Aligning Observed and Modeled Behavior

An overview of alignments, current research and future challenges

by

Boudewijn van Dongen

Eindhoven University of Technology
Process Mining

The goal of process mining is to extract gain insights into business processes by analyzing operational behavior captured in event logs, through:

- Discovering processes
  - How do people behave?
- Compliance oriented
  - Where and why do people deviate?
- Performance oriented
  - Where are bottlenecks in my processes?
Deviations in Business Processes

The commonality between a process manual and a concrete barrier: it’s been there for years and everybody walks around it.

#olifantenpaadjes
Outline

• Introduction to alignments
• Computing alignments
• Active research
• Online alignments
• Conclusions
• Future challenges
Typical Deviations

Unnecessary deviations

Necessary deviations

Interfering deviations

Illegal deviations
Alignments: A way to explain deviations

An alignment contains the most likely execution of the model, corresponding to an observed execution.

An alignment shows where deviations occurred and why these deviations are considered as such.
Computing Alignments

• The search space is a “product” of the statespace of the model and the trace

• Each node is a combination of a state in the model and the remaining events in the trace

• Each arc is a move on model, move on log or a synchronous move

• A heuristic function estimates the remaining distance to the final node (i.e. model in a final state and all events executed)

• We find a shortest path!
Computing Alignments (A*)

Initialize PriorityQueue q
While (peek(q) is not target t)
    VisitedNode n = head(q)
    add n to the considered nodes
    For each edge in the graph from node(n) to m
        If m was considered before, continue
        If m is in the queue with lower cost, continue
        If m is in the queue with higher cost, update and reposition it
        If m is new,
            compute an estimate for the remaining distance to t
            V = new VisitedNode(m)
            set n a predecessor for v
            add v to the priority queue
    Return head(q)
Computing Alignments (A* with fast lowerbound)

Initialize PriorityQueue q
While (peek(q) is not target t)
    VisitedNode n = head(q)
    If the estimate for node(n) is not exact
        Compute the exact estimate for the remaining distance to t and
        add n to the priority queue
        continue
    add n to the considered nodes
    For each edge in the graph from node(n) to m
        If m was considered before, continue
        If m is in the queue with lower cost, continue
        If m is in the queue with higher cost, update and reposition it
        If m is new,
            compute a fast lowerbound for the remaining distance to t
            v = new VisitedNode(m)
            set n a predecessor for v
            add v to the priority queue
    Return head(q)
Estimating remaining distance

• Naïve estimation: 0 Trivial
• Smarter estimation: trace Trivial
• LP-based estimation: “Polynomial”
  • Minimize $c \cdot x$
    Where $A \cdot x = r$
    $c \cdot x \geq c \cdot x'$
• ILP-based estimation: Exponential
• Hybrid ILP (1 sec for “I” part) Exponential
• Caching LP basis solutions Exponential

Little to no difference in practice
Implementations: Estimator Versions

**Petri nets:**
- Dijkstra (estimator 0)
- Naive (estimator parikh)
- ILP (estimator using ILP)
- Basis caching
- Fast lowerbounds

**Process Trees:**
- Naive (estimator parikh)
- LP (estimator using LP)
- Hybrid ILP (ILP & LP)
- Special constraints for OR
- Fast lowerbounds
- Basis caching within LP
- Statespace reduction (stubborn sets)
Time vs. Space

- **Time per trace**
- **Unique states**
- **Queued states**
- **Computed Estimates**

### Computing alignments

- **No ILP**
  - Time in seconds: 1,071,585
  - Time per trace: 0.03064
  - Queued states: 57,181
  - Unique states: 49,464
  - Computed Estimates: 46,812

- **LP**
  - Time in seconds: 59,723
  - Time per trace: 0.01031
  - Queued states: 58,330
  - Unique states: 46,812
  - Computed Estimates: 46,812

- **ILP**
  - Time in seconds: 57,181
  - Time per trace: 0.00901
  - Queued states: 58,327
  - Unique states: 48,077
  - Computed Estimates: 46,368

- **HLP**
  - Time in seconds: 58,330
  - Time per trace: 0.00913
  - Queued states: 58,327
  - Unique states: 48,077
  - Computed Estimates: 46,368

- **HLP C**
  - Time in seconds: 58,327
  - Time per trace: 0.00914
  - Queued states: 58,327
  - Unique states: 48,077
  - Computed Estimates: 46,368

- **HLP C S**
  - Time in seconds: 49,464
  - Time per trace: 0.00725
  - Queued states: 46,812
  - Unique states: 46,812
  - Computed Estimates: 46,812

- **HCLP C S O**
  - Time in seconds: 46,812
  - Time per trace: 0.00716
  - Queued states: 48,077
  - Unique states: 48,077
  - Computed Estimates: 48,077

- **LP S O**
  - Time in seconds: 48,077
  - Time per trace: 0.00812
  - Queued states: 48,077
  - Unique states: 48,077
  - Computed Estimates: 48,077

- **ILP S O**
  - Time in seconds: 46,368
  - Time per trace: 0.00702
  - Queued states: 46,368
  - Unique states: 46,368
  - Computed Estimates: 46,368
Time vs. fitness

Computing optimal alignments is a time-intensive task
Active research on alignments I

- Conformance checking foundations
  - Fitness (Arya Adriansyah)
  - Precision (Jorge Munoz)

- Process discovery
  - Genetic algorithms (PhD work Joos Buijs)
  - Model Repair (Dirk Fahland, Artem Polyvyany)
  - Alignments without models (Xixi Lu)

- Automated compliance checking
  - Needs $n$-to-$m$ mappings for activities / transitions (Elham Rhamezani)
Active research on alignments II

• Data/Resource aware alignments
  • Include data into the statespace (Felix Mannhardt)
  • First control-flow, then data (Massimiliano De Leoni)
  • Multi perspective

• Process model animation
  • Animate models based on alignments (Sander Leemans)

• Online Alignments
  • Requires “constant” time and memory (Andrea Burattin, me)
Alignments – Some Insights

Consider the model and trace on the right

\[\text{Online alignments}\]
Online conformance checking

Requirements:

• Compute alignments without knowledge of the future
• Use constant time per event
• Use constant space

For this, we introduce a representation of alignments, such that between each pair of synchronous moves:

• Log moves are done first,
• Model moves are postponed until strictly necessary
Aligning process models – search space

• Aligning process models requires traversing a state space which is the synchronous product of the model’s state space and the trace.
Aligning process models – search space

< A C B A >
Aligning process models – search space

\(<A\ C\ B\ A>\)
Aligning process models – search space

Online alignments
Properties of the shortest path

For a given label A, there is a limited number of shortest paths that end in a synchronous move on some activity A between a state S and another state S′.
Redefined search space

• For each state $s$ and label $A$ compute the shortest paths ending in a synchronous move on some activity labelled $A$.
• The maximum number of model moves allowed in this path is limited by a given parameter “depth”

To execute an activity $D$ in the state after $A$ was executed, we need a sequence of model moves consisting of $<C>$ with length 1.

Another possibility is a path $<C,D,C>$ with length 3, hence this is not the shortest sequence.
Online conformance checking

• Suppose we have an oracle that produces, given a state S and an label A some reachable states $S'$ trough a series of model moves followed by a synchronous move on $A$, and the cost of reaching $S'$

• For each trace, we keep (an ordered list of) partial alignment(s) for the events in the trace we have observed so far

• For each new event, we update all partial alignments using the oracle
Online conformance checking - example

- For each event, we build a partial alignment.
- Between two synchronous moves, there’s always first log moves, then model moves (if any).
- We may need to keep more than one partial alignment.
- We may even need to keep sub-optimal partial alignments.

On the image:

- The diagram illustrates the process of online conformance checking with partial alignments.
- The notation `<A,D,E>` indicates a sequence of events where `A`, `D`, and `E` are aligned.
- The alignment process involves aligning events from the log and the model to ensure conformance.

In the diagram:

- The symbols `A`, `B`, `C`, `D`, `E` represent events.
- The process involves aligning these events to check for conformance.
- The partial alignments are shown in green for the log and yellow for the model.
Online conformance checking - parameters

• The oracle needs a *depth* parameter, indicating the maximum number of model moves allowed to get to a synchronous move.

• The replayer needs to keep a number of possible partial alignments in a *queue* (sorted by cost).

• Per event, the oracle needs to look through the search space in a depth first manner, until maximum depth is reached. Hence the time per event is bounded by a constant (given the model).

• The number of partial alignments to keep per trace is bounded by the size of the statespace.
Online conformance checking - guarantees

Given sufficient depth and sufficient queue size:

• Each optimal partial alignment gives a lower bound on the total alignment cost.
• The final alignment (after ending the trace) is an optimal alignment.
• We need constant time and space, as long as the actual alignment is not required, but only the cost of deviating.
• If the actual alignment is needed, the space required scales linearly in the length of the trace.
An example

• Maximum depth: 11

• Statespace:
  • 82,916 nodes
  • 719,521 edges

• Log:
  • 500 random traces
  • 16,446 events
  • Fitness 0.61
An example

<table>
<thead>
<tr>
<th>Depth</th>
<th>Queue</th>
<th>Time</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>82916</td>
<td>28 s</td>
</tr>
<tr>
<td>1</td>
<td>82916</td>
<td>44 s</td>
</tr>
<tr>
<td>2</td>
<td>82916</td>
<td>64 s</td>
</tr>
<tr>
<td>3</td>
<td>82916</td>
<td>94 s</td>
</tr>
<tr>
<td>5</td>
<td>1935</td>
<td>115 s</td>
</tr>
<tr>
<td>8</td>
<td>1935</td>
<td>143 s</td>
</tr>
</tbody>
</table>

From depth 5, with queue size 1935, we can guarantee optimality (on 500 random traces)

Time needed to align log offline: +/- 50 minutes

(same model with log fitness 0.81: 136 sec)
Conclusions

• Alignments explain relations between event logs and process models
• Alignments are foundational to process mining
• Computing alignments is computationally hard
• Many variants exist, but all guarantee:
  • The projection to the log provides the observed trace
  • The projection to the model provides a valid run thereof
Future challenges

• Computational complexity
  • Especially with data, resources
  • Find tight bounds for online alignments
  • Fast “approximate” alignments

• Determinism of alignments
  • Same choices across alignments

• Obtaining the right models,
  • Translate informal text to formal models
  • Alignments on BPMN

• Industry adaptation
Questions?

Stay clear of trees, align with roads instead!