

The Quantitative Linear-Time–Branching-Time Spectrum

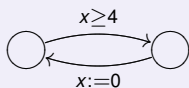
Uli Fahrenberg Axel Legay Claus Thrane

IRISA/INRIA Rennes, France / Aalborg University, Denmark

LaBRI March 2012

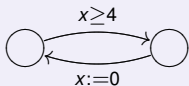
Quantitative Analysis

Quantitative Models



Quantitative Quantitative Analysis

Quantitative *Models*

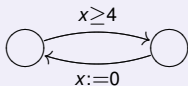


Quantitative *Logics*

$$\Pr_{\leq .1}(\diamond error)$$

Quantitative Quantitative Quantitative Analysis

Quantitative Models



Quantitative Logics

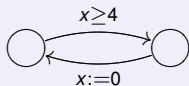
$$\Pr_{\leq .1}(\diamond error)$$

Quantitative Verification

$$\llbracket \varphi \rrbracket (s) = 3.14$$
$$d(s, t) = 42$$

Quantitative Quantitative Quantitative Analysis

Quantitative Models



Quantitative Logics

$$\Pr_{\leq .1}(\diamond error)$$

Quantitative Verification

$$\begin{aligned} \llbracket \varphi \rrbracket (s) &= 3.14 \\ d(s, t) &= 42 \end{aligned}$$

Boolean world

Trace equivalence \equiv

Bisimilarity \sim

$s \sim t$ implies $s \equiv t$

$s \models \varphi$ or $s \not\models \varphi$

$s \sim t$ iff $\forall \varphi : s \models \varphi \Leftrightarrow t \models \varphi$

“Quantification”

Linear distances d_L

Branching distances d_B

$d_L(s, t) \leq d_B(s, t)$

$\llbracket \varphi \rrbracket (s)$ is a quantity

$d_B(s, t) = \sup_{\varphi} d(\llbracket \varphi \rrbracket (s), \llbracket \varphi \rrbracket (t))$

Quantitative Quantitative Quantitative Analysis

Problem: For processes with quantities, lots of different ways to measure distance

- point-wise

$$d_T(\sigma, \tau) = \sup_i |\sigma_i - \tau_i|$$

- accumulating

$$d_T(\sigma, \tau) = \sum_i |\sigma_i - \tau_i|$$

- limit-average

$$d_T(\sigma, \tau) = \limsup_N \frac{1}{N} \sum_{i=0}^N |\sigma_i - \tau_i|$$

- discounting

$$d_T(\sigma, \tau) = \sum_i \lambda^i |\sigma_i - \tau_i|$$

- maximum-lead

$$d_T(\sigma, \tau) = \sup_N \left| \sum_{i=0}^N \sigma_i - \sum_{i=0}^N \tau_i \right|$$

- Cantor

$$d_T(\sigma, \tau) = 1 / (1 + \inf \{j \mid \sigma_j \neq \tau_j\})$$

- etc

Upshot

Two ideas:

- For an application, it is easiest to define distance between **system traces** (executions)
- Use **games** to convert this *linear* distance to *branching* distances

Or:

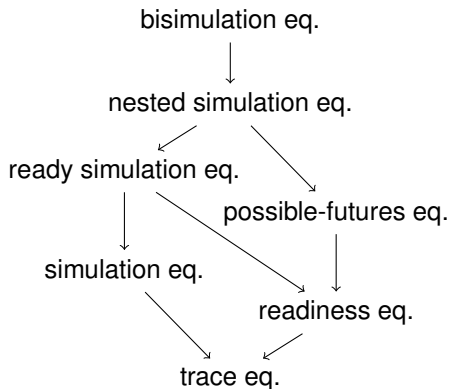
- **If you give us a distance between strings, we give you back a bunch of distances between systems.**

- 1 Background: Quantitative analysis
- 2 The Linear-Time–Branching-Time Spectrum via Games
- 3 From Trace Distances to Branching Distances via Games
- 4 Further Results
- 5 Conclusion

- 1 Background: Quantitative analysis
- 2 The Linear-Time–Branching-Time Spectrum via Games**
- 3 From Trace Distances to Branching Distances via Games
- 4 Further Results
- 5 Conclusion

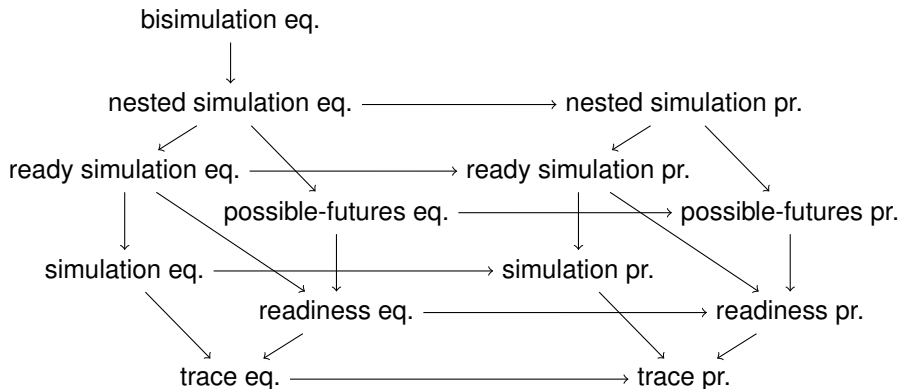
The Linear-Time–Branching-Time Spectrum

van Glabbeek, 2001 (excerpt):



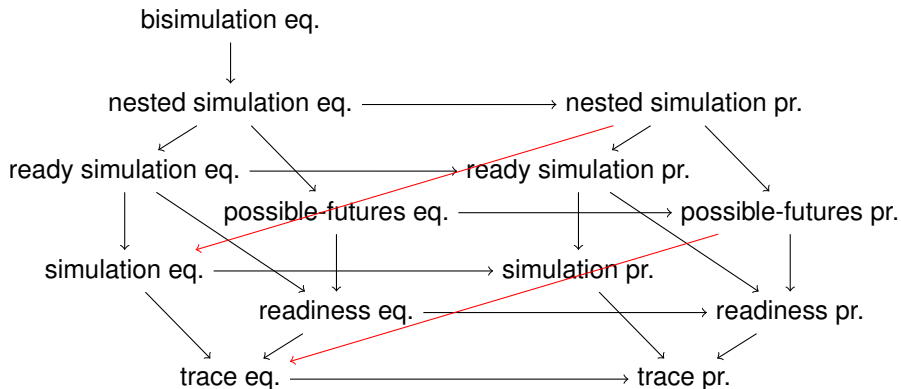
The Linear-Time–Branching-Time Spectrum

van Glabbeek, 2001 (excerpt):

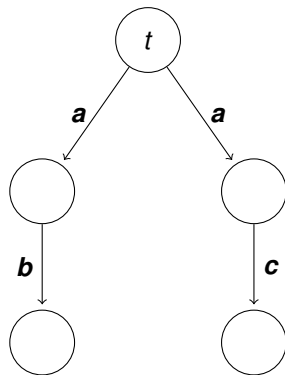
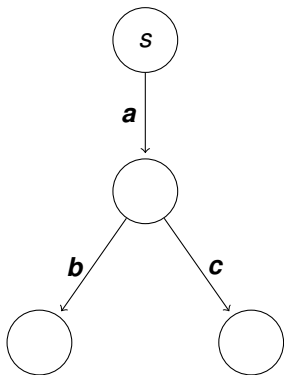


The Linear-Time–Branching-Time Spectrum

van Glabbeek, 2001 (excerpt):

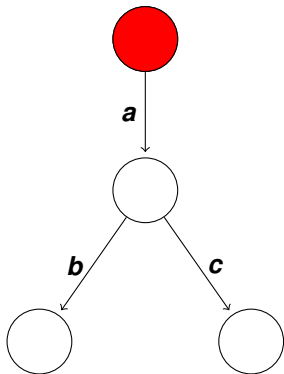


The Simulation Game

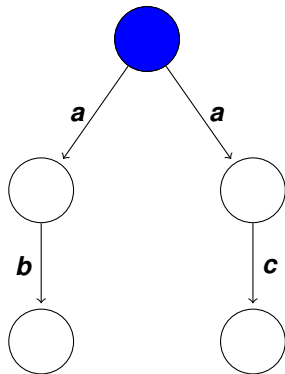


The Simulation Game

Spoiler

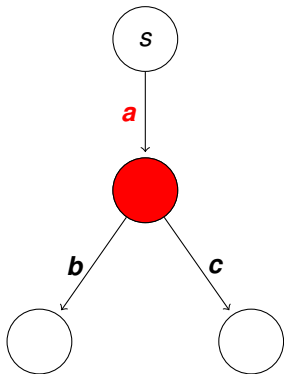


Duplicator

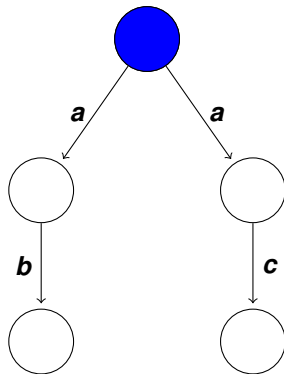


The Simulation Game

Spoiler

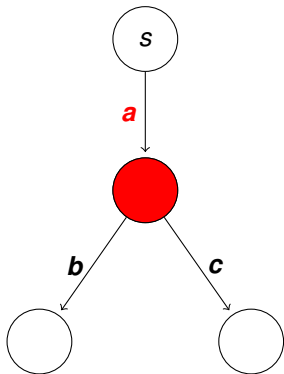


Duplicator

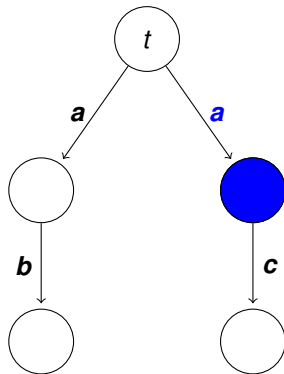


The Simulation Game

Spoiler

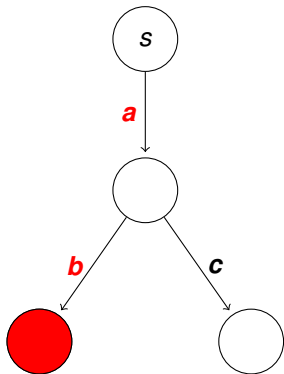


Duplicator

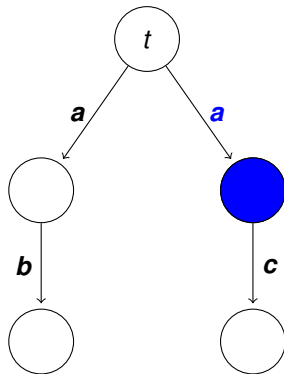


The Simulation Game

Spoiler

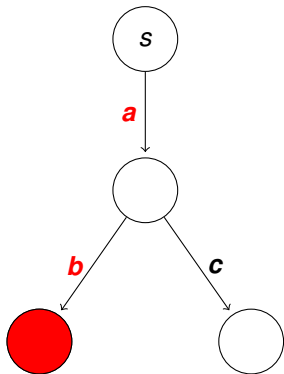


Duplicator

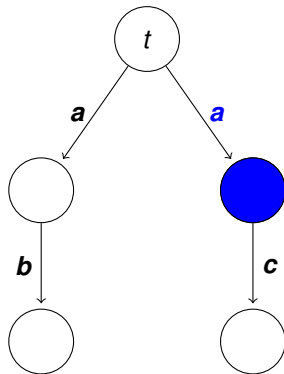


The Simulation Game

Spoiler



Duplicator

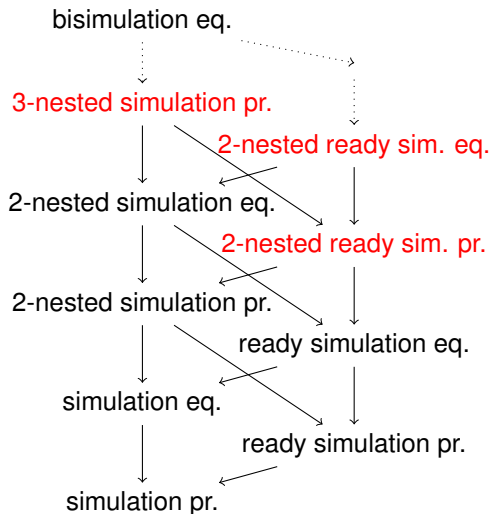


Spoiler wins

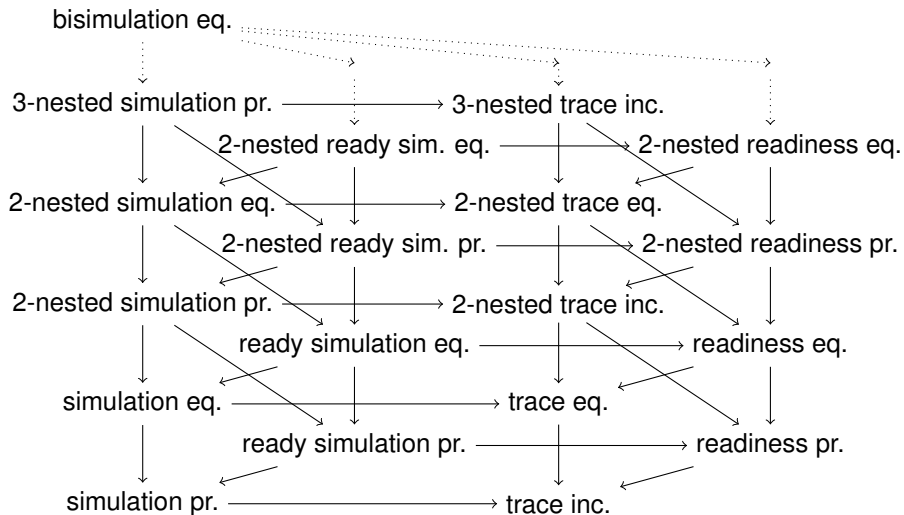
The Simulation Game

1. Player 1 (“**Spoiler**”) chooses edge from s (leading to s')
 2. Player 2 (“**Duplicator**”) chooses matching edge from t (leading to t')
 3. Game continues from configuration s', t'
- ω : If Player 2 can always answer: YES, t simulates s .
Otherwise: NO

The Linear-Time–Branching-Time Spectrum, Reordered



The Linear-Time–Branching-Time Spectrum, Reordered



- 1 Background: Quantitative analysis
- 2 The Linear-Time–Branching-Time Spectrum via Games
- 3 From Trace Distances to Branching Distances via Games**
- 4 Further Results
- 5 Conclusion

The Simulation Game, Revisited

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'
- ω : If Player 2 can always answer: YES, t simulates s .
Otherwise: NO

Or, as an 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 (maybe after infinitely many rounds!), **compare the chosen traces**:
If the trace chosen by t matches the one chosen by s : YES
Otherwise: NO

Quantitative Ehrenfeucht-Fraïssé Games

The quantitative setting:

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

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 $d_T(\sigma, \tau)$

This can be done for all the games in the LTBT spectrum.

Quantitative EF Games: The Gory Details – 1

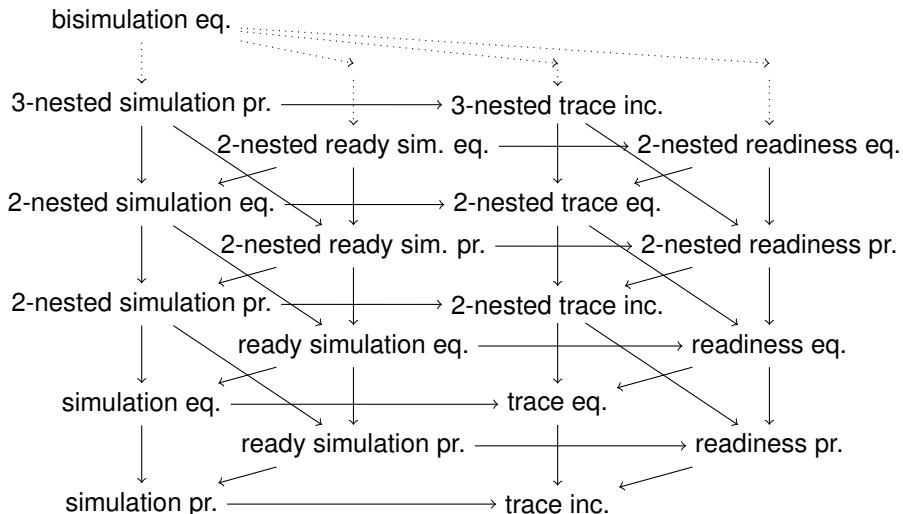
- **Configuration** of the game: (π, ρ) : π the Player-1 choices up to now; ρ the Player-2 choices
- **Strategy**: mapping from configurations to next moves
 - Θ_i : set of Player- i strategies
- **Simulation** strategy: Player-1 moves allowed from **end of π**
- **Bisimulation** strategy: Player-1 moves allowed from end of π or end of ρ
 - (hence π and ρ 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 (σ, τ)
- **Bisimulation distance**: $\sup_{\theta_1 \in \Theta_1} \inf_{\theta_2 \in \Theta_2} d_T(\sigma, \tau)$
- **Simulation distance**: $\sup_{\theta_1 \in \Theta_1^0} \inf_{\theta_2 \in \Theta_2} d_T(\sigma, \tau)$ (**restricting Player 1's capabilities**)

Quantitative EF Games: The Gory Details – 2

- **Blind Player-1 strategies:** depend only on the **end** of ρ
 - (“cannot see Player-2 moves”)
 - $\tilde{\Theta}_1$: set of blind Player-1 strategies
- **Trace inclusion distance:** $\sup_{\theta_1 \in \tilde{\Theta}_1^0} \inf_{\theta_2 \in \Theta_2} d_T(\sigma, \tau)$
- For **nesting:** count the number of times Player 1 choses edge from **end of ρ**
 - Θ_1^k : k choices from end of ρ 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:** $\sup_{\theta_1 \in \tilde{\Theta}_1^1} \inf_{\theta_2 \in \Theta_2} d_T(\sigma, \tau)$
- For **ready:** allow extra “I’ll see you” Player-1 transition from end of ρ

The Quantitative Linear-Time–Branching-Time Spectrum

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



- 1 Background: Quantitative analysis
- 2 The Linear-Time–Branching-Time Spectrum via Games
- 3 From Trace Distances to Branching Distances via Games
- 4 Further Results**
- 5 Conclusion

Transfer Principle

- Given two equivalences or preorders in the *qualitative* setting for which there is a *counter-example* which separates them, then the two corresponding distances are **topologically inequivalent**
- (under certain mild conditions for the trace distance)
- (And the proof uses precisely the same counter-example!)

Recursive Characterization

- If the trace distance $d : (\sigma, \tau) \mapsto d(\sigma, \tau)$ has a decomposition $d = g \circ f : \text{Tr} \times \text{Tr} \rightarrow L \rightarrow \mathbb{R}_{\geq 0} \cup \{\infty\}$ through a complete lattice L ,
- and f has a **recursive formula**
- *i.e.* such that $f(\sigma, \tau) = F(\sigma_0, \tau_0, f(\sigma^1, \tau^1))$ for some $F : \Sigma \times \Sigma \times L \rightarrow L$ (which is *monotone* in the third coordinate)
- (where $\sigma = \sigma_0 \cdot \sigma^1$ is a split of σ into first element and tail)
- **then** all distances in the QLTBT are given as **least fixed points** of some functionals using F

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

Recursive Characterization: Theorem

The endofunction I on $(\mathbb{N}_+ \cup \{\infty\}) \times \{1, 2\} \rightarrow L^{S \times S}$ defined by

$$I(h_{m,p})(s, t) = \begin{cases} \max \begin{cases} \sup_{s \xrightarrow{x} s'} \inf_{t \xrightarrow{y} t'} F(x, y, h_{m,1}(s', t')) \\ \sup_{t \xrightarrow{y} t'} \inf_{s \xrightarrow{x} s'} F(x, y, h_{m-1,2}(s', t')) \end{cases} & \text{if } m \geq 2, p = 1 \\ \sup_{s \xrightarrow{x} s'} \inf_{t \xrightarrow{y} t'} F(x, y, h_{m,1}(s', t')) & \text{if } m = 1, p = 1 \\ \max \begin{cases} \sup_{t \xrightarrow{y} t'} \inf_{s \xrightarrow{x} s'} F(x, y, h_{m,2}(s', t')) \\ \sup_{s \xrightarrow{x} s'} \inf_{t \xrightarrow{y} t'} F(x, y, h_{m-1,1}(s', t')) \end{cases} & \text{if } m \geq 2, p = 2 \\ \sup_{t \xrightarrow{y} t'} \inf_{s \xrightarrow{x} s'} F(x, y, h_{m,2}(s', t')) & \text{if } m = 1, p = 2 \end{cases}$$

has a least fixed point $h^* : (\mathbb{N}_+ \cup \{\infty\}) \times \{1, 2\} \rightarrow L^{S \times S}$, and if the LTS (S, T) is finitely branching, then $d^{k\text{-sim}} = g \circ h_{k,1}^*$ for all $k \in \mathbb{N}_+ \cup \{\infty\}$.

Conclusion & Further Work

- We show how to convert any (typically application-given) distance on system traces to (almost) any type of branching distance in the LTBT spectrum
- “Adding an extra dimension to the LTBT spectrum”
- Application to different scenarios (How does it work in concrete cases? Do we get sensible algorithms? Approximations?)
- Application to real-time and hybrid systems
 - Replace “finitely branching” by “compactly branching”?
- Quantitative LTBT with silent moves?
- What about probabilistic systems?