

The Quantitative Linear-Time–Branching-Time Spectrum

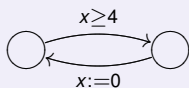
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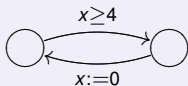
Quantitative Analysis

Quantitative Models



Quantitative Quantitative Analysis

Quantitative *Models*

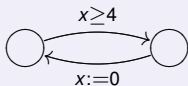


Quantitative *Logics*

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

Quantitative Quantitative Quantitative Analysis

Quantitative Models



Quantitative Logics

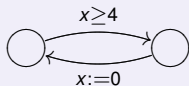
$$\Pr_{\leq .1}(\diamond error)$$

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:

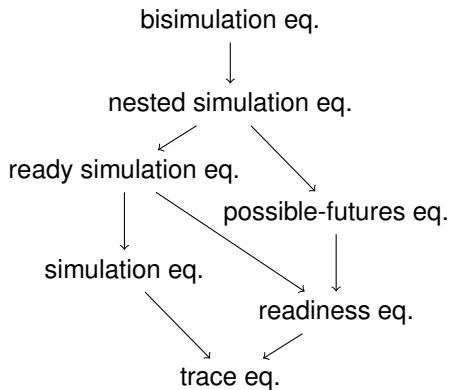
- **If you give us a distance between strings, we give you back a bunch of distances between systems.**

- 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

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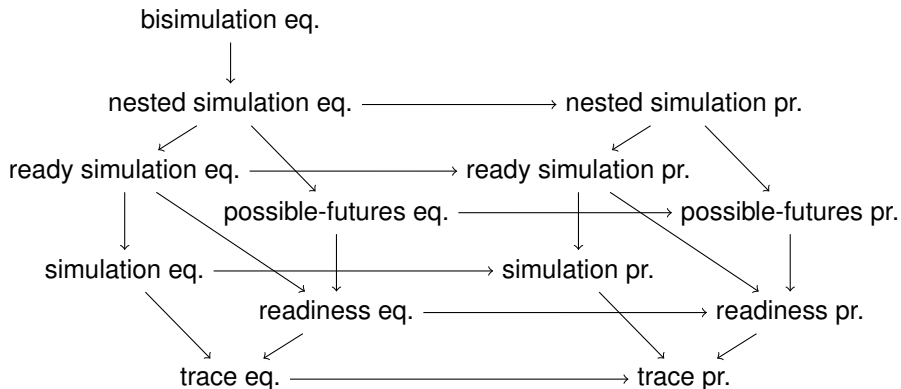
The Linear-Time–Branching-Time Spectrum

van Glabbeek, 2001 (excerpt):



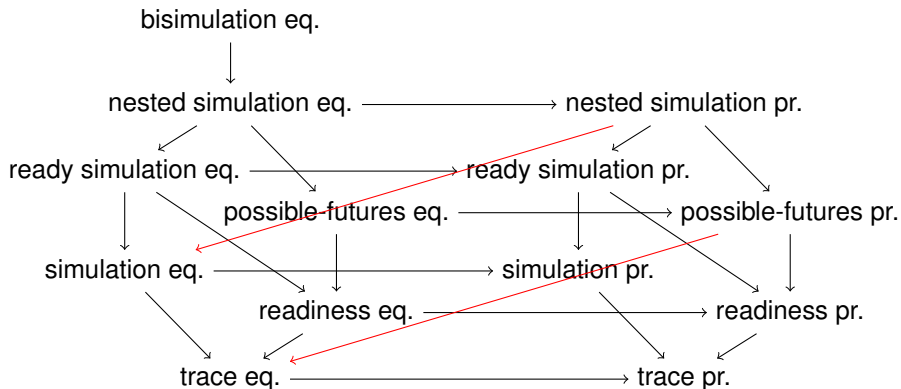
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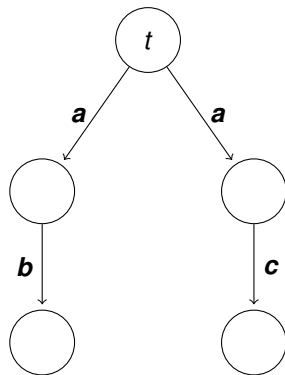
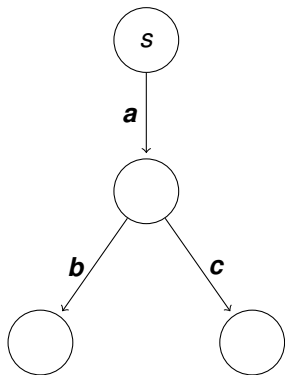


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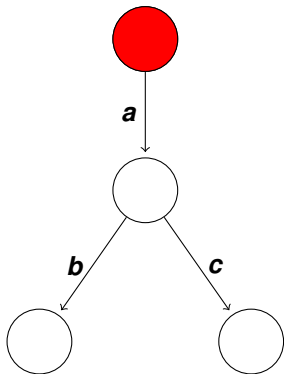


The Simulation Game

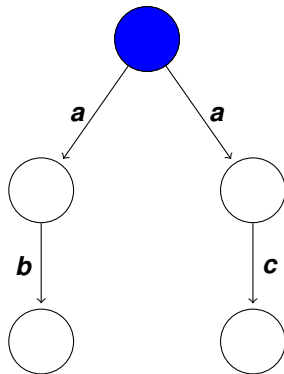


The Simulation Game

Spoiler

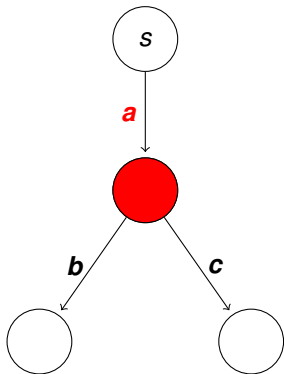


Duplicator

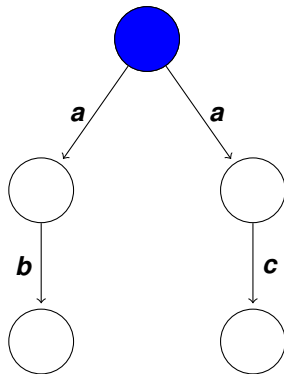


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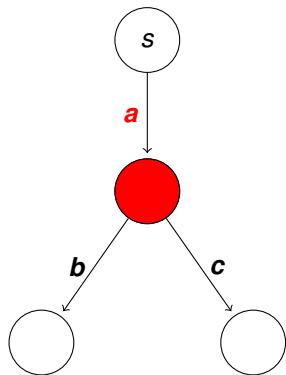


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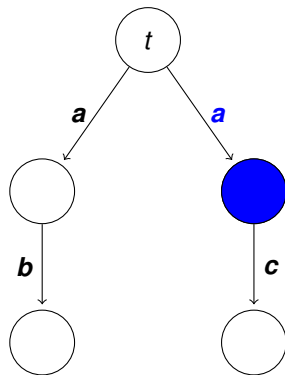


The Simulation Game

Spoiler

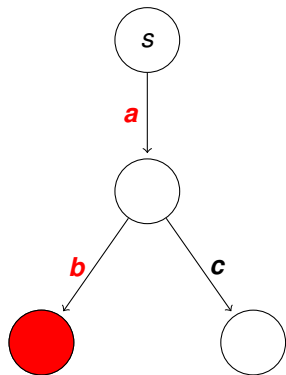


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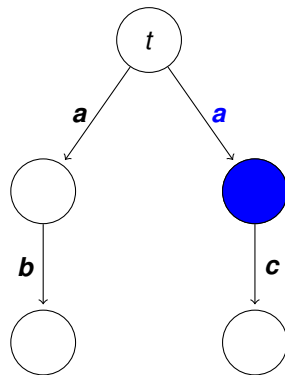


The Simulation Game

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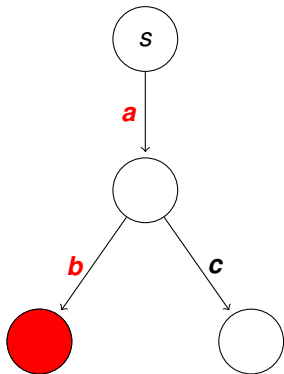


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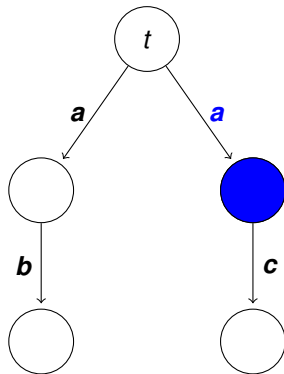


The Simulation Game

Spoiler



Duplicator

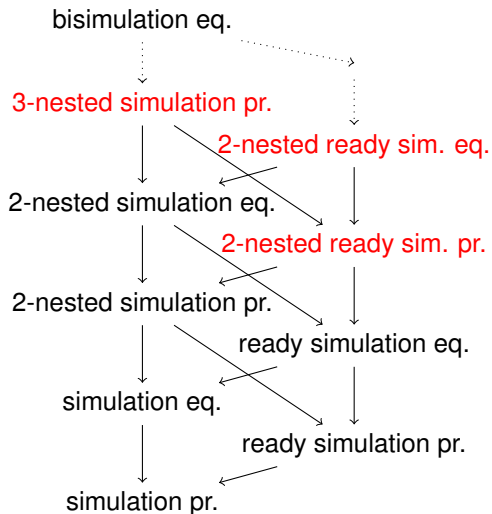


Spoiler wins

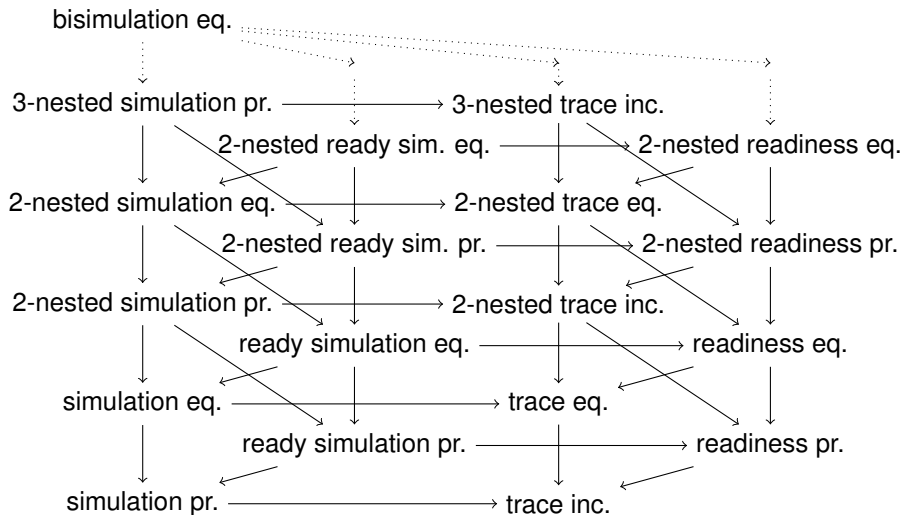
The Simulation Game

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



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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

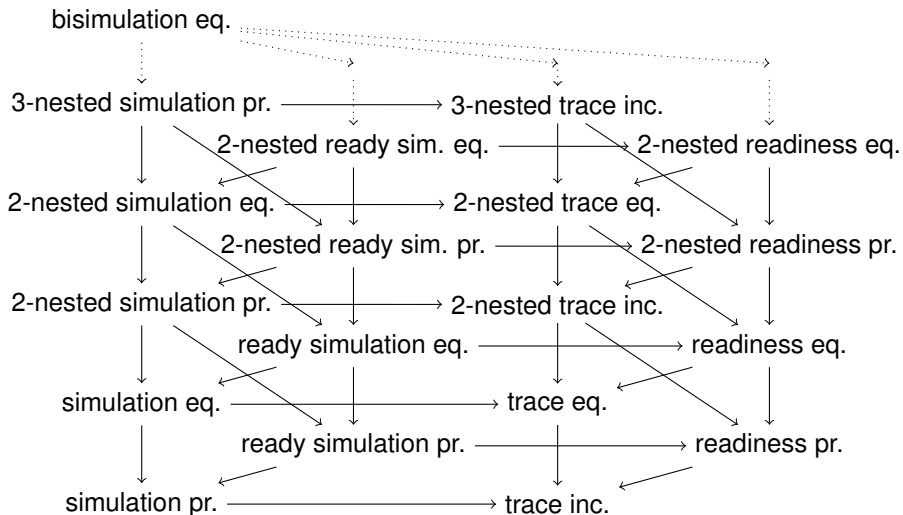
- **Configuration** of the game: (π, ρ) : π the Player-1 choices up to now; ρ the Player-2 choices
- **Strategy**: mapping from configurations to next moves
 - Θ_i : set of Player- i strategies
- **Simulation** strategy: Player-1 moves allowed from **end of π**
- **Bisimulation** strategy: Player-1 moves 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 $d : (\sigma, \tau) \mapsto d(\sigma, \tau) \in \mathbb{R}_{\geq 0} \cup \{\infty\}$:



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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?