{24}) & E(t_{25}) & B(t_{26}) \end{array} \right)$
$SA_2 =$	$\left( \begin{array}{cccccccccccc} A(t_1) & B(t_2) & F(t_3) & B(t_4) & E(t_5) & - & D(t_6) & C(t_7) & A(t_8) & E(t_9) & B(t_{10}) \\ A(t_{11}) & B(t_{12}) & - & - & E(t_{13}) & C(t_{14}) & D(t_{15}) & - & A(t_{16}) & E(t_{17}) & B(t_{18}) \\ A(t_{19}) & B(t_{20}) & - & - & E(t_{21}) & C(t_{22}) & D(t_{23}) & - & A(t_{24}) & E(t_{25}) & B(t_{26}) \end{array} \right)$

with experts for examples).

To illustrate the notion of class of equivalence, let us take the *Edit* activity of the "writing a scientific paper" process that can be made by different students and professors. The *Edit* activities can then be labeled differently according to the performer with a set of classes of the form:  $C = \{C^{E1} = \{(s_1, \delta_1)\}, C^{E2} = \{(s_2, \delta_2)\}, \dots\}$ , where the variables  $s_i$  and  $p_i$  denotes respectively students and professors. In this case, the resulting model of the process will be complex without necessity. One of the interesting features of the TOM4L Approach is the notion of discrete event class. This notion can be used to define abstract classes of the form  $C^{\phi_i} = \{(\phi_i, \delta_1), \dots, (\phi_i, \delta_n)\}$  where  $\phi_i$  denotes an abstract variable and the set  $\{\delta_j\}, j = 1 \dots n$ , is an arbitrary set of constants. This property allows defining classes of equivalence that simplifies a process model. For example, an abstract class  $C^{\phi_i}$  be defined as an equivalent class of the set of classes  $C$  of the "writing a scientific paper" process. Doing this way allows constituting a set of sequences coming from different students and professors.

### 3.2 Class of Equivalence

By definition, a large scale process consist on a lot of steps. Some of these steps differs only with some characteristics but realizes similar treatments.

**Definition 1.** Given a model  $M = \{R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+])\}$  build with a set  $\Omega = \{\omega_i\}$  of discrete event class occurrences  $\omega_i$ , a class  $C^{\phi_i} = \{(\phi_i, \delta_1), (\phi_i, \delta_2), \dots, (\phi_i, \delta_n)\}$  is an equivalence class of a sub set of classes  $C = \{C^j\}$ ,  $j = 1 \dots n$ ,  $C^j = \{(x_j, \delta_j)\}$ , of the set of classes  $C_M$  of a model  $M$  iff:

$$\begin{aligned}
 & \forall C^a, C^b \in C \\
 & \forall C^k \in C_M, \\
 & (\exists R(C^k, C^a, [\tau_{ka}^-, \tau_{ka}^+]) \in M \wedge \exists R(C^k, C^b, [\tau_{kb}^-, \tau_{kb}^+]) \in M) \\
 & \quad \wedge \\
 & \forall C^r \in C_M, \\
 & (\exists R(C^a, C^r, [\tau_{ar}^-, \tau_{ar}^+]) \in M \wedge \exists R(C^b, C^r, [\tau_{br}^-, \tau_{br}^+]) \in M)
 \end{aligned} \tag{1}$$

This definition means that every classes  $C^j$  of the subset  $C \subseteq C_M$  are linked with the same classes

in  $M$ . When this condition is verified, each occurrences  $C^j(t_s)$  of the classes  $C^j$  in the sequences  $\omega_i$  of  $\Omega$  can be rewritten as occurrences  $C^{\phi_i}(t_s)$  of the equivalence class  $C^{\phi_i}$ . The abstract variable  $\phi_i$  has no *a priori* meaning:  $\phi_i$  can be substituted with the corresponding concrete variable  $x_i$  in any occurrences  $C^{\phi_i}(t_s)$ . Consequently, the set of uphill relations  $\{R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+])\}$  of  $M$  will become  $\{R(C^i, C^{\phi_i}, [\tau_{i\phi_i}^-, \tau_{i\phi_i}^+])\}$  and the set of downhill relations  $\{R(C^j, C^k, [\tau_{jk}^-, \tau_{jk}^+])\}$  of  $M$  will become  $\{R(C^{\phi_i}, C^k, [\tau_{\phi_i k}^-, \tau_{\phi_i k}^+])\}$ . In practice, an equivalence class can be used to represent related discrete event classes. In the application presented in the next section, a discrete event class represents a set of treatments that can be made on an object using a particular machine. Equivalence classes are then used to represent this set of treatments made by different machines: in that case, the machines are equivalents because the same set of treatments can be done on each of the machines.

Given a set of sequences  $\Omega = \{\omega_i\}_{i=\{1, \dots, n\}}$ , the algorithm for defining equivalence classes find all the equivalence classes and rewrite the corresponding occurrences in each sequence  $\omega_i$  (Algorithm 1):

1. Build a model  $M$  given a set of sequences  $\Omega = \{\omega_i\}_{i=\{1, \dots, n\}}$ .
2. Find all the subset of classes  $C$  verifying the equation 1.
3. For all the subsets  $C$ .
  - Create an equivalence class  $C^{\phi_i}$ .
  - For all  $C^j \in C$ , rewrite all the occurrences  $C^j(t_s)$  in all the sequences  $\omega_i \in \Omega$  with the rewriting rule:  $C^j(t_s) \equiv C^{\phi_i}(t_s)$ .
4. Build a new model  $M'$  with  $\Omega$ .

### 3.3 Process Phase

The information contained in a series of manufacturing messages is concerned with both the state of the manufactured object and the manufacturing process that makes evolving this state from an initial state up to a final state. But generally, the state of the manufactured object is not provided with the messages. So we

propose to capture indirectly this dimension with the notion of *Process Phase*. However, before introducing this new concept, we must define formally what is called a *Sequence Alignment*.

### 3.3.1 Sequence Alignment

A sequence alignment consists in a way of arranging sequences to identify regions of similarity between them. Aligned sequences are usually represented as rows within a matrix  $M = [m_{i,j}]_{i \in \{1, \dots, n\}, j \in \{1, \dots, r\}}$ . Gaps ('-') are inserted between the residues so that identical or similar residues are aligned in successive columns.

**Definition 2.** *Given a model  $M = \{R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+])\}$  build with a set  $\Omega = \{\omega_i\}$  of discrete event class occurrences. A sequence alignment  $SA$  between all sequences  $\omega_i \in \Omega$  is defined as follows:*

$$\begin{aligned} SA(\Omega) = \{ & M_{n \times r} | \\ & \forall \omega_j \in \Omega, r \geq \text{card}(\omega_j) \wedge \\ & \forall m_{ij} \in M_{n \times r}, m_{ij} \in C_M \cup \{-\} \wedge \\ & \forall m_{ij}, m_{i(j+1)} \in M_{n \times r}, \\ & (m_{ij} = C^j(t_k), C^j(t_k) \in \omega_i \wedge \\ & m_{i(j+1)} = C^l(t_x), C^l(t_x) \in \omega_i \wedge \\ & \quad t_x > t_k) \} \end{aligned} \quad (2)$$

where:

- $\text{card}(s)$  is the length of sequence  $s$ .

There are many possible alignments between  $n$  sequences. There exists many algorithms for creating sequence alignments based on different objectives and strategies (Thompson et al., 1994; Notredame et al., 2000; Needleman and Wunsch, 1970; Smith and Waterman, 1981). However, it is necessary to provide a similarity matrix  $S$  to these algorithms so that they are able to calculate when to align or not residues of the different sequences. If we use the same alignment algorithm over the same set of sequences with different similarity matrices the resulting alignments will be probably not the same. So, it is very important to use a similarity matrix that agrees with the semantics of the sequences treated. We will illustrate this problem in section 4, over STMicroelectronics' manufacturing process sequences.

Given a set of sequences  $\Omega = \{\omega_i\}_{i=1, \dots, n}$  and a sequence alignment  $SA(\Omega)$ , we propose an algorithm for renaming classes. This algorithm traverses all sequences and renames a class if in the alignment it is aligned with classes different from itself or with gaps (Algorithm 2):

- $\forall \omega_i \in \Omega, \forall C^j(t_k) \in \omega_i$  do:

- Rename the occurrence  $C^j(t_k)$  when:
  - $\exists C^i(t_s) \in \omega_m, \omega_m \in \Omega, C^j(t_k)$  is aligned with  $C^i(t_s)$  in  $SA(\Omega)$  and  $C^i \neq C^j$ ; or
  - $C^j(t_k)$  is aligned with a gap ('-') in  $SA(\Omega)$ .

In Table 3 we will show the corresponding series from Table 1 after applying Algorithm 2 on them, considering sequence alignment  $SA_1$ .

### 3.3.2 Process Phase Concept

As it was said before, the concept of process phase will be used to capture the different stages that an object must go through in the manufacturing process to achieve the final expected state. For this task we need to count with a 'good' sequence alignment of the sequences that we use to build the model. The reason is that this alignment provides a guideline to distinguish normal treatments from special tasks performed in particular executions of the process. Aligned tasks are supposed to be regular tasks whereas not aligned tasks are not.

**Definition 3.** *Given a set of sequences  $\Omega = \{\omega_i\}_{i=1, \dots, n}$  and a sequence alignment  $SA(\Omega)$ , a process phase of the model  $M$  constructed using Algorithm 2 is a submodel  $M' = \{R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+]) | C^i, C^j \in C_{M'}\} \subseteq M$ , so that there is no path  $P = \{R(C^i, C^{i+1}, [\tau_{i+1}^-, \tau_{i+1}^+])\} \in M', i = 1 \dots n$ , where:*

$$\forall j, k \in \mathbb{N}, 1 \leq j < k \leq n, C^j = C^k \quad (3)$$

Algorithm 3 aims at cutting a set  $\Omega = \{\omega_i\}_{i=1, \dots, n}$  of sequences in subsequences  $\omega_k^i$  that respects the equation 3 (i.e.  $\omega_k^i$  does not contain two occurrences of the same class):

1.  $\forall \omega_i \in \Omega$  do
  - Remove  $\omega_i$  from  $\Omega$ .
  - Cut up  $\omega_i$  in a set  $\Omega_i = \{\omega_k^i\}$  of sub sequences  $\omega_k^i$  verifying the equation 3.
2.  $\forall \omega_k^i \in \Omega_i$  do
  - Add an occurrence of the  $C^0$  and  $C^1$  classes at the beginning and the end of  $\omega_k^i$ .
3.  $\Omega = \bigcup \Omega_i$ .

An occurrence of an abstract start class  $C^0$  and an occurrence of an abstract final class  $C^1$  are added at the beginning and the end of each sub sequences  $\omega_k^i$  so that the *BJT4G* algorithm automatically identifies the process phases. For example, when applying the algorithm 3 on the three sequences of Table 1, the *BJT4G* algorithm will find two process phases. The second process phase

Table 3: Series of event after renaming with Algorithm 2.

$A(t_1)$	$B(t_2)$	$F_{1,3}(t_3)$	$B_{1,4}(t_4)$	$E(t_5)$	$D_{1,6}(t_6)$	$C(t_7)$	$A(t_8)$	$E(t_9)$	$B(t_{10})$	
$A(t_{11})$	$B(t_{12})$			$E(t_{13})$		$C(t_{14})$	$D_{2,15}(t_{15})$	$A(t_{16})$	$E(t_{17})$	$B(t_{18})$
$A(t_{19})$	$B(t_{20})$			$E(t_{21})$		$C(t_{22})$	$D_{3,23}(t_{23})$	$A(t_{24})$	$E(t_{25})$	$B(t_{26})$

will be:  $\{R(C^0, C^A, [\tau_{0A}^-, \tau_{0A}^+]), R(C^A, C^E, [\tau_{AE}^-, \tau_{AE}^+]), R(C^E, C^B, [\tau_{EB}^-, \tau_{EB}^+]), R(C^B, C^1, [\tau_{B1}^-, \tau_{B1}^+])\}$ .

It can be easily deduced from the definition of process phase that process phases are dependent on the sequence alignment chosen. It is necessary to count with a 'good' sequence alignment for being able to correctly identify the different stages in the process.

### 3.4 Potential Cycles

When looking the model of figure 1, it is clear that the classes  $C$  and  $D$  introduce a cycle. Cycles present a strong problem of interpretation, making hard to understand the resulting models. This explains why there is a lot of works aiming at avoiding cycles in process models (cf. (Cook and Wolf, 2004), (Schimm, 2004) (van der Aalst et al., 2004), (Pinter and Golani, 2004), (Weijters and van der Aalst, 2003) or (Agrawal et al., 1998) for examples). But these works make assumptions about the process or impose constraints about the constitution of the sequences. In all the case, this consists in having some *a priori* knowledge about the process to be modeled or the set of programs that write the messages in the process data base.

The aim of the TOM4L Approach is to provide models of sequences without any *a priori* knowledge about the process and the set of programs that have generated the occurrences. One difficulty is that cycles often appear when mining a process because of the transitivity property of the sequential binary relations.

**Property 1.** *The timed sequential binary relations  $R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+])$  of a given abstract chronicle model  $M = \{R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+])\}$  are transitives.*

$$\begin{aligned} \forall R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+]) \in M \wedge \forall R(C^j, C^k, [\tau_{jk}^-, \tau_{jk}^+]) \in M \\ R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+]) \wedge R(C^j, C^k, [\tau_{jk}^-, \tau_{jk}^+]) \\ \Rightarrow \exists R^T(C^i, C^k, [\tau_{ik}^-, \tau_{ik}^+]) \end{aligned} \quad (4)$$

**Definition 4.** *Given a process model  $M$ , two discrete event classes  $C^i$  and  $C^j$  are not ordered when:*

$$M \vdash R^T(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+]) \wedge M \vdash R^T(C^j, C^i, [\tau_{ji}^-, \tau_{ji}^+]) \quad (5)$$

Two classes  $C^i$  and  $C^j$  that can not be ordered in a model will be denoted  $C^i \parallel C^j \equiv C^j \parallel C^i$ .

For examples the three sequences of the Table 1 do not provide any order between the classes  $C^C$  and  $C^D$  (Figure 1). Consequently:  $C^C \parallel C^D \equiv C^C \parallel C^D$ .

**Property 2.** *The set of discrete event classes  $C^{\parallel} = \{C^1, C^2, \dots, C^n\}$  can not be ordered when:*

$$\forall C^i \in C^{\parallel}, \forall C^j \in C^{\parallel}, C^i \parallel C^j. \quad (6)$$

In the theory of graphs, the classes of a set  $C^{\parallel}$  are strongly connected components. The algorithm 4 aims at detecting a potential cycle (i.e. a set  $C^{\parallel}$  is defined):

1. Build a process model  $M$  from  $\Omega$  with the *BJT4G* algorithm.
2. Build the set  $C^{\parallel} = \{C_i^{\parallel}\}$  of the sets  $C_i^{\parallel}$  of classes without order with the equation 5
3.  $\forall C_i^{\parallel} \in C^{\parallel}$  do
  - Remove the relations  $R(C^i, C^j, [\tau_{ij}^-, \tau_{ij}^+])$  of  $M$  where  $C^i \in C_i^{\parallel}$  or  $C^j \in C_i^{\parallel}$ .
  - Generate all the pathes  $P = \{P_k\}$  with  $P_k = \{R(C^i, C^{i+1}, [\tau_{ii+1}^-, \tau_{ii+1}^+])\}$  where  $C^i \in C_i^{\parallel}$  and  $C^{i+1} \in C_i^{\parallel}$ .
  - Insert the relations of the pathes  $P$  in  $M$ .

To avoid the adding of *a priori* knowledge about the process or the programs, the algorithm 4 computes all the paths linking the classes in  $C^{\parallel}$  (cf. model of Figure 1 with the classes  $C$  and  $D$ ). It is clear that if  $Card(C^{\parallel}) = n$ , there is  $n!$  possible paths. But it is a simple way to put the emphasis on potential cycles.

### 3.5 Modeling a Large Scale Process

The algorithm 5 aims at modeling a large scale manufacturing process. It simply uses the four algorithms provided in the preceding subsections. Given a set of sequences  $\Omega = \{\omega_i\}_{i=\{1, \dots, n\}}$  and a sequence alignment  $SA(\Omega)$ , the algorithm 5 finds a process model  $M$  with the *BJT4G* algorithm:

1. Rewrite the sequences of  $\Omega$  with the algorithm 1.
2. Rewrite the sequences of  $\Omega$  with the algorithm 2 using information from  $SA(\Omega)$ .
3. Produce the sets  $\Omega_k$  of subsequences  $\omega_k^i$  with the algorithm 3.
4.  $\forall \Omega_k$  do
  - Build a process model  $M_k$  of the phase  $k$  with the algorithm 4.
5.  $M = \bigcup M_k$

Applied to the sequences of the Table 1, this algorithm provides the model of the Figure 1. This algorithm has also been used to model the wafer manufacturing process of the Rousset (France) plant of the STMicroelectronics company.

## 4 APPLICATION

The aim of STMicroelectronics Company is to improve the control of the wafer manufacturing processes through the definition of human scale process models and a better knowledge of the timed constraints between the different steps of manufacturing. A "wafer" is a silicon plate used for the construction of STMicroelectronics' products. A wafer manufacturing process is a series of elementary treatments called "recipes" that are executed on machines called "equipments". An "operation" is a particular serie of recipes and each recipe is associated with different equipments, those that are qualified for performing the treatment. A complete series of operations is called "manufacturing route". The STMicroelectronics' plant situated in the south of France, in Rousset, counts with more than 10.000 different recipes, each manufacturing route is composed of about 400 operations and there are actually more than 300 equipments in the plant. The supervision system of the wafer manufacturing process describes a manufacturing route with messages providing the name of a recipe, the machine on which the recipe is executed, the corresponding operation and the start and finish times of the recipe.

For validating the approach presented in this paper, a set  $\Omega$  containing 5 sequences  $\omega_i$ ,  $i = \{1, \dots, 5\}$ , has been extracted from STMicroelectronics' databases. These sequences contain data from real production of the Company. For the construction of this example we considered an extract of each of these 5 sequences  $\omega_i$ ,  $i = \{1, \dots, 5\}$ . Only the 200 first executed recipes were used.

The Algorithm 5 has been applied at the equipment level so that a process model  $M$  represents a manufacturing route by a serie of equipments. Even though we worked with extracts of real routes, we found a large volume of data. In the example constructed, 127 discret event classes have been found. A class is defined with a singleton  $C^i = \{(\phi_i, i)\}$  where the constant  $i$  is a natural number in the interval  $[1000, \dots, 1126]$ . Each of these classes represent a particular equipment in STMicroelectronics' plant.

We will illustrate the application of the approach proposed in this paper showing how it works over two subsequences of sequences  $\omega_1$  and  $\omega_2$  used for the

example construction. These two subsequences can be seen in Figure 2.

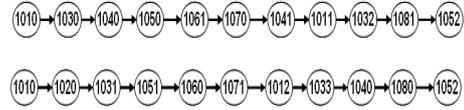


Figure 2: Subsequence of  $\omega_1$  (up) and subsequence of  $\omega_2$  (down).

During the example construction, 67 different equivalent classes have been found. The equivalent classes created using Algorithm 1 are singletons of the form  $C^j = \{(\phi_j, j)\}$  with  $j \in [6000, 6066]$ . Consider the equivalent class  $C^{6008} = \{(\phi_{6008}, 6008)\}$  contains the discret event classes  $\{C^{1031}, C^{1032}, C^{1033}\}$  and that  $\{C^{1010}, C^{1011}, C^{1012}\} \subseteq C^{6009}$ ,  $\{C^{1050}, C^{1051}, C^{1052}\} \subseteq C^{6031}$ ,  $\{C^{1040}, C^{1041}\} \subseteq C^{6040}$ ,  $\{C^{1020}\} \subseteq C^{6042}$ ,  $\{C^{1060}, C^{1061}\} \subseteq C^{6046}$ ,  $\{C^{1080}, C^{1081}\} \subseteq C^{6056}$ ,  $\{C^{1070}, C^{1071}\} \subseteq C^{6065}$ . The algorithm 1 rewrites the two subsequences of Figure 2 and produces the subsequences of Figure 3.

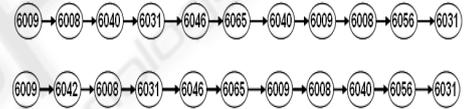


Figure 3: Subsequences of  $\omega_1$  and  $\omega_2$  rewritten with Algorithm 1.

Suppose now that we construct alignments between the two subsequences of  $\omega_1$  and  $\omega_2$ , using an optimal algorithm (Needleman and Wunsch, 1970) and two different similarity matrices  $S_1$  and  $S_2$  shown in table 4. The main difference between these two matrices is that  $S_2$  penalizes the alignment of different residues whereas  $S_1$  does not. A '-1' penalty is used for the introduction of gaps.

The alignment corresponding to the two subsequences using the similarity matrix  $S_1$  is shown in Figure 4. The alignment for similarity matrix  $S_2$  can be seen in Figure 5.

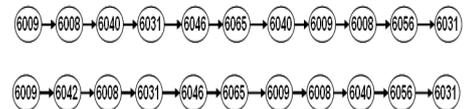


Figure 4: Subsequences of  $\omega_1$  and  $\omega_2$  aligned using similarity matrix  $S_1$ .

After analysing the alignment with STMicroelectronics' experts we could corroborate that only the alignment constructed using matrix  $S_2$  respects the semantics of the sequences. This tiny example shows

Table 4: Two possible similarity matrices for residues of subsequences of  $\omega_1$  and  $\omega_2$  considered.

Similarity matrices		
$S_1 =$	$\begin{pmatrix} - & 6008 & 6009 & 6031 & 6040 & 6042 & 6046 & 6056 & 6065 \\ 6008 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 6009 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 6031 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 6040 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 6042 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 6046 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 \\ 6056 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ 6065 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \end{pmatrix}$	
	$S_2 =$	$\begin{pmatrix} - & 6008 & 6009 & 6031 & 6040 & 6042 & 6046 & 6056 & 6065 \\ 6008 & 1 & -1 & -1 & -1 & -1 & -1 & -1 & -1 \\ 6009 & -1 & 1 & -1 & -1 & -1 & -1 & -1 & -1 \\ 6031 & -1 & -1 & 1 & -1 & -1 & -1 & -1 & -1 \\ 6040 & -1 & -1 & -1 & 1 & -1 & -1 & -1 & -1 \\ 6042 & -1 & -1 & -1 & -1 & 1 & -1 & -1 & -1 \\ 6046 & -1 & -1 & -1 & -1 & -1 & 1 & -1 & -1 \\ 6056 & -1 & -1 & -1 & -1 & -1 & -1 & 1 & -1 \\ 6065 & -1 & -1 & -1 & -1 & -1 & -1 & -1 & 1 \end{pmatrix}$

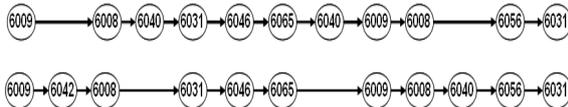


Figure 5: Subsequences of  $\omega_1$  and  $\omega_2$  aligned using similarity matrix  $S_2$ .

the importance of choosing the right matrix to calculate the alignment between all sequences considered. An error in this step will introduce a problem that will be carried over the following steps. Working with a wrong sequence alignment will probably conduce to errors when cutting sequences in process phases.

After applying Algorithm 2 for renaming classes according to the alignment shown in Figure 5, the subsequences of  $\omega_1$  and  $\omega_2$  are rewritten as it is shown in Figure 6.



Figure 6: Subsequences of  $\omega_1$  and  $\omega_2$  rewritten using Algorithm 2.

Algorithm 3 finds two process phases ( $i$  and  $i + 1$ ) for the subsequence considered of  $\omega_1$  and for the subsequence considered of  $\omega_2$ . Process phase  $i$  can be seen in Figure 7 and process phase  $i + 1$  is shown in Figure 8.

Using Algorithm 4 over the phases found, the corresponding models were constructed. Following the two subsequences used for illustrating the approach, we construct the models shown in Figures 9 and 10.

During the construction of the hole example

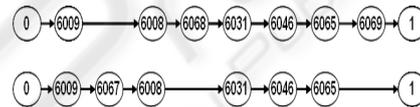


Figure 7:  $\omega_1^i$  (up) and  $\omega_2^i$  (down).

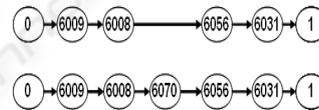


Figure 8:  $\omega_1^{i+1}$  (up) and  $\omega_2^{i+1}$  (down).

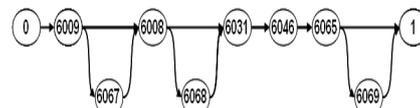


Figure 9: Model for phase  $i$  obtained from  $\omega_1^i$  and  $\omega_2^i$ .

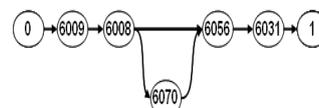


Figure 10: Model for phase  $i + 1$  obtained from  $\omega_1^{i+1}$  and  $\omega_2^{i+1}$ .

model, 19 process phases were found. The resulting model contains a total of 302 nodes.

In Figure 11 the resulting model constructed for the 5th phase found is shown. The volume of data in this small example is large even though we worked with extracts of manufacturing routes. The model obtained using this method is, however, simple and easy to understand. The resulting model has already been

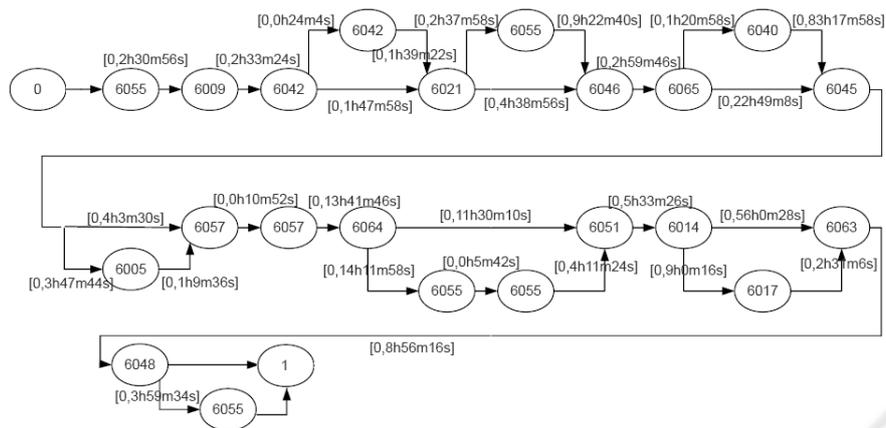


Figure 11: Phase 5 of the model.

validated by STMicroelectronics' experts.

The next step is to build a model of for complete sequences and for bigger sets (more than 5 sequences). Afterwards, it would be valuable to construct models for other levels of granularity (level of operations or recipe level are some options).

## 5 CONCLUSIONS

This paper presents the TOM4L Approach for modeling manufacturing processes from timed data contained in a supervision system database. One of the interesting features of this approach is the notion of discrete event class. This notion is used to define a process phase concept and discrete event classes of equivalence that are required for large scale manufacturing processes. The definition of these concepts leads to a global algorithm that has been applied to modeling STMicroelectronics' (Rousset) manufacturing processes. This concrete application shows the operational flavor of the extensions of the TOM4L Approach. The importance of calculating a similarity matrix that corresponds to sequence semantics has also been shown in the applicative section. The construction of these models will be a valuable tool for STMicroelectronics to control production and to alarm experts when the real activity of the Company does not follow their theoretical definitions.

## 6 CURRENT WORKS

Current works are devoted to find an analogy of the work presented in (Gauthier et al., 2008), works done on the social science field. Our idea is to find a way

to calculate the similarity values between different events classes in any kind of sequence, without introducing experts knowledge about their contents.

## REFERENCES

- Agrawal, R., Gunopulos, D., and Leymann, F. (1998). Mining process models from workflow logs. *In Sixth International Conference on Extending Database Technology*, pages 469–483.
- Benayadi, N., Le Goc, M., and Bouché, P. (2008). Using the stochastic approach framework to model large scale manufacturing processes. *Proceedings of the 3rd International Conference on Software and Data Technologies (ICSoft 2008)*.
- Cook, E. J. and Wolf, A. L. (1998). Discovering models of software processes from event-based data. *ACM Transactions on Software Engineering and Methodology*, 7:215–249.
- Cook, E. J. and Wolf, A. L. (2004). Event-based detection of concurrency. *In Proceedings of the 6th ACM SIGSOFT international symposium on Foundations of software engineering*, volume 53, pages 35–45.
- Gauthier, J. A., Widmer, E. D., Bucher, P., and Notredame, C. (2008). How much does it cost? optimization of costs in sequence analysis of social science data. *Sociological Methods and Research*.
- Ghallab, M. (1996). On chronicles: Representation, on-line recognition and learning. *Proc. Principles of Knowledge Representation and Reasoning, Aiello, Doyle and Shapiro (Eds.) Morgan-Kaufman*, pages 597–606.
- Le Goc, M. (2006). *Notion d'observation pour le diagnostic des processus dynamiques: Application à Sachem et à la découverte de connaissances temporelles*. Hdr, Faculté des Sciences et Techniques de Saint Jérôme.
- Le Goc, M., Bouché, P., and Giambiasi, N. (2005). Stochastic modeling of continuous time discrete event se-

- quence for diagnosis. *16th International Workshop on Principles of Diagnosis (DX'05)*, California, USA.
- Needleman, S. B. and Wunsch, C. D. (1970). A general method applicable to the search of similarities in the amino acid sequence of two proteins. *J. Mol. Biol.*, 48:443–453.
- Notredame, C., Higgins, D. G., and Heringa, J. (2000). T-coffee: A novel method for fast and accurate multiple sequence alignment. *J. Mol. Biol.*, 302:205–217.
- Pinter, S. and Golani, M. (2004). Discovering workflow models from activities' lifespans. In *Special issue: Process/workflow mining*, volume 53, pages 283–296.
- Schimm, G. (2004). Mining exact models of concurrent workflows. In *Computers in Industry*, volume 53(3), pages 265–281.
- Smith, T. F. and Waterman, M. S. (1981). Identification of common molecular subsequences. *J. Mol. Biol.*, 147:195–197.
- Thompson, J. D., Higgins, D. G., and Gibson, T. J. (1994). Clustal w: improving the sensitivity of progressive multiple sequence alignment through sequence weighting, position-specific gap penalties and weight matrix choice. *Nucleic Acid Research*, 22:4673–4680.
- van der Aalst, W., Weijters, T., and Maruster, L. (2004). Workflow mining: Discovering process models from event logs. In *IEEE Transactions on Knowledge and Data Engineering*, volume 16, pages 1128–1142.
- van der Aalst, W. M. P. and Weijters, A. J. M. M. (2004). Process mining. *Special issue of Computers in Industry*, 53:231–244.
- Weijters, A. and van der Aalst, W. (2003). Rediscovering workflow models from event-based data using little thumb. In *Integrated Computer-Aided Engineering*, volume 10(2), pages 151–162.