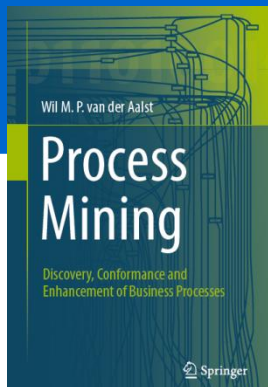


Process Mining: Data Science in Action

Token-Based Replay: Some Examples

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www.processmining.org

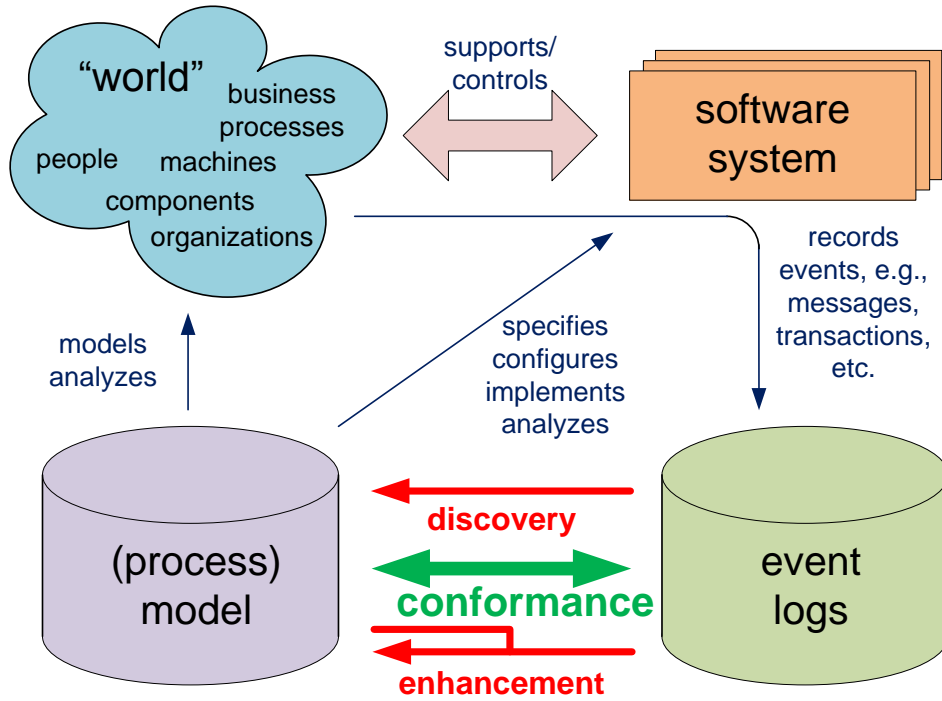


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Eindhoven
University of Technology

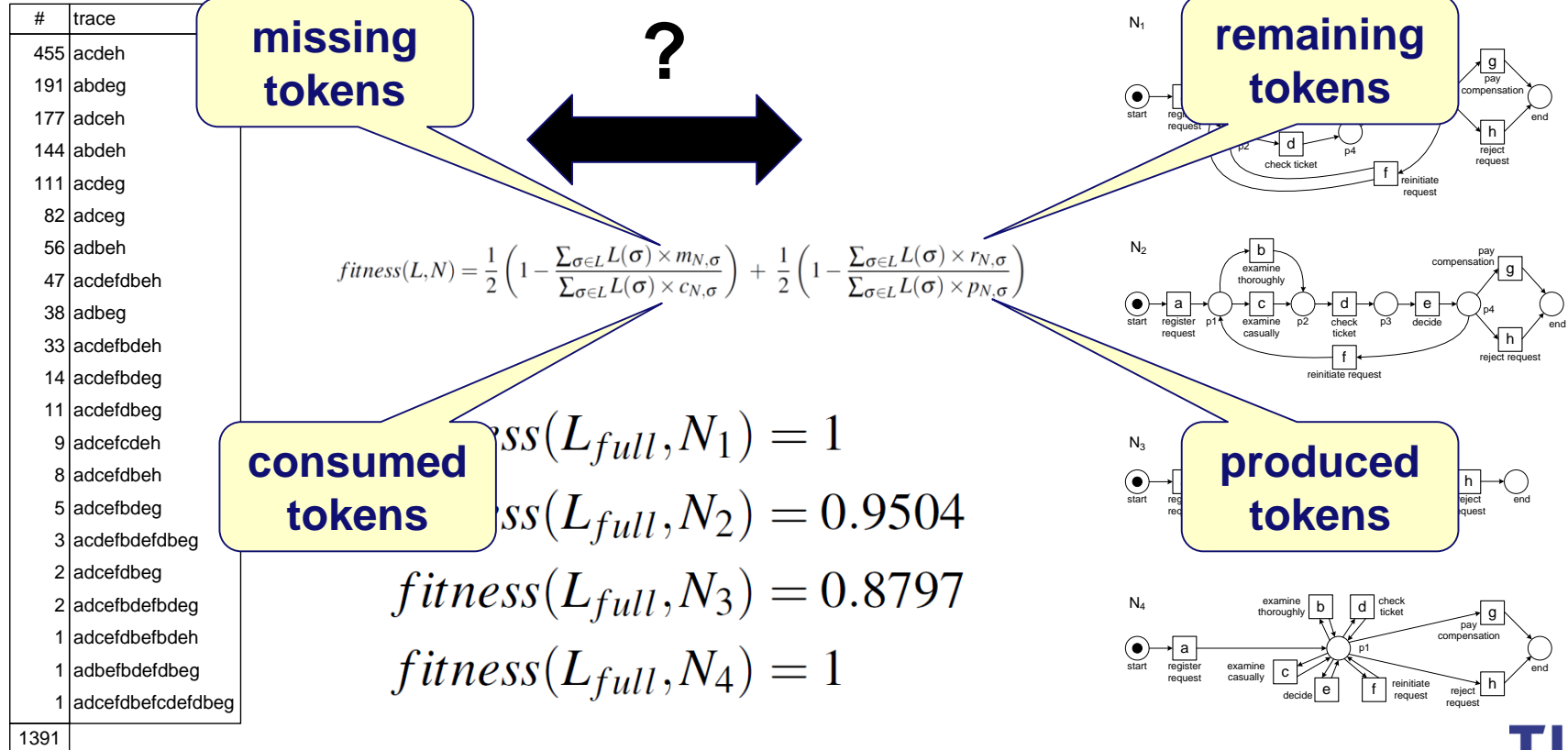
Where innovation starts

Conformance checking



1. Conformance checking using causal footprints.
2. Conformance checking based on **token-based replay**.
3. Alignment-based conformance checking.

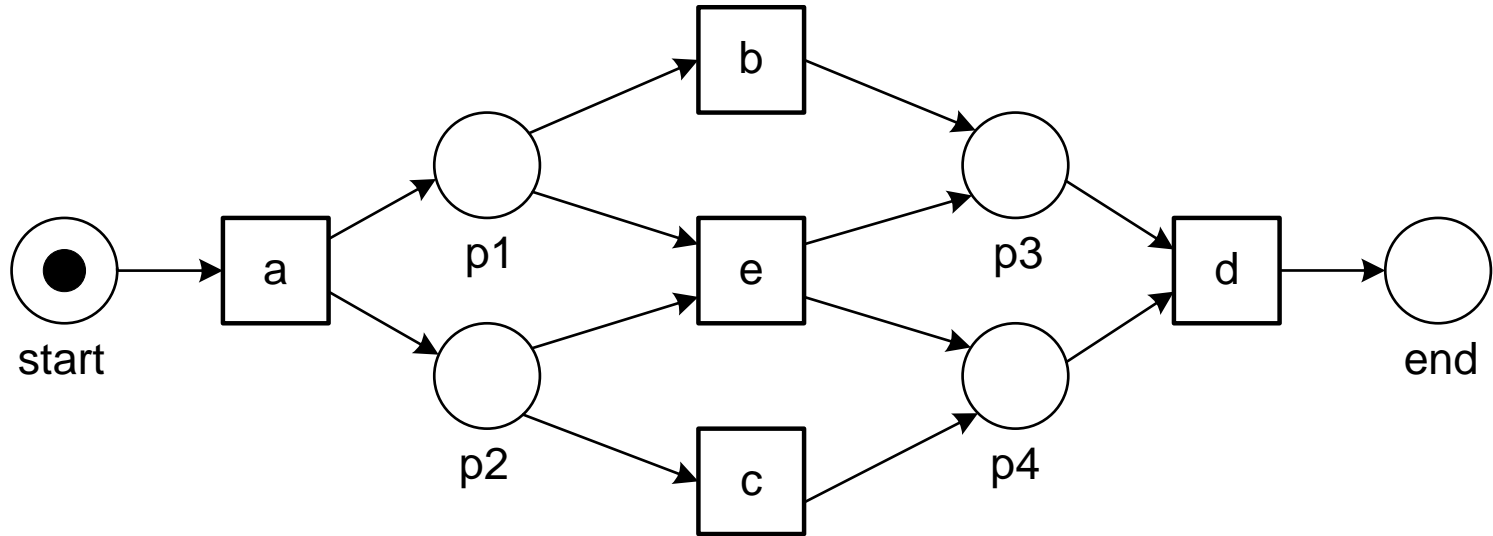
Last lecture: Token-based replay



Question (may take some time)

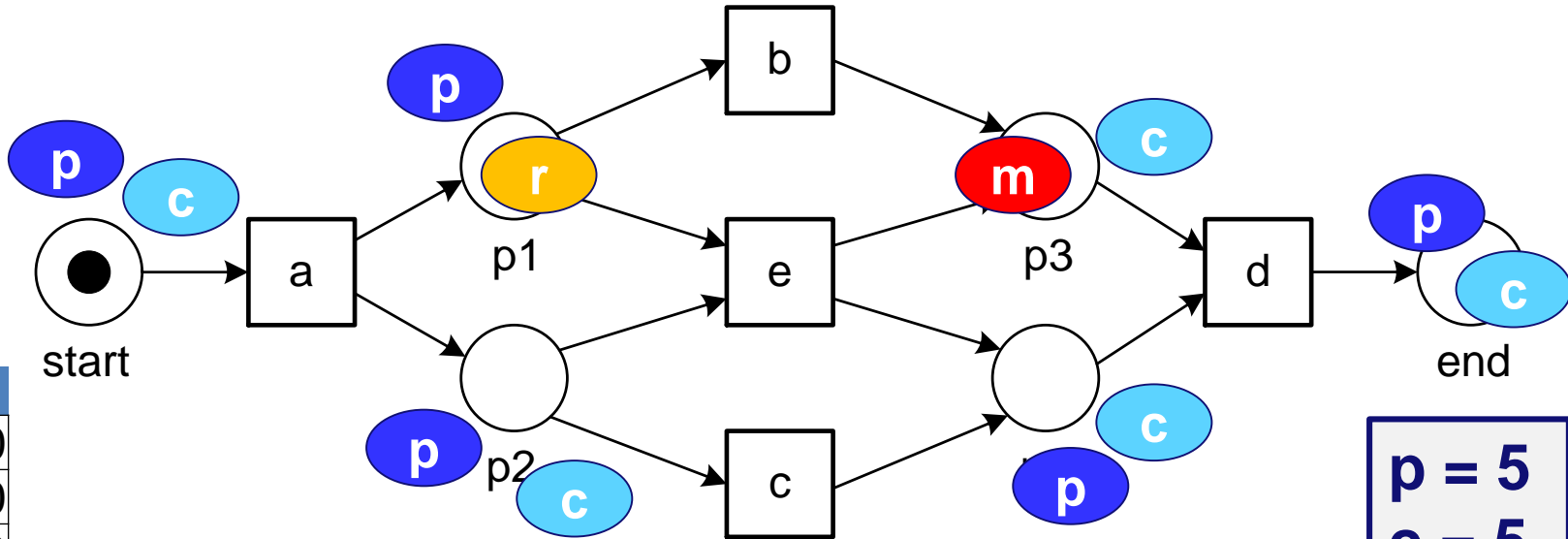
Compute fitness using missing and remaining tokens

trace	frequency
abcd	10
acbd	10
aed	10
abd	2
acd	1
ad	1
abbd	1



- Consider the event log containing 35 cases.
- What is the fitness?

Let us pick one trace: acd



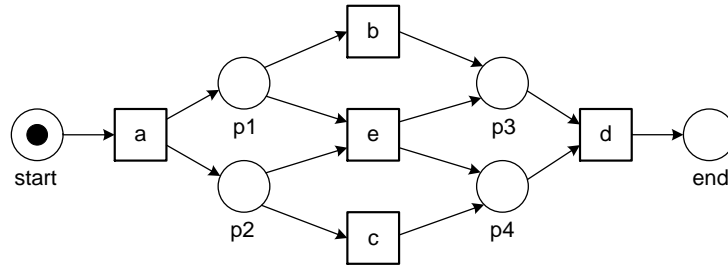
trace	frequency
abcd	10
acbd	10
aed	10
abd	2
acd	1
ad	1
abbd	1

$$fitness(L, N) = \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times m_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times c_{N, \sigma}} \right) + \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times r_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times p_{N, \sigma}} \right)$$

p = 5
c = 5
m = 1
r = 1

Fitness = 0.9658

trace	frequency	produced tokens (p)	remaining tokens (r)	consumed tokens (c)	missing tokens (m)	produced tokens (p')	remaining tokens (r')	consumed tokens (c')	missing tokens (m')
abcd	10	6	0	6	0	60	0	60	0
acbd	10	6	0	6	0	60	0	60	0
aed	10	6	0	6	0	60	0	60	0
abd	2	5	1	5	1	10	2	10	2
acd	1	5	1	5	1	5	1	5	1
ad	1	4	2	4	2	4	2	4	2
abbd	1	6	2	6	2	6	2	6	2



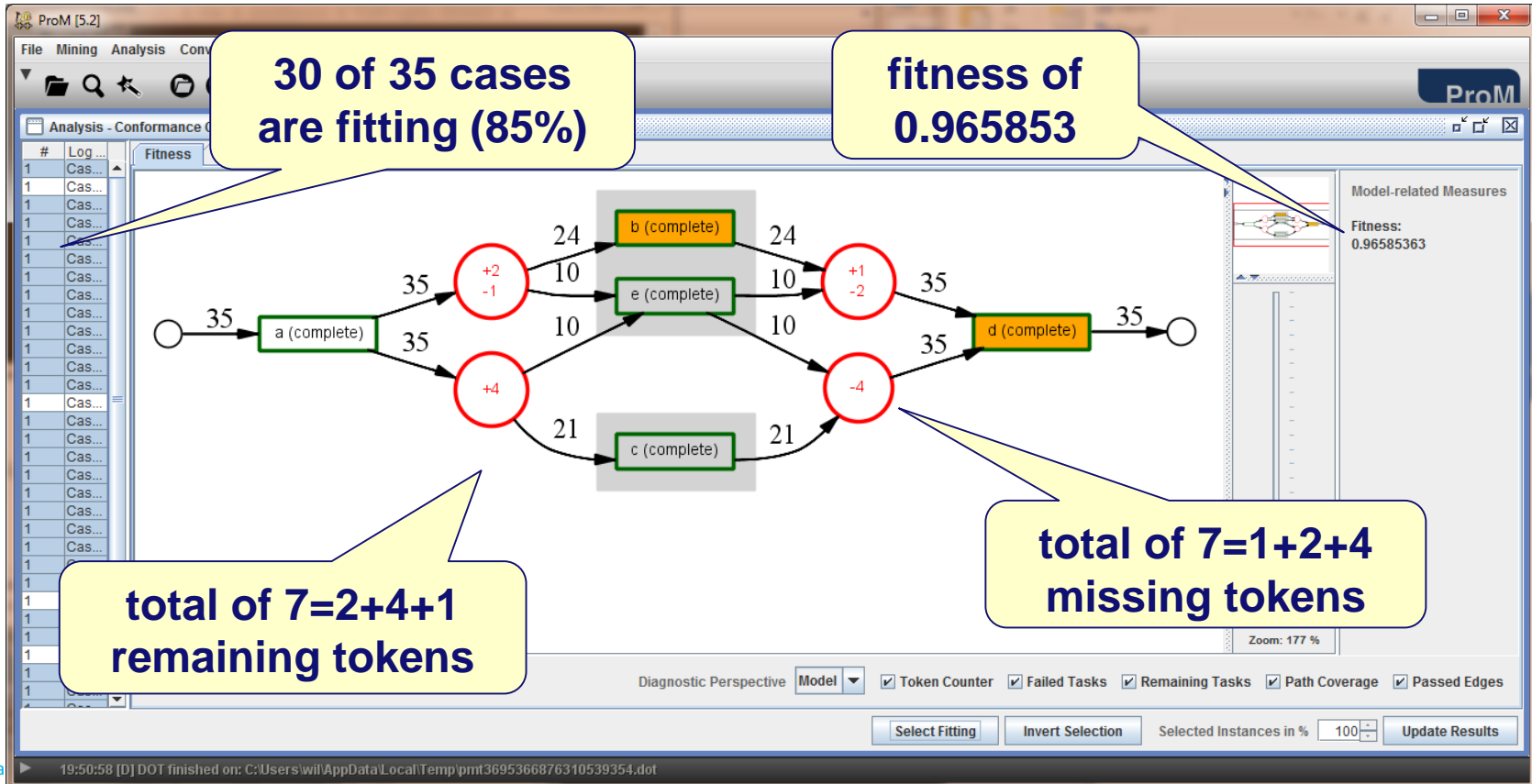
205	7	205	7
sum p	sum r	sum c	sum m

fitness	0.965853659
---------	--------------------

$$fitness(L, N) = \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times m_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times c_{N, \sigma}} \right) + \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times r_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times p_{N, \sigma}} \right)$$

ProM 5.2 output

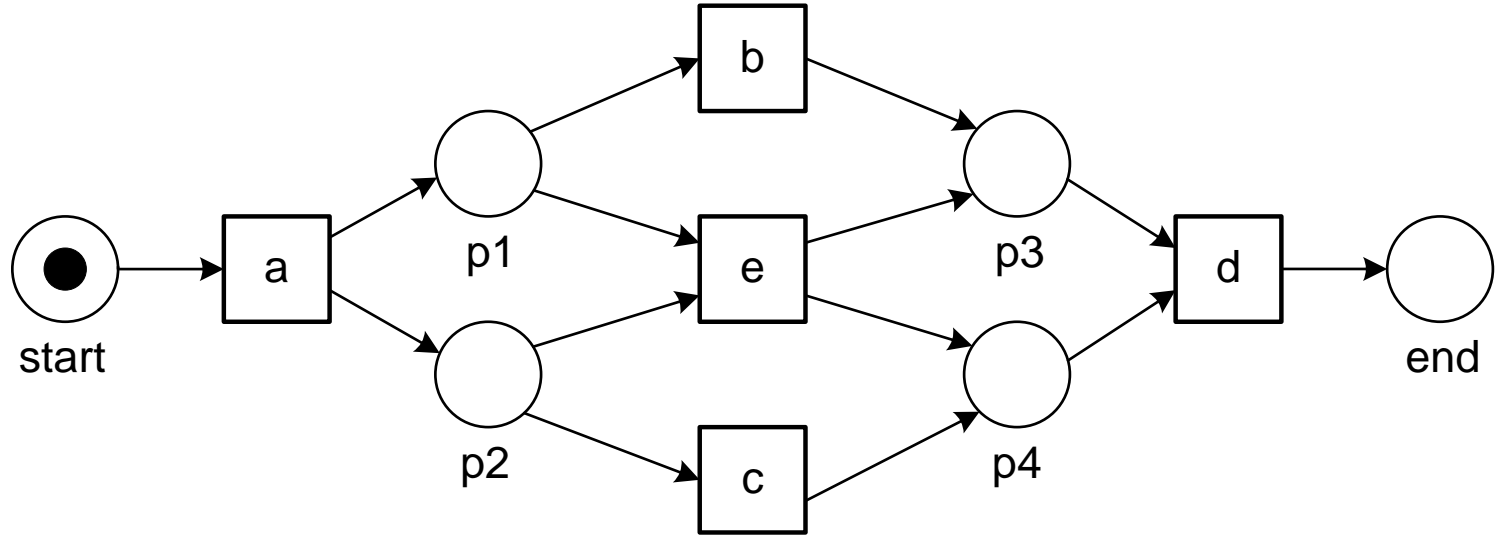
(ProM 6 only supports more advanced conformance checking techniques)



Question

Compute fitness using missing and remaining tokens

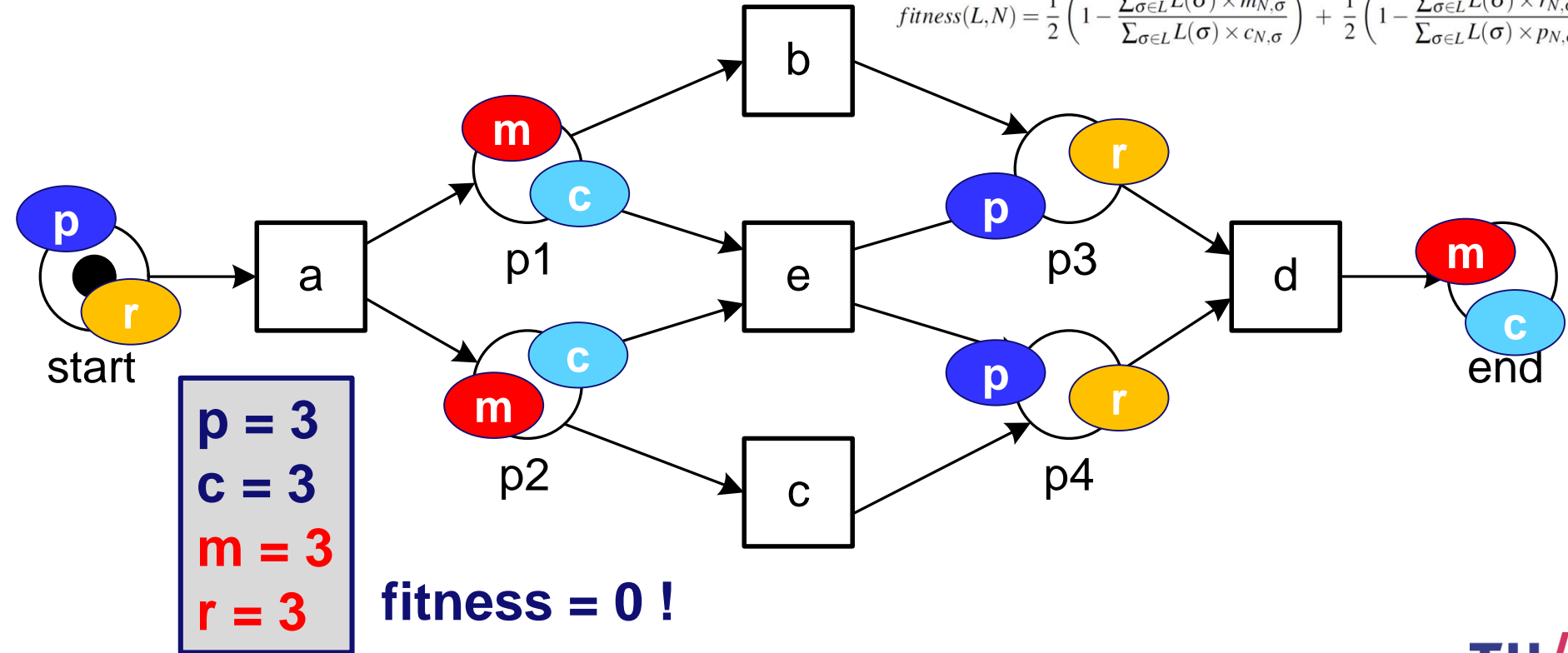
trace	frequency
e	1



- Consider the event log containing just one case: $L = [\langle e \rangle]$.
- What is the fitness (using token-based replay)?

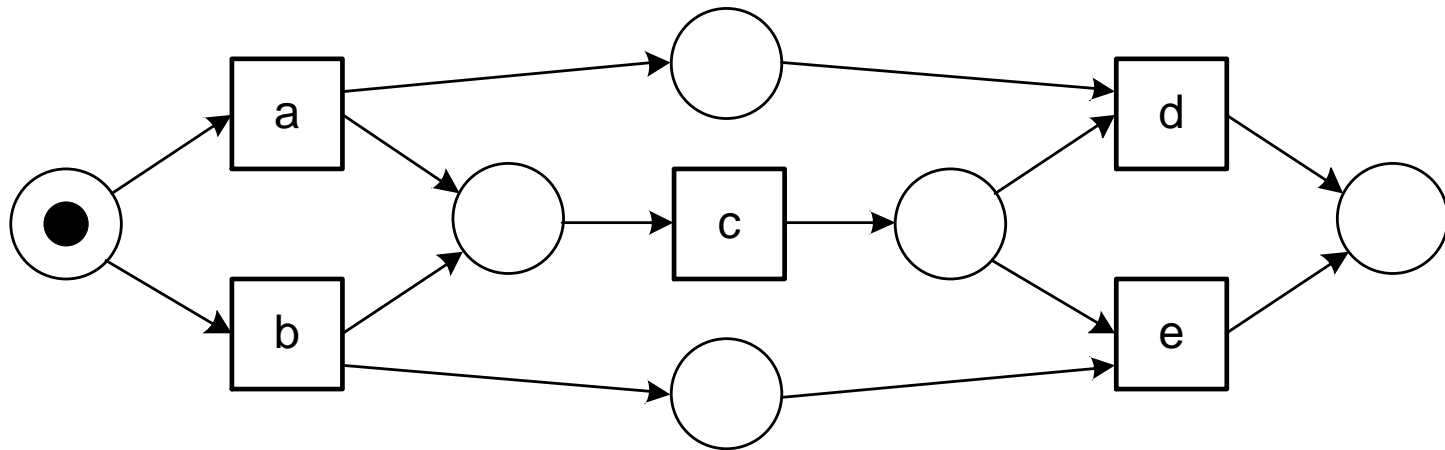
Answer obtained by replaying $\langle e \rangle$

$$\text{fitness}(L, N) = \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times m_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times c_{N, \sigma}} \right) + \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times r_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times p_{N, \sigma}} \right)$$



Another example

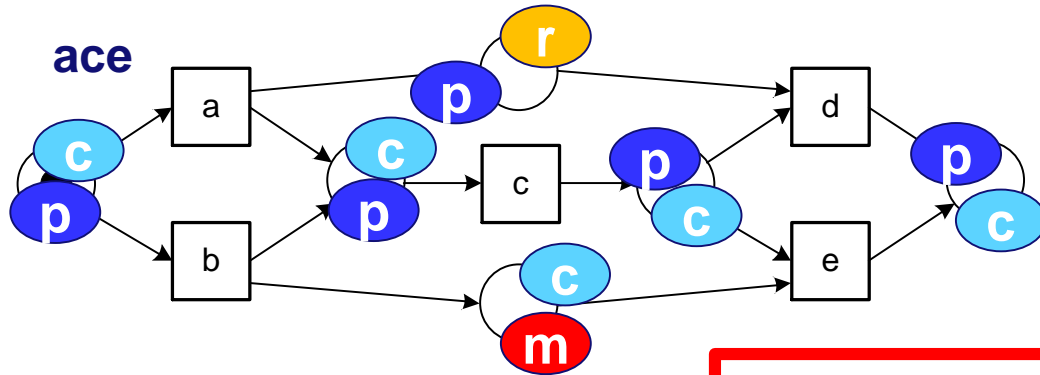
trace	frequency
acd	10
bce	10
ace	5
bcd	5
dca	1
abd	1
d	1



- Consider the event log containing 33 cases.
- What is the fitness?

Fitness = 0.895705521

trace	frequency	produced tokens (p)	remaining tokens (r)	consumed tokens (c)	missing tokens (m)	produced tokens (all)	remaining tokens (all)	consumed tokens (all)	missing tokens (all)
acd	10	5	0	5	0	50	0	50	0
bce	10	5	0	5	0	50	0	50	0
ace	5	5	1	5	1	25	5	25	5
bcd	5	5	1	5	1	25	5	25	5
dca	1	5	3	5	3	5	3	5	3
abd	1	6	3	5	2	6	3	5	2
d	1	2	1	3	2	2	1	3	2

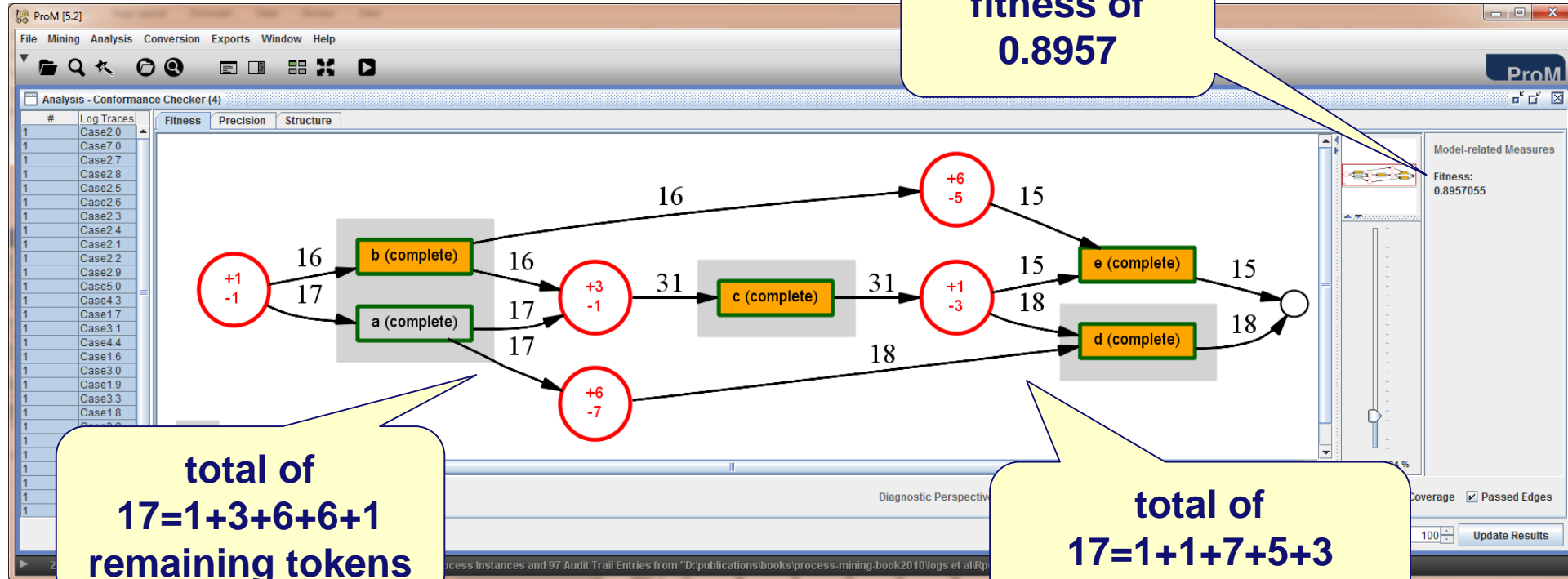


163	17	163	17
sum p	sum r	sum c	sum m

fitness	0.895705521
---------	--------------------

$$fitness(L, N) = \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times m_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times c_{N, \sigma}} \right) + \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times r_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times p_{N, \sigma}} \right)$$

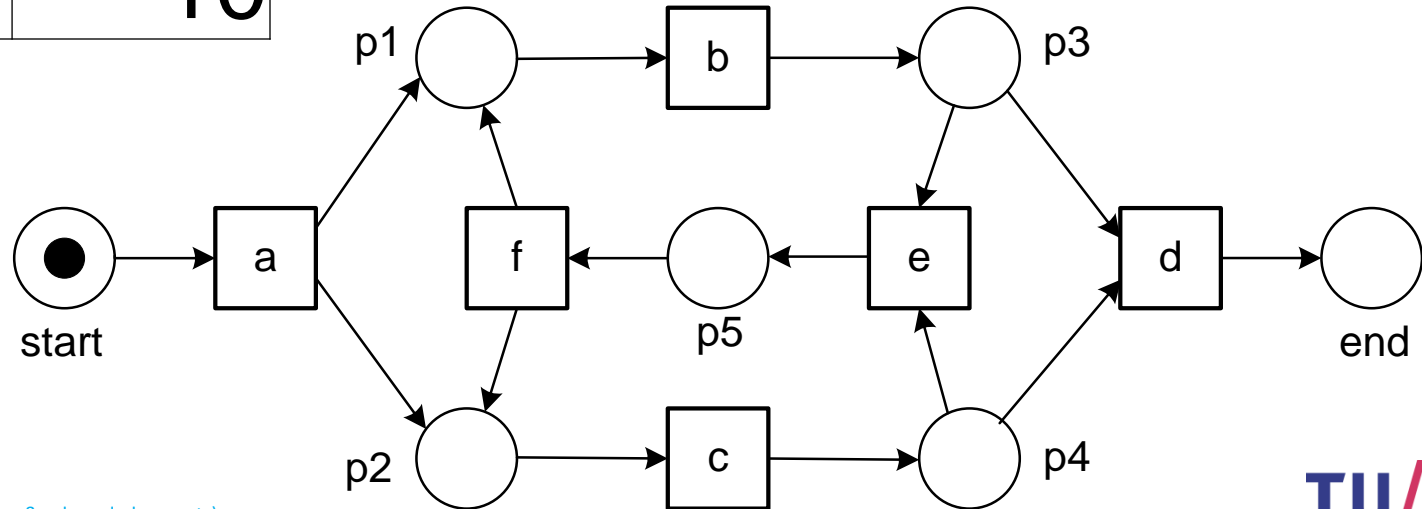
ProM 5.2 diagnostics



Another example

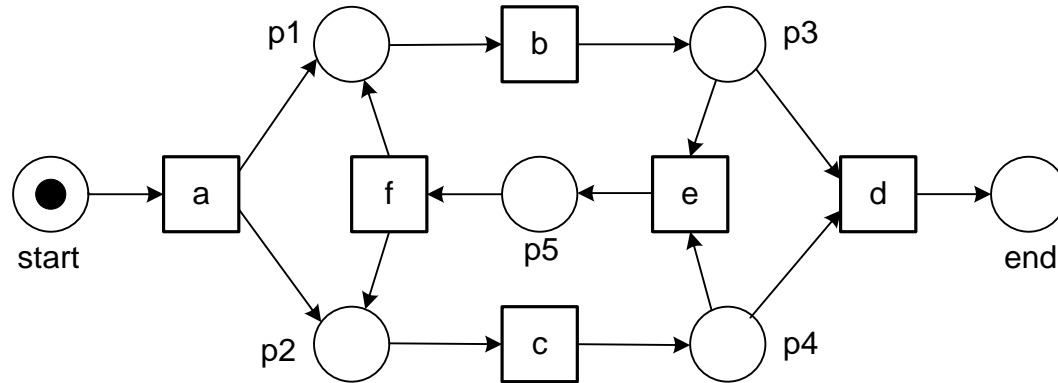
trace	frequency
abefcd	10
abbefccd	10

- Consider the event log containing 20 cases.
- What is the fitness?



Fitness = 0.8

trace	frequency	produced tokens (p)	remaining tokens (r)	consumed tokens (c)	missing tokens (m)	produced tokens (all)	remaining tokens (all)	consumed tokens (all)	missing tokens (all)
abefcd	10	9	2	9	2	90	20	90	20
abbefccd	10	11	2	11	2	110	20	110	20



200
sum p

40
sum r

200
sum c

40
sum m

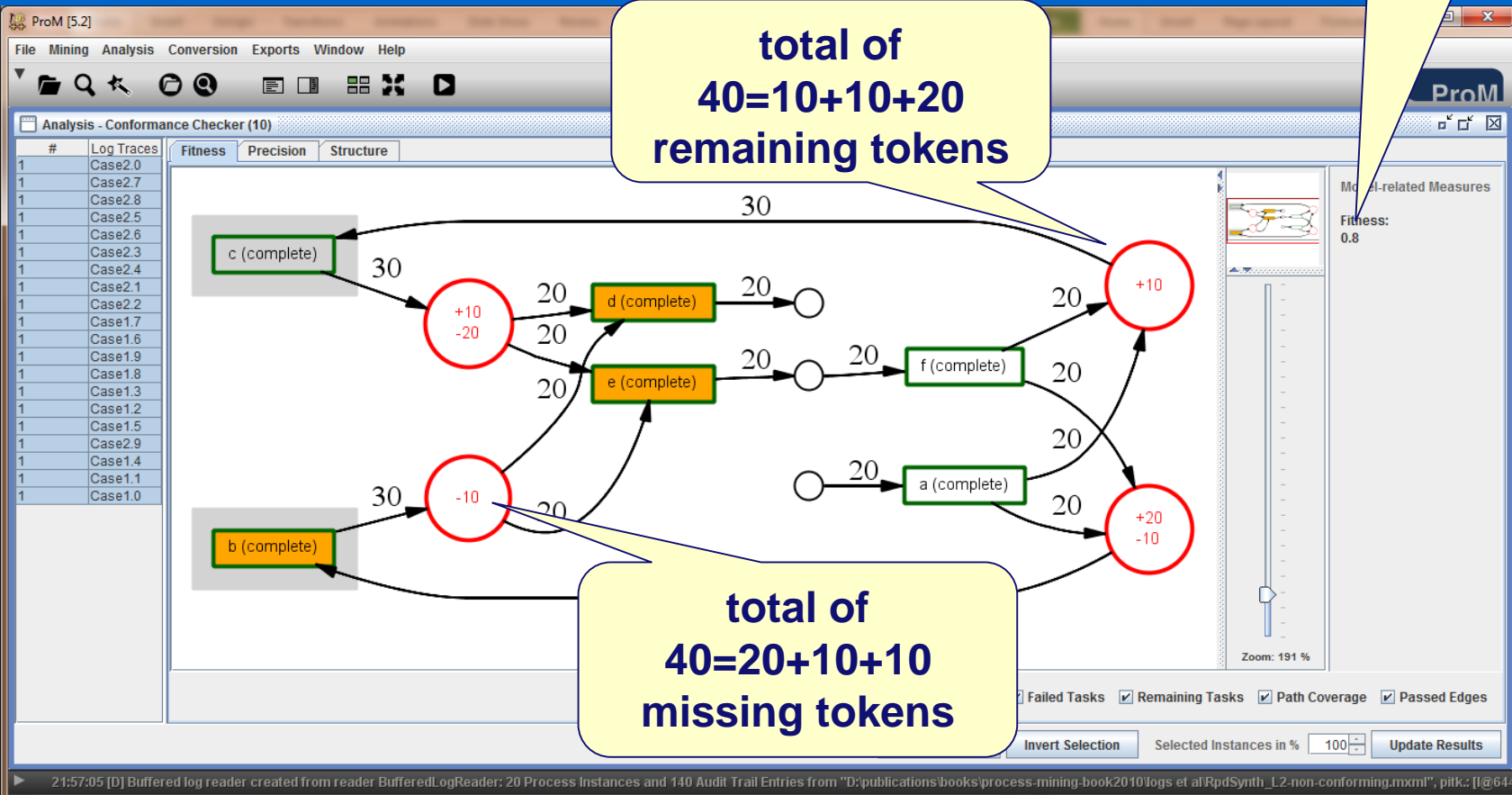
fitness

0.8

$$fitness(L, N) = \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times m_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times c_{N, \sigma}} \right) + \frac{1}{2} \left(1 - \frac{\sum_{\sigma \in L} L(\sigma) \times r_{N, \sigma}}{\sum_{\sigma \in L} L(\sigma) \times p_{N, \sigma}} \right)$$

ProM 5.2 diagnostics

fitness of 0.8



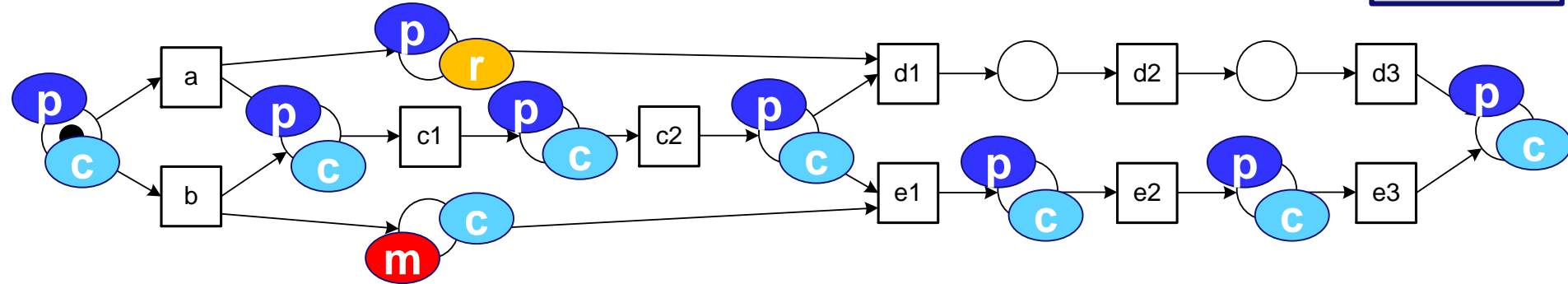
Limitations

- Basic replay approach assumes **visible & uniquely labeled** transitions.
- ProM implementation uses **heuristics** to deal with silent transitions and multiple transitions having the same label.
- Conformance values sometimes **too optimistic** due to "token flooding".
- Local decision making may lead to misleading results.

Local decision making is not enough ...

p = 8
c = 8
m = 1
r = 1
f = 0.875

$\langle a, c1, c2, e1, e2, e3 \rangle$



- Replay technique does **not** provide a corresponding path through the model (vital for conformance/performance analysis and other diagnostics).
- We would like to see the "closest path", i.e., $\langle \mathbf{b}, c1, c2, e1, e2, e3 \rangle$.

Next: alignments

$\langle a, c1, c2, e1, e2, e3 \rangle$

move in
log only

move in
model
only

synchronous
move
(move in both)

top row
corresponds to
trace in event log

bottom row
corresponds to
path in model

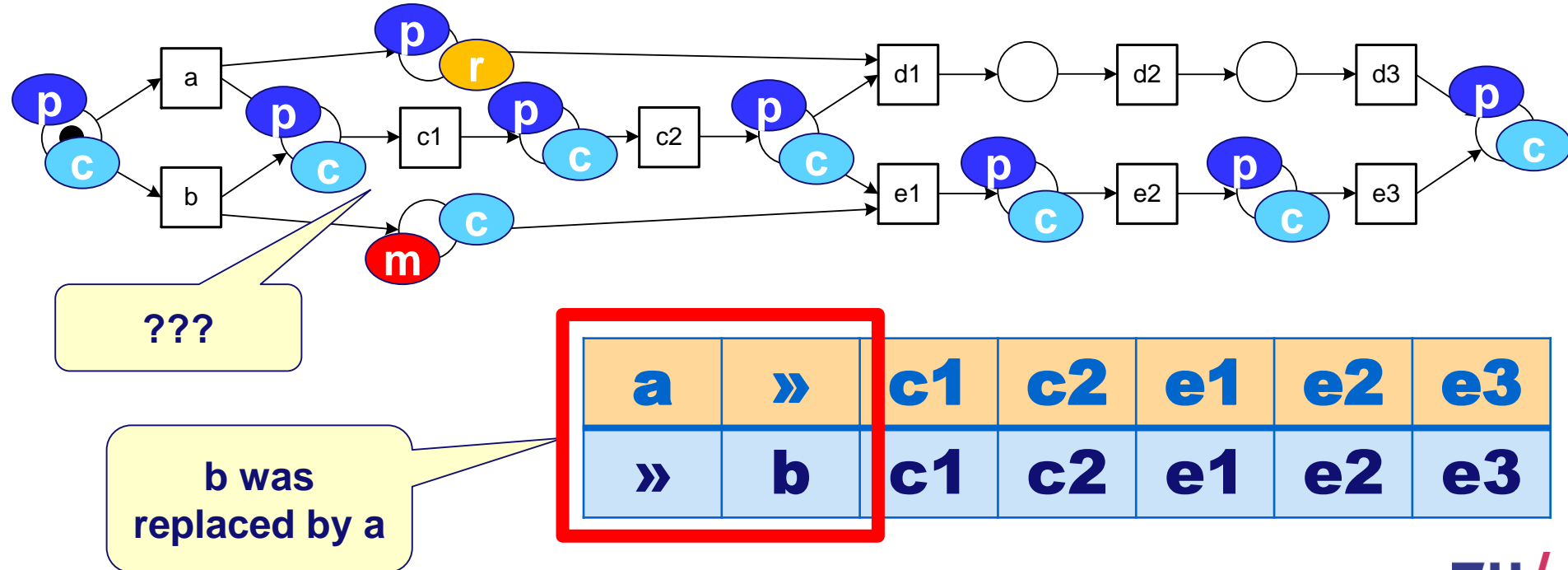
a	»	c1	c2	e1	e2	e3
»	b	c1	c2	e1	e2	e3

"no move"

"no move"

Alignments provide better diagnostics

$\langle a, c1, c2, e1, e2, e3 \rangle$



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Analysis

Chapter 3
Data Mining

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Conformance
Checking

Chapter 8
Mining Additional
Perspectives

Chapter 9
Operational Support

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Getting the Data

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Introduction

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Chapter 12
Analyzing "Spaghetti
Processes"

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Navigation

Chapter 14
Epilogue

