Quantitative Verification The Good, The Bad and The Ugly

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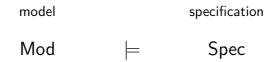




# Nice People

Sebastian S. Bauer, Nikola Beneš, Zoltán Ésik, Lisbeth Fajstrup, Eric Goubault, Line Juhl, Jan Křetínský, Kim G. Larsen, Martin Raussen, Georg Struth, Claus Thrane, Louis-Marie Traonouez

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Model Checkin	g			

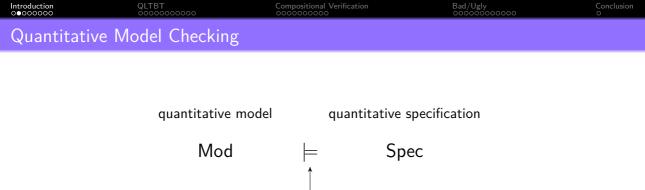


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 Quantitative Model Checking
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quantitative model quantitative specification

 $\mathsf{Mod} \models \mathsf{Spec}$ 





replace by

# $\models_{\varepsilon}$

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Claus T:	Quantitative Quan	titative Quantitative	Analysis	
Quantit	ative <i>Models</i> Q	uantitative <i>Logics</i>	Quantitative Verification	

Quantitative <i>Models</i>	Quantitative <i>Logics</i>	Quantitative Verification
$x \ge 4$ x := 0	$Pr_{\leq .1}(\Diamond \mathit{error})$	$[\![\phi]\!](s) = 3.14$ d(s,t) = 42

Boolean world	"Quantification"
Trace equivalence $\equiv$	Linear distances $d_L$
Bisimilarity $\sim$	Branching distances <i>d</i> <sub>B</sub>
$s \sim t$ implies $s \equiv t$	$d_L(s,t) \leq d_B(s,t)$
$\pmb{s} \models \phi \text{ or } \pmb{s} \not\models \phi$	$\llbracket \phi \rrbracket (s)$ is a quantity
$oldsymbol{s} \sim oldsymbol{t}$ iff $orall \phi : oldsymbol{s} \models \phi \Leftrightarrow oldsymbol{t} \models \phi$	$d_B(s,t) = \sup_{\phi} d(\llbracket \phi \rrbracket(s), \llbracket \phi \rrbracket(t))$

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Compositiona	al Verification			

model specification Mod = Spec

• 
$$\mathsf{Mod} \models \mathsf{Spec}_1 \& \mathsf{Spec}_1 \leq \mathsf{Spec}_2 \Longrightarrow \mathsf{Mod} \models \mathsf{Spec}_2$$

 $\bullet \; \mathsf{Mod} \models \mathsf{Spec}_1 \, \& \, \mathsf{Mod} \models \mathsf{Spec}_2 \Longrightarrow \, \mathsf{Mod} \models \mathsf{Spec}_1 \wedge \mathsf{Spec}_2$ 

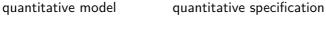
 $\bullet \ \mathsf{Mod}_1 \models \mathsf{Spec}_1 \And \mathsf{Mod}_2 \models \mathsf{Spec}_2 \Longrightarrow \mathsf{Mod}_1 \parallel \mathsf{Mod}_2 \models \mathsf{Spec}_1 \parallel \mathsf{Spec}_2$ 

 $\bullet \; \mathsf{Mod}_1 \models \mathsf{Spec}_1 \And \mathsf{Mod}_2 \models \mathsf{Spec}/\mathsf{Spec}_1 \Longrightarrow \mathsf{Mod}_1 \, \| \, \mathsf{Mod}_2 \models \mathsf{Spec}$ 

bottom-up and top-down

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 $\mathsf{Mod} \models_{\varepsilon} \mathsf{Spec}$ 

$$\bullet \ \mathsf{Mod} \models_{\varepsilon} \mathsf{Spec}_1 \And \mathsf{Spec}_1 \leq_{\varepsilon} \mathsf{Spec}_2 \Longrightarrow \mathsf{Mod} \models_{\varepsilon} \mathsf{Spec}_2$$

 $\bullet \; \mathsf{Mod} \models_{\varepsilon} \mathsf{Spec}_1 \, \& \, \mathsf{Mod} \models_{\varepsilon} \mathsf{Spec}_2 \Longrightarrow \, \mathsf{Mod} \models_{\varepsilon} \mathsf{Spec}_1 \wedge \mathsf{Spec}_2$ 

- $\bullet \ \mathsf{Mod}_1 \models_{\varepsilon} \mathsf{Spec}_1 \And \mathsf{Mod}_2 \models_{\varepsilon} \mathsf{Spec}_2 \Longrightarrow \mathsf{Mod}_1 \parallel \mathsf{Mod}_2 \models_{\varepsilon} \mathsf{Spec}_1 \parallel \mathsf{Spec}_2$
- $\bullet \ \mathsf{Mod}_1 \models_{\varepsilon} \mathsf{Spec}_1 \And \mathsf{Mod}_2 \models_{\varepsilon} \mathsf{Spec}/\mathsf{Spec}_1 \Longrightarrow \mathsf{Mod}_1 \parallel \mathsf{Mod}_2 \models_{\varepsilon} \mathsf{Spec}$
- surely not the same  $\varepsilon$  everywhere!?

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User Stories				

### "In your quantitative verification, what type of distances do you use?"

• point-wise	$D(\sigma, au) = {{{{sup}_i}} {\sigma _i} - { au _i} }$
<ul> <li>accumulating</li> </ul>	$D(\sigma,  au) = \sum_i  \sigma_i -  au_i $
<ul> <li>limit-average</li> </ul>	$D(\sigma, au) = \limsup_{N rac{1}{N}} \sum_{i=0}^{N}  \sigma_i -  au_i $
<ul> <li>discounted</li> </ul>	$D(\sigma, au) = \sum_i {oldsymbol{\lambda}^i   \sigma_i -  au_i  }$
<ul> <li>maximum-lead</li> </ul>	$D(\sigma, \tau) = \sup_{N} \left  \sum_{i=0}^{N} (\sigma_i - \tau_i) \right $
• Cantor	$D(\sigma, au) = 1/(1 + \inf\{j \mid \sigma_j  eq  au_j\})$
• discrete	$D(\sigma, au)=0$ if $\sigma= au;$ $\infty$ otherwise

Introduction

# Asarin-Basset-Degorre 2018

$$D(\sigma,\tau) = \max \begin{cases} \sup_{i} \inf_{j} \{|t_i - s_j| \mid a_i = b_j\} \\ \sup_{j} \inf_{i} \{|t_i - s_j| \mid a_i = b_j\} \end{cases}$$

- In quantitative verification, lots of different distances
- Develop theory to cover all/most of them
  - idea: use bisimulation games
- $\Rightarrow$  The Quantitative Linear-Time–Branching-Time Spectrum
  - QAPL 2011, FSTTCS 2011, TCS 2014

Challenge (ca. 2012):

- How to make this compositional?
- Still not satisfied!

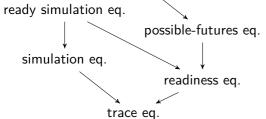


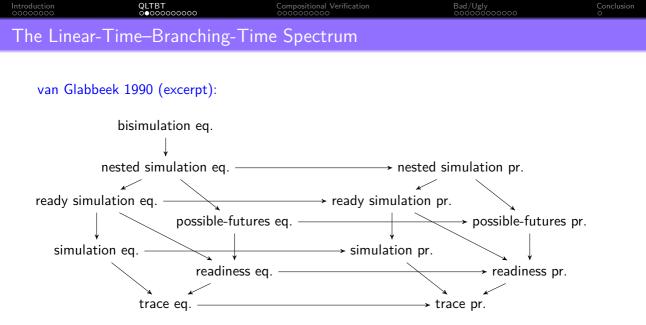
2 The Quantitative Linear-Time-Branching-Time Spectrum

Compositional Verification



QLTBT Compositional Verification 00000000000 The Linear-Time-Branching-Time Spectrum van Glabbeek 1990 (excerpt): bisimulation eq. nested simulation eq.



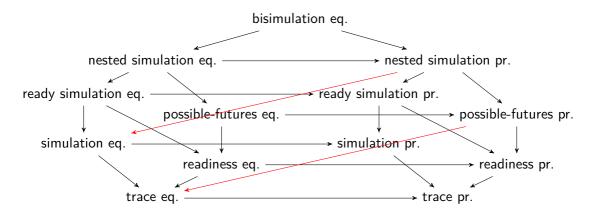


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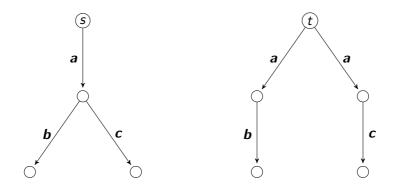
 The Linear-Time-Branching-Time Spectrum

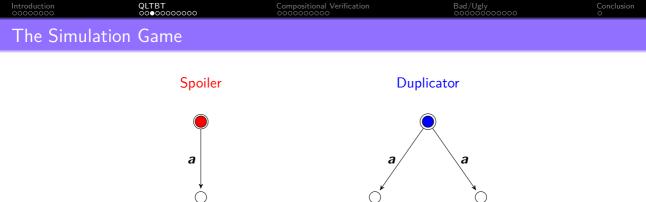
## van Glabbeek 1990 (excerpt):



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# The Simulation Game



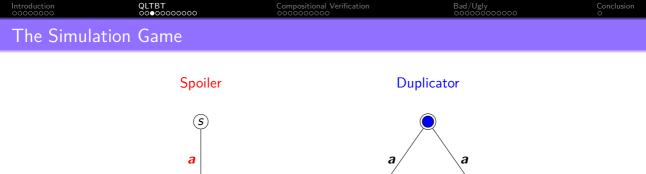


b

С

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b



b

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b

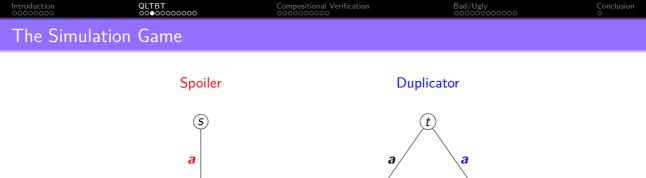


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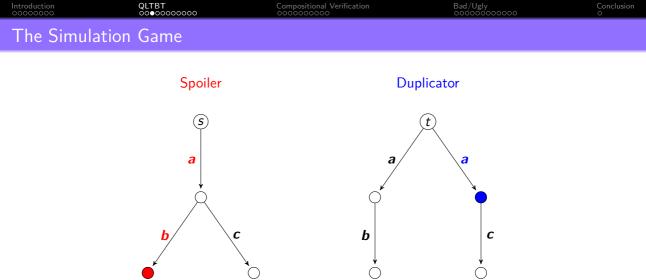


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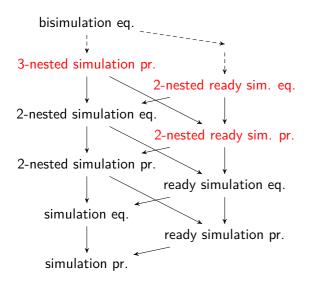
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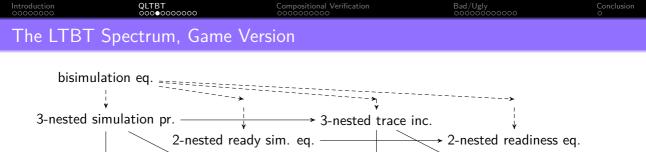
b



Spoiler wins







 $\searrow$  2-nested trace inc.

ready simulation eq.

ready simulation pr. –

2-nested simulation eq.  $\rightarrow$  2-nested trace eq.

2-nested simulation pr. -

simulation eq.

simulation pr. -

→ trace inc.

2-nested ready sim. pr.  $\longrightarrow$  2-nested readiness pr.

readiness eq.

→ readiness pr.

# The Simulation Game, Revisited

QLTBT

Introduction

- 1. Player 1 chooses edge from s (leading to s')
- 2. Player 2 chooses matching edge from t (leading to t')
- 3. Game continues from configuration s', t'
- $\omega$ . If Player 2 can always answer: YES, *t* simulates *s*. Otherwise: NO
- Or, as an Ehrenfeucht-Fraïssé game ("delayed evaluation"):
  - 1. Player 1 chooses edge from s (leading to s')
  - 2. Player 2 chooses edge from t (leading to t')
  - 3. Game continues from new configuration s', t'
  - $\omega$ . At the end (maybe after infinitely many rounds!), compare the chosen traces: If the trace chosen by t matches the one chosen by s: YES Otherwise: NO

Compositional Verification

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# Quantitative Ehrenfeucht-Fraïssé Games

The quantitative setting:

- Assume we have a way, possibly application-determined, to measure distances of (finite or infinite) traces
- a hemimetric  $D: (\sigma, \tau) \mapsto D(\sigma, \tau) \in \mathbb{R}_{\geq 0} \cup \{\infty\}$

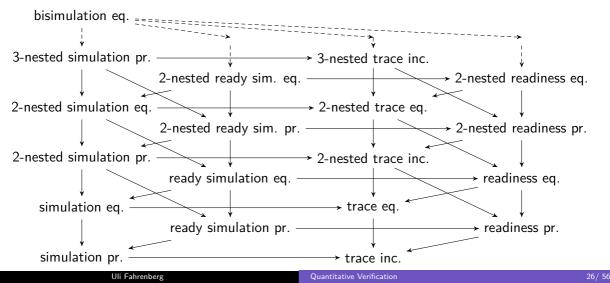
The quantitative Ehrenfeucht-Fraïssé game:

- 1. Player 1 chooses edge from s (leading to s')
- 2. Player 2 chooses edge from t (leading to t')
- 3. Game continues from new configuration s', t'
- ω. At the end, compare the chosen traces σ, τ: The simulation distance from s to t is defined to be D(σ, τ)
  - Player 1 plays to maximize  $D(\sigma, \tau)$ ; Player 2 plays to minimize

This can be generalized to all the games in the LTBT spectrum.



For any trace distance  $D: (\sigma, \tau) \mapsto D(\sigma, \tau) \in \mathbb{R}_{\geq 0} \cup \{\infty\}$ :



# Quantitative EF Games: Some Details

- Configuration of the game:  $(\pi, \rho)$ :  $\pi$  the Player-1 choices up to now;  $\rho$  the Player-2 choices
- Strategy: mapping from configurations to next moves
  - $\Theta_i$ : set of Player-*i* strategies
- $\bullet$  Simulation strategy: Player-1 moves allowed from end of  $\pi$
- $\bullet$  Bisimulation strategy: Player-1 moves allowed from end of  $\pi$  or end of  $\rho$ 
  - (hence  $\pi$  and  $\rho$  are generally not paths "mingled paths")
- Pair of strategies  $\implies$  (possibly infinite) sequence of configurations
- Take the limit; unmingle  $\implies$  pair of (possibly infinite) traces  $(\sigma, \tau)$
- Bisimulation distance:  $\sup_{\theta_1 \in \Theta_1} \inf_{\theta_2 \in \Theta_2} d_T(\sigma, \tau)$
- Simulation distance:  $\sup_{\theta_1 \in \Theta_1^0} \inf_{\theta_2 \in \Theta_2} d_{\mathcal{T}}(\sigma, \tau)$

(restricting Player 1's capabilities)

# Quantitative EF Games: Some Details – II

 $\bullet$  Blind Player-1 strategies: depend only on the end of  $\rho$ 

- ("cannot see Player-2 moves")
- $\tilde{\Theta}_1$ : set of blind Player-1 strategies
- Trace inclusion distance: sup inf  $d_T(\sigma, \tau)$  $\theta_1 \in \tilde{\Theta}_1^0 \theta_2 \in \Theta_2$
- $\bullet\,$  For nesting: count the number of times Player 1 switches between end of  $\pi$  and end of  $\rho$ 
  - $\Theta_1^k$ : k switches allowed
- Nested simulation distance:  $\sup_{\theta_1 \in \Theta_1^1} \inf_{\theta_2 \in \Theta_2} d_T(\sigma, \tau)$
- Nested trace inclusion distance:  $\sup_{\theta_1 \in \tilde{\Theta}_1^1 \, \theta_2 \in \Theta_2} \inf d_T(\sigma, \tau)$  (!)
- $\bullet\,$  For ready: allow extra "I'll see you" Player-1 transition from end of  $\rho$

# Transfer Theorem

#### Theorem

If two equivalences or preorders are inequivalent in the qualitative setting, and the trace distance D is separating, then the corresponding QLTBT distances are topologically inequivalent.

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Recu	sive Characterization			
Tł	eorem			
d	the trace distance $D : (\sigma, \tau) \mapsto$ = $g \circ f : Tr \times Tr \rightarrow L \rightarrow \mathbb{R}_{\geq 0} \cup$ d f has a recursive characteriza	$\downarrow \{\infty\}$ through a complete	lattice L,	or some

then all distances in the corresponding QLTBT spectrum are given as least fixed points of some

functionals using F.

All trace distances I know can be expressed recursively like this.

 $F: \Sigma \times \Sigma \times L \rightarrow L$  which is monotone in the third coordinate,

• Example: simulation distance:

$$d_{sim}(s,t) = \sup_{s \xrightarrow{a} s'} \inf_{t \xrightarrow{b} t'} F(a,b,d_{sim}(s',t'))$$
(I.f.p.)

- *L* is "memory"
- also gives relation family characterization

#### Uli Fahrenberg

#### Quantitative Verification



2 The Quantitative Linear-Time-Branching-Time Spectrum

## Compositional Verification



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Specification	n Theories			

Let Mod be a set of models with an equivalence  $\sim$ .

### Definition

A complete specification theory for (Mod,  $\sim$ ) is (Spec,  $\leq$ ,  $\parallel$ ,  $\chi$ ) such that

- $\leq$  is a refinement preorder on Spec
- $\chi : \mathsf{Mod} \to \mathsf{Spec}$  picks out characteristic specifications
  - *i.e.*,  $\forall \mathcal{M}_1, \mathcal{M}_2 \in \mathsf{Mod} : \mathcal{M}_1 \sim \mathcal{M}_2 \iff \chi(\mathcal{M}_1) \leq \chi(\mathcal{M}_2)$

• (Spec,  $\leq$ ,  $\parallel$ ) forms a bounded commutative distributive residuated lattice up to  $\leq$   $\cap$   $\geq$ 

 $\Rightarrow~\lor$  and  $\land$  on Spec; double distributivity;  $\bot,\top\in$  Spec

 $\bullet$  everything up to modal equivalence  $\equiv = \leq \cap \geq$ 

 $\Rightarrow$  || distributes over  $\lor$ , has unit U, has residual / (up to  $\equiv$ )

• 
$$\mathcal{S}_1 \| \mathcal{S}_2 \leq \mathcal{S}_3 \iff \mathcal{S}_2 \leq \mathcal{S}_3 / \mathcal{S}_1$$

- Disjunctive modal transition systems
- Acceptance automata
- Hennessy-Milner logic with maximal fixed points
- CONCUR 2013, ICTAC 2014, I&C 2020 (all with bisimulation as model equivalence  $\sim$ )

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Acceptance Au	tomata			

## Let $\Sigma$ be a finite alphabet.

### Definition

A (nondeterministic) acceptance automaton (AA) is a structure  $\mathcal{A} = (S, S^0, \text{Tran})$ , with  $S \supseteq S^0$  finite sets of states and initial states and Tran :  $S \to 2^{2^{\Sigma \times S}}$  an assignment of *transition constraints*.

- standard labeled transition system (LTS): Tran :  $S \rightarrow 2^{\Sigma \times S}$  (coalgebraic view)
- (for AA:) Tran(s) = { $M_1, M_2, \dots$ }: provide  $M_1$  or  $M_2$  or  $\dots$
- a disjunctive choice of conjunctive constraints
- J.-B. Raclet 2008 (but deterministic); see also H. H. Hansen 2003
- note multiple initial states

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Refinement				

### Definition

Let 
$$A_1 = (S_1, S_1^0, \text{Tran}_1)$$
 and  $A_2 = (S_2, S_2^0, \text{Tran}_2)$  be AA.  
A relation  $R \subseteq S_1 \times S_2$  is a modal refinement if:  
**a**  $\forall s_1^0 \in S_1^0 : \exists s_2^0 \in S_2^0 : (s_1^0, s_2^0) \in R$   
**a**  $\forall (s_1, s_2) \in R : \forall M_1 \in \text{Tran}_1(s_1) : \exists M_2 \in \text{Tran}_2(s_2) :$   
**b**  $\forall (a, t_1) \in M_1 : \exists (a, t_2) \in M_2 : (t_1, t_2) \in R$   
**a**  $\forall (a, t_2) \in M_2 : \exists (a, t_1) \in M_1 : (t_1, t_2) \in R$   
Write  $A_1 \leq A_2$  if there exists such a modal refinement.

• for any constraint choice  $M_1$  there is a bisimilar choice  $M_2$ 

- $\mathcal{A}_1$  has fewer choices than  $\mathcal{A}_2$
- no more choices  $\hat{=}$  only one  $M \in \operatorname{Tran}(s) \hat{=} \mathsf{LTS}$
- formally: an embedding  $\chi$  : LTS  $\hookrightarrow$  AA such that  $\chi(\mathcal{L}_1) \leq \chi(\mathcal{L}_2)$  iff  $\mathcal{L}_1$  and  $\mathcal{L}_2$  are bisimilar

(init) (tran)

Compositional Verification Introduction QLTBT Bad/Ugly Conclusion 00000000000 Logical Operations Let  $A_1 = (S_1, S_1^0, \text{Tran}_1)$  and  $A_2 = (S_2, S_2^0, \text{Tran}_2)$  be AA. Disjunction:  $\mathcal{A}_1 \lor \mathcal{A}_2 = (S_1 \stackrel{+}{\cup} S_2, S_1^0 \stackrel{+}{\cup} S_2^0, \operatorname{Tran}_1 \stackrel{+}{\cup} \operatorname{Tran}_2)$ Conjunction: define  $\pi_i: 2^{\Sigma \times S_1 \times S_2} \to 2^{\Sigma \times S_i}$  by  $\pi_1(M) = \{(a, s_1) \mid \exists s_2 \in S_2 : (a, s_1, s_2) \in M\}$  $\pi_2(M) = \{(a, s_2) \mid \exists s_1 \in S_1 : (a, s_1, s_2) \in M\}$ Let  $\mathcal{A}_1 \wedge \mathcal{A}_2 = (S_1 \times S_2, S_1^0 \times S_2^0, \text{Tran})$  with  $\operatorname{Tran}((s_1, s_2)) = \{M \subseteq \Sigma \times S_1 \times S_2 \mid \pi_1(M) \in \operatorname{Tran}_1(s_1), \pi_2(M) \in \operatorname{Tran}_2(s_2)\}$ 

#### Theorem

For all LTS  $\mathcal{L}$  and AA  $\mathcal{A}_1, \mathcal{A}_2$ :

$$\mathcal{L} \models \mathcal{A}_1 \lor \mathcal{A}_2 \iff \mathcal{L} \models \mathcal{A}_1 \text{ or } \mathcal{L} \models \mathcal{A}_2$$
  
 $\mathcal{L} \models \mathcal{A}_1 \land \mathcal{A}_2 \iff \mathcal{L} \models \mathcal{A}_1 \And \mathcal{L} \models \mathcal{A}_2$ 

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 Structural Operations:
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#### Structural Operations: Composition

Let 
$$\mathcal{A}_1 = (S_1, S_1^0, \operatorname{Tran}_1)$$
 and  $\mathcal{A}_2 = (S_2, S_2^0, \operatorname{Tran}_2)$  be AA

For  $M_1 \subseteq \Sigma \times S_1$  and  $M_2 \subseteq \Sigma \times S_2$ , define

$$M_1 || M_2 = \{ (a, (t_1, t_2)) \mid (a, t_1) \in M_1, (a, t_2) \in M_2 \}$$

(assumes CSP synchronization, but can be generalized)

Let  $\mathcal{A}_1 \| \mathcal{A}_2 = (S_1 \times S_2, S_1^0 \times S_2^0, \mathsf{Tran})$  with

 $\mathsf{Tran}((s_1, s_2)) = \{M_1 | M_2 | M_1 \in \mathsf{Tran}_1(s_1), M_2 \in \mathsf{Tran}_2(s_2)\}$ 

Theorem (independent implementability)

For all AA  $A_1$ ,  $A_2$ ,  $A_3$ ,  $A_4$ :

$$\mathcal{A}_1 \leq \mathcal{A}_3 \ \& \ \mathcal{A}_2 \leq \mathcal{A}_4 \implies \mathcal{A}_1 \| \mathcal{A}_2 \leq \mathcal{A}_3 \| \mathcal{A}_4$$

Compositional Verification

# Structural Operations: Quotient

Let 
$$\mathcal{A}_1 = (S_1, S_1^0, \text{Tran}_1)$$
 and  $\mathcal{A}_2 = (S_2, S_2^0, \text{Tran}_2)$  be AA.  
Define  $\mathcal{A}_1/\mathcal{A}_2 = (S, S^0, \text{Tran})$ :  
•  $S = 2^{S_1 \times S_2}$   
• write  $S_2^0 = \{s_2^{0,1}, \dots, s_2^{0,p}\}$  and let  $S^0 = \{\{(s_1^{0,q}, s_2^{0,q}) \mid q \in \{1, \dots, p\}\} \mid \forall q : s_1^{0,q} \in S_1^0\}$   
• Tran =

Compositional Verification

# Structural Operations: Quotient

Let 
$$A_1 = (S_1, S_1^0, \text{Tran}_1)$$
 and  $A_2 = (S_2, S_2^0, \text{Tran}_2)$  be AA.  
Define  $A_1/A_2 = (S, S^0, \text{Tran})$ :  
•  $S = 2^{S_1 \times S_2}$   
• write  $S_2^0 = \{s_2^{0,1}, \dots, s_2^{0,p}\}$  and let  $S^0 = \{\{(s_1^{0,q}, s_2^{0,q}) \mid q \in \{1, \dots, p\}\} \mid \forall q : s_1^{0,q} \in S_1^0\}$   
• Tran =

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# Structural Operations: Quotient

Let 
$$\mathcal{A}_1 = (S_1, S_1^0, \text{Tran}_1)$$
 and  $\mathcal{A}_2 = (S_2, S_2^0, \text{Tran}_2)$  be AA.  
Define  $\mathcal{A}_1/\mathcal{A}_2 = (S, S^0, \text{Tran})$ :  
•  $S = 2^{S_1 \times S_2}$   
• write  $S_2^0 = \{s_2^{0,1}, \dots, s_2^{0,p}\}$  and let  $S^0 = \{\{(s_1^{0,q}, s_2^{0,q}) \mid q \in \{1, \dots, p\}\} \mid \forall q : s_1^{0,q} \in S_1^0\}$   
• Tran = ...

#### Theorem

For all AA  $A_1$ ,  $A_2$ ,  $A_3$ :

$$\mathcal{A}_1 \| \mathcal{A}_2 \leq \mathcal{A}_3 \iff \mathcal{A}_2 \leq \mathcal{A}_3 / \mathcal{A}_1$$

• up to  $\equiv$ , / is the adjoint (or residual) of  $\parallel$ 

#### Quantitative Specification Theories?

#### Definition (recall)

A complete specification theory for (Mod,  $\sim$ ) is (Spec,  $\leq$ ,  $\parallel$ ,  $\chi$ ) such that

•  $\leq$  is a refinement preorder on Spec

• 
$$\mathcal{M}_1 \sim \mathcal{M}_2 \iff \chi(\mathcal{M}_1) \leq \chi(\mathcal{M}_2)$$

• (Spec,  $\leq$ ,  $\parallel$ ) forms a b.c.d. residuated lattice up to  $\equiv$ 

• generalize  $\sim$  by pseudometric  $d_{Mod}$ 

• 
$$d_{\mathsf{Mod}}(\mathcal{M}_1, \mathcal{M}_2) = 0$$
 iff  $\mathcal{M}_1 \sim \mathcal{M}_2$ 

• generalize  $\leq$  by hemimetric d

• 
$$d_{\text{Mod}}(\mathcal{M}_1, \mathcal{M}_2) = d(\chi(\mathcal{M}_1), \chi(\mathcal{M}_2))$$

• 
$$d(\mathcal{M},\mathcal{S}) = d(\chi(\mathcal{M}),\mathcal{S})$$

 $\bullet$  still want (Spec,  $\leq, \parallel)$  to be a b.c.d. residuated lattice up to  $\equiv$ 

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Acceptance Automata

For  $DMTS/AA/HML_{max}$ :

- $d_{Mod}$ : any bisimulation distance
- d: corresponding modal refinement distance
- transitivity  $\rightsquigarrow$  triangle ineq.:  $d(\mathcal{S}_1, \mathcal{S}_2) + d(\mathcal{S}_2, \mathcal{S}_3) \geq d(\mathcal{S}_1, \mathcal{S}_3)$
- $d(S, S_1 \land S_2) = \max(d(S, S_1), d(S, S_2))$  or  $\infty$
- $d(\mathcal{S}_1 \lor \mathcal{S}_2, \mathcal{S}) = \max(d(\mathcal{S}_1, \mathcal{S}), d(\mathcal{S}_2, \mathcal{S}))$  or  $\infty$
- quotient is quantitative residual:  $d(S_1||S_2, S_3) = d(S_2, S_3/S_1)$
- for  $\|$  itself, uniform continuity: a function  $P : \mathbb{R}_{\geq 0} \times \mathbb{R}_{\geq 0} \to \mathbb{R}_{\geq 0}$  such that  $d(\mathcal{S}_1 \| \mathcal{S}_2, \mathcal{S}_3 \| \mathcal{S}_4) \leq P(d(\mathcal{S}_1, \mathcal{S}_3), d(\mathcal{S}_2, \mathcal{S}_4))$

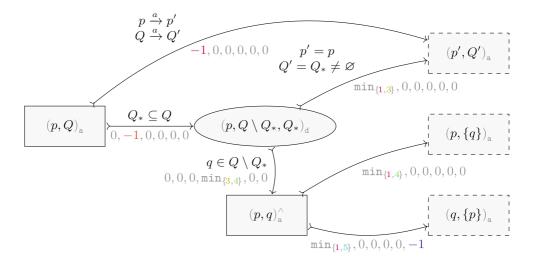
The Bad and/or Ugly QLTBT

# Silent Moves in QLTBT?

Introduction

- Any serious spectrographer needs to think about silent moves
- (van Glabbeek 1993: LTBT II)
- Bisping, Jansen 2023: Energy games for the weak spectrum
  - but uses power set for linear part (recall: we use blindness instead)
  - difficult to reconcile power set with quantitative setting
- otherwise, some coalgebra approaches:
  - Sprunger, Katsumata, Dubut, Hasuo 2021: Fibrational bisimulations and quantitative reasoning
  - Ford, Milius, Schröder, Beohar, König 2022: Graded monads and behavioural equivalence games
  - Beohar, Gurke, König, Messing 2023: Hennessy-Milner theorems via Galois connections
  - again, power set seems very popular ...
- status: IT'S COMPLICATED

94 B. Bisping



**Fig. 7.** Schematic spectroscopy game  $\mathcal{G}_{\bigtriangleup}$  of Definition 10.

QLIBI

Compositional Verification

Bad/Ugly 000●00000000

Conclusion

## Asarin-Basset-Degorre Distance

Recall:

$$D(\sigma,\tau) = \max \begin{cases} \sup_{i} \inf_{j} \{|t_i - s_j| \mid a_i = b_j\} \\ \sup_{j} \inf_{i} \{|t_i - s_j| \mid a_i = b_j\} \end{cases}$$

QLTBT

Introduction

On the practical side, if we observed timed words with some finite precision (say 0.01s), then it would be difficult to distinguish the order of close events, e.g. detect the difference between

 $w_1 = (a, 1), (b, 2), (c, 2.001)$  and  $w_2 = (a, 1.001), (c, 1.999), (b, 2.001).$ 

Moreover, it is even difficult to count the number of events that happen in a short lapse of time, e.g. the words  $w_1, w_2$  look very similar to

$$w_3 = (a, 1), (c, 1.999), (c, 2), (b, 2.001), (c, 2.0002).$$

A slow observer, when receiving timed words  $w_1, w_2, w_3$  will just sense an a at the date  $\approx 1$  and b and c at the date  $\approx 2$ .

As the main contribution of this paper, we introduce a metric on timed words (with non-fixed number of events) for which  $w_1, w_2, w_3$  are very close to each Uli Fahrenberg Quantitative Verification Introduction QLTBT Co

### Asarin-Basset-Degorre Distance

Recall:

$$D(\sigma,\tau) = \max \begin{cases} \sup_{i} \inf_{j} \{|t_i - s_j| \mid a_i = b_j\} \\ \sup_{j} \inf_{i} \{|t_i - s_j| \mid a_i = b_j\} \end{cases}$$

- takes into account permutations of symbols which are close in timing
- but in a way which may lose symbols
- relation to timed pomsets? Amrane, Bazille, Clement, UF 2024: Languages of HDTA
- status: HOPEFUL

Introduction 00000000	QLTBT 0000000000	Compositional Verification	Bad/Ugly 000000●00000	Conclusion 0
Robustness				

A quantitative system is robust if

 small changes in inputs

 small environment perturbations

 small measuring errors

Formulate using uniform continuity: there is a constant K such that

 $d_{ ext{behavior}}(S,S') \leq K \, d_{ ext{syntax}}(S,S')$ 

for all perturbations S' of S.

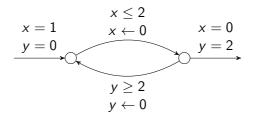
- Standard formulation in control theory
- Generally want systems to be robust

Introduction	QLTBT	Compositional Verification	Bad/Ugly	Conclusion
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Debustness	look of			

#### Robustness, lack of

Our quantitative models are not robust:

(Merci, Nico)



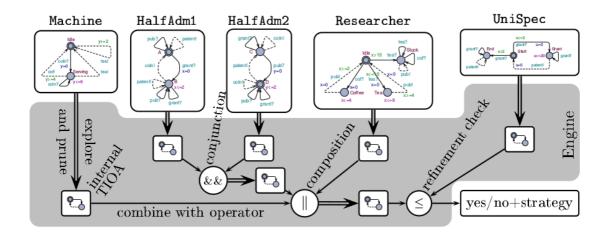
- restrict to robust models?
- quantify robustness?
- use different models?

• . . .

#### Timed input-output automata:

- David, Larsen, Legay, Nyman, Traonouez, Wąsowski 2015: Real-time specifications
- Goorden, Larsen, Legay, Lorber, Nyman, Wąsowski 2023: Timed I/O Automata: It is never too late to complete your timed specification theory
- complete, with quotient, but without disjunction
- only deterministic specifications
- tool support: ECDAR / Uppaal TiGa (Aalborg)
- some work on robustness and implementability: Larsen, Legay, Traonouez, Wąsowski 2014: Robust synthesis for real-time systems

## Timed Input-Output Automata



# Specification Theories for Real-Time Systems, contd.

Modal event-clock specifications:

QLTBT

Introduction

- Bertrand, Legay, Pinchinat, Raclet 2012: Modal event-clock specifications for timed component-based design
- complete, with quotient, but without disjunction
- only deterministic specifications
- some work on robustness: UF, Legay 2012: A robust specification theory for modal event-clock automata

Synchronous time-triggered interface theories:

- Delahaye, UF, Henzinger, Legay, Ničković 2012: Synchronous interface theories and time triggered scheduling
- no quotient, dubious conjunction, no implementation
- relation to **BIP**

Specification Theories for Hybrid Systems

ttion QLTBT Compositional Verifica

Specification Theories for Hybrid Systems

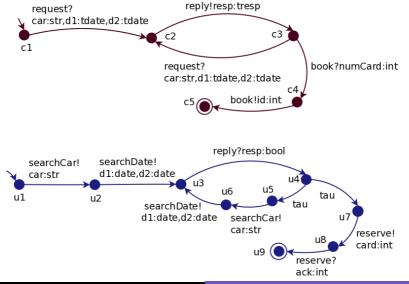
#### • Quesel, Fränzle, Damm 2011: Crossing the bridge between similar games

Introduction	QLTBT	Compositional Verification	Bad/Ugly	Conclusion
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Conclusion				

<ul> <li>general theory of quantitative verification</li> </ul>	✓
<ul> <li>general theory of compositional quantitative verification</li> </ul>	¯\_(シ)_/¯
<ul> <li>algebraic properties</li> </ul>	✓
<ul> <li>quantitative algebraic properties</li> </ul>	×
<ul> <li>silent moves</li> </ul>	×
<ul> <li>for real-time systems</li> </ul>	¯\_(シ)_/¯
<ul> <li>robustness</li> </ul>	¯\_(シ)_/¯
<ul> <li>compositionality</li> </ul>	¯\_(シ)_/¯
<ul> <li>robust compositionality</li> </ul>	×
<ul> <li>for hybrid systems</li> </ul>	×



# Interface Compatibility

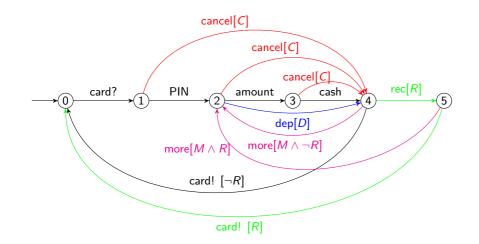


### Interface Compatibility

- Use discounted (bi)simulation distances for measuring interface compatibility
- With A. Legay, M. Ouederni, G. Salaün
  - bisimulation d. for symmetric compatibility
  - ready simulation d. for asymmetric compatibility
- With tool support:

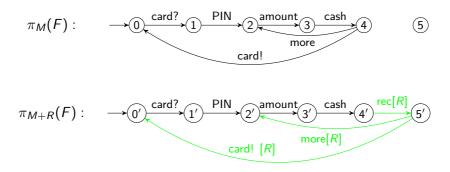
Results of the Compatibility Measure							
Example	medicalSystem						
STSs	MedServer.xml Client.xml						
Compatibility Notion Unspecified Receptions							
			c1	c2	c3	c4	
		s1	1.0	0.06	0.01	0.01	
		s2	0.05	1.0	0.35	0.01	
Compatibility Matrix		s3	0.01	0.26	1.0	0.01	
		s4	0.01	0.01	0.01	1.0	4
		s5	0.01	0.26	0.64	0.01	¥ //
Global Compatibility	1						
Mismatches	Download						

### Behaviour Interactions in Product Lines



#### Behaviour Interactions in Product Lines

- Use a variant of Cantor bisimulation distance for counting the number of behaviour interactions in feature transition systems
- With J. Atlee, S. Beidu, A. Legay
- Use projections to products and compute Cantor bisimulation distance without repetitions:



## Inter-Textual Distances in Statistical NLP

- Use discounted bisimulation distance to measure differences between texts
- With F. Biondi, S. Kongshøj, A. Legay
  - texts are very simple transition systems!
- Implementation
- Works better for some cases than standard distances used in statistical natural language processing
- New collaboration with NLP people in Grenoble

### Inter-Textual Distances in Statistical NLP

- Let  $A = (a_1, a_2, \dots, a_{N_A})$  and  $B = (b_1, b_2, \dots, b_{N_B})$  be texts
- Write  $\delta_{i,j} = [\text{ if } a_i = b_j \text{ then } 0 \text{ else } 1] (word match indicator)$
- position match ( $\lambda$ : discounting factor,  $0 \le \lambda < 1$ ):

$$d_{\mathsf{pm}}(i,j) = \delta_{i,j} + \lambda \delta_{i+1,j+1} + \lambda^2 \delta_{i+2,j+2} + \lambda^3 \delta_{i+3,j+3} + \cdots$$

- "try to match *n*-grams for *n* as high as possible, but don't be too sad if very long phrases don't match"
- global distance:

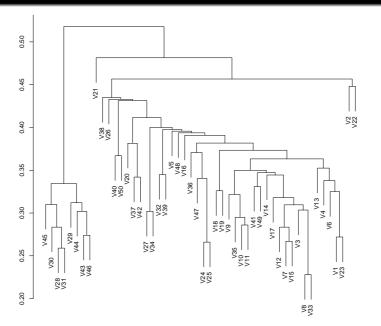
$$d_3(A,B) = \frac{1-\lambda}{N_A} \sum_{i=1}^{N_A} \min_{j=1,\ldots,N_B} d_{pm}(i,j)$$

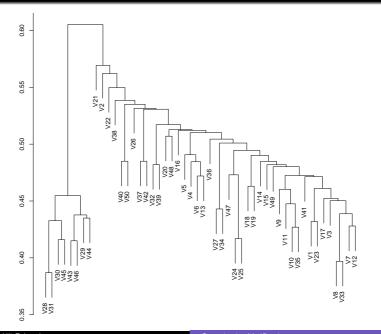
- find best possible match for each position in A, average, and scale
- and symmetrize:  $d_4(A, B) = \max(d_3(A, B), d_3(B, A))$

### Inter-Textual Distances in Statistical NLP

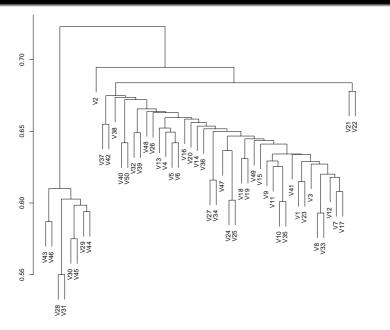
Experiment:

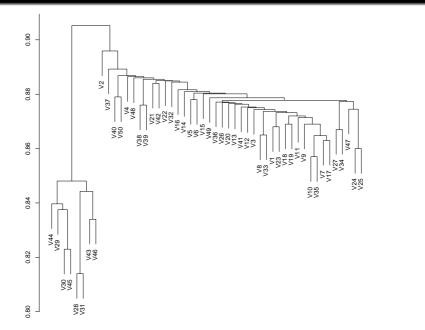
- 42 genuine scientific papers, all about modeling and verification
- 8 automatically generated "fake" papers ("SciGen")





 $\lambda = .4$ 





 $\lambda = .95$ 

#### Uli Fahrenberg

#### Quantitative Verification