

Models and Languages for Computational Systems Biology

Lecture 8: Mix, match and use temporal logics

Ian Stark

School of Informatics
The University of Edinburgh

Thursday 4 February 2010
Semester 2 Week 4



HML

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p$$

Assertions about states in an LTS and possible transitions between them.



Matthew Hennessy and Robin Milner

On observing nondeterminism and concurrency.

In *Automata, Languages and Programming: Proceedings of the Seventh Colloquium ICALP '80*, Lecture Notes in Computer Science 85, pages 299–309. Springer-Verlag, 1980.

HML

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p$$

Assertions about states in an LTS and possible transitions between them.

Hennessy-Milner logic is a branching-time logic for labelled transition systems.

Like CTL, it can talk about all the possible paths out of a state, and also includes information about the labels on transitions.

Unlike LTL and CTL, it does not make assertions about extended runs of programs, but only single steps at a time.

HML

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p$$

Assertions about states in an LTS and possible transitions between them.

Semantics:

$s \models \langle a \rangle p$ There is a transition $s \xrightarrow{a} s'$ for some $s' \models p$.

$s \models [a]p$ For any s' , if $s \xrightarrow{a} s'$ then $s' \models p$.

Note: The $\langle a \rangle$ and $[a]$ echo classic temporal modalities for possibility \diamond and necessity \square ; which are also occasionally used for **A** and **G** in CTL/CTL*; as are \forall and \exists .

HML

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p$$

Assertions about states in an LTS and possible transitions between them.

Some properties:

$$\neg(\langle a \rangle p) = [a](\neg p)$$

$$[a]p \wedge [a]q = [a](p \wedge q)$$

$$\langle a \rangle p \vee \langle a \rangle q = \langle a \rangle(p \vee q)$$

$$\langle a \rangle p \wedge [a]q \Rightarrow \langle a \rangle(p \wedge q)$$

HML

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p$$

Assertions about states in an LTS and possible transitions between them.

Some derived forms:

$$[-]p \stackrel{\text{def}}{=} \forall a \in \text{Label} . [a]p \qquad \langle - \rangle p \stackrel{\text{def}}{=} \exists a \in \text{Label} . \langle a \rangle p$$

$$[-b]p \stackrel{\text{def}}{=} \forall a \in (\text{Label} \setminus b) . [a]p \qquad \langle -b \rangle p \stackrel{\text{def}}{=} \exists a \in (\text{Label} \setminus b) . \langle a \rangle p$$

$$[K]p \stackrel{\text{def}}{=} \forall a \in K \subseteq \text{Label} . [a]p \qquad \langle K \rangle p \stackrel{\text{def}}{=} \exists a \in K \subseteq \text{Label} . \langle a \rangle p$$

$$[-K]p \stackrel{\text{def}}{=} \forall a \in (\text{Label} \setminus K) . [a]p \qquad \langle -K \rangle p \stackrel{\text{def}}{=} \exists a \in (\text{Label} \setminus K) . \langle a \rangle p$$

If the label set is finite then these do not extend HML expressiveness.

HML

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p$$

Assertions about states in an LTS and possible transitions between them.

Examples:

 $s \models [-]\text{False}$

Deadlock

 $s \models \langle - \rangle \text{True}$

s can make some progress

 $s \models \langle - \rangle \text{True} \wedge [-a]\text{False}$

a must happen next

 $s \models \langle a \rangle \text{True} \wedge [a](\langle b \rangle \text{True} \wedge [b][-]\text{False})$

Transition a then b is possible, but will lead to certain deadlock

Modal μ

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p \mid Z \mid \mu X.p \mid \nu Y.p$$

-  Dexter Kozen
Results on the propositional μ -calculus.
Theoretical Computer Science 27(3):333–354.

Modal μ

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p \mid Z \mid \mu X.p \mid \nu Y.p$$

Incorporates propositional variables X, Y, Z, \dots , and two fixpoint operators with the following properties:

$\mu X.p(X)$ is the least M such that $M \subseteq p(M)$

$\nu Y.p(y)$ is the greatest N such that $N \supseteq p(N)$

There is an additional constraint that X and Y must be nested within an even number of negations.

Fixpoint logics are notorious for being incomprehensible. (Bradfield & Stirling, 2001)

Modal μ

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p \mid Z \mid \mu X.p \mid \nu Y.p$$

Some equivalences and examples:

$$\neg(\mu X.p(X)) = \nu Y.\neg(p(\neg Y))$$

$$\mathbf{EF}p = \mu X.(p \vee \langle - \rangle X)$$

$$\mathbf{AG}q = \nu Y.(q \wedge [-]Y)$$

$$\mathbf{E}(p \mathbf{U} q) = \mu X.(q \vee (p \wedge \langle - \rangle X))$$

Modal μ -calculus is sufficiently expressive to encode CTL*, and more besides. (See Bradfield & Stirling 2001 for examples)

Monadic Second-Order Logic

MSOL

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \forall x.p \mid \exists x.p \mid (x \xrightarrow{a} y) \mid Z \mid \forall X.p \mid \exists Y.p$$

Monadic second-order logic is a very general and expressive logic. It uses quantification over states ($\forall x, \exists y$) as well as over sets of states ($\forall X, \exists Y$), which is sufficient to encode everything we have seen so far.

However, in a precise sense, the modal μ -calculus already captures all distinctions in observable behaviour between systems (although particular behaviours might still be simpler to write in MSOL).

In practice, both modal μ and MSOL are regarded as hard to read and write, and computationally challenging to verify; although they may form a powerful backend to some more friendly syntax for describing behaviours.

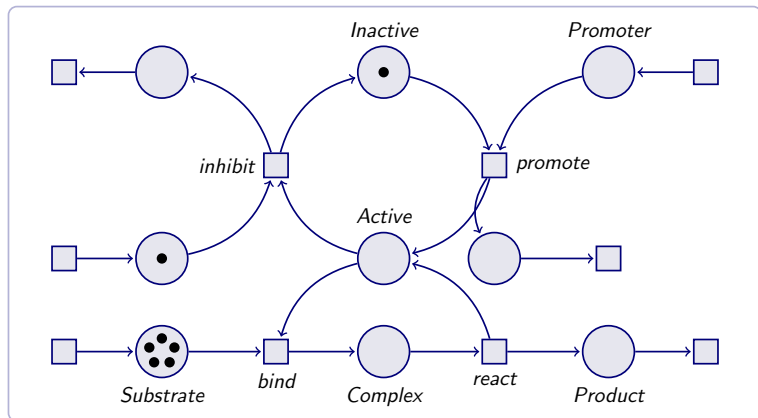
CTL—

$$p ::= \text{Atom} \mid p \wedge q \mid p \vee q \mid p \Rightarrow q \mid \neg p \mid \text{True} \mid \text{False} \\ \mid \langle a \rangle p \mid [a]p \mid \mathbf{AG}p \mid \mathbf{EF}p \mid \dots$$

One way to make temporal logic more conveniently expressive is to incorporate a range of modalities. Some care may be required to maintain feasibility of model-checking — although comprehensibility is perhaps a deeper problem.

As well as mixing various modalities, the underlying (non-modal) part of the logic may be richer: with assertions over real numbers expressing concentrations of reactants, or quantifiers over numeric values, possibly interlaced with the temporal modalities.

Example



$AG[\textit{promote}](\textit{Substrate} \Rightarrow \langle \textit{complex} \rangle \langle \textit{react} \rangle \textit{Product})$

Excerpts from:



F. Fages and A. Rizk.

On the analysis of numerical data time series in temporal logic.

In *Computational Methods in Systems Biology: Proceedings of the International Conference CMSB 2007*, LNBI 4695, pp. 48–63.

Springer-Verlag, 2007.

Reachability : $F([A] \geq p)$, what threshold p species A attain in the trace ?

Checkpoints : $\text{not } (([A] < p_1) \cup ([B] > p_2))$, for which thresholds p_1 and p_2 is it false that $[A]$ is lower than p_1 until $[B]$ is above p_2 , i.e., for which p_1 and p_2 $[A] \geq p_1$ is a checkpoint of $[B] > p_2$?

Stability : $G([A] = < p_1 \ \& \ [A] \geq p_2)$, what is the range of values taken by $[A]$? This range can be looked for in some context given by a condition like in $G(\text{Time} > 10 \rightarrow ([A] < p_1 \ \& \ [A] > p_2))$.

Oscillation : $F((d([A])/dt > 0 \ \& \ [A] > v_1) \ \& \ (F((d([A])/dt < 0 \ \& \ [A] < v_2))))$, what amplitude $(v_1 - v_2)$ is attained in at least one oscillation ? An oscillation is defined as the change of sign of the derivative. This formula can be extended for more oscillations and is abbreviated by `oscil(M,K,p)`. It states that M must have amplitude P in at least K oscillations. By applying the algorithm for each value of K , beginning with 1, we can find the number of oscillations in the trace and minimal amplitude P attained by K oscillations for any K .

Influence : $G(d[A]/dt > p_1 \rightarrow d^2[B]/dt^2 \geq 0)$, above which threshold does the derivative of A have an influence on B ? The influence is positive if a high value of $d[A]/dt$ entails a positive second derivative of $[B]$. It is worth noticing that, as multiple species might influence B , this formula only indicates a correlation between the value of the derivative of A and the second derivative of B and gives no proof of direct influence.

For stability, let us find the range of values taken by [Cdc2] in the last third part of the trace:

```
biocham: trace_analyze(G(Time>66 -> ([Cdc2]=<v1 & [Cdc2]>=v2))).  
[[v1>=0.479, v2=<0.338]]
```

The domain is defined by the conjunction of the two constraints $v1 \geq 0.479$ and $v2 \leq 0.338$. These values are the maximum and minimum values attained by [Cdc2] in the last third part of the trace. The results for the other species are given in Table [1](#).

An oscillation query may compute several interval domains:

```
biocham: trace_analyze(oscil(Cdc2,1)).  
[[v2>=0.338, v1=<0.479], [v2>=0.341, v1=<0.479]]
```

The result is the union of two boxes. In such domains, the most relevant point is not obvious. Here we look for the maximum amplitude $v1 - v2$. The maximum is obtained in the domain with $v1 - v2 = 0.479 - 0.338 = 0.141$. This result states that at least one oscillation of Cdc2 has an amplitude greater or equal to 0.141. The number of oscillations is then incremented until obtaining an empty validity domain. It is obtained for Cdc2 with the query `oscil(Cdc2,3)`, stating that there are only two oscillations of Cdc2 in the trace.



P. T. Monteiro, D. Ropers, R. Mateescu, A. T. Freitas, and H. de Jong.

Temporal logic patterns for querying dynamic models of cellular interaction networks.

Bioinformatics, 24(16):227–233, 2008.



H. de Jong.

Qualitative modeling and simulation of bacterial regulatory networks.

Computational Methods in Systems Biology: Proceedings of the 6th International Conference, CMSB 2008, Lecture Notes in

Bioinformatics 5307, Springer-Verlag, 2008

(with slides)

References (1/2)

These are the papers introducing CTL*, HML, and the modal μ -calculus, respectively.

 E. Allen Emerson and Joseph Y. Halpern

“Sometimes” and “not never” revisited: on branching versus linear time temporal logic.

Journal of the ACM 33(1):151–178.

 Matthew Hennessy and Robin Milner

On observing nondeterminism and concurrency.

In *Automata, Languages and Programming: Proceedings of the Seventh Colloquium ICALP '80*, Lecture Notes in Computer Science 85, pages 299–309. Springer-Verlag, 1980.

 Dexter Kozen

Results on the propositional μ -calculus.

Theoretical Computer Science 27(3):333–354.

References (2/2)

Bradfield and Stirling give a good overview of temporal logics and model-checking; Vardi argues for the comprehensibility of LTL over CTL; and Janin and Walukiewicz show that the modal μ -calculus is all you need for observable behaviour.



Julian Bradfield and Colin Stirling

Modal logics and mu-calculi: an introduction.

In *Handbook of Process Algebra*, pages 293–330. Elsevier, 2001.



Moshe Vardi

Branching vs. Linear Time: Final Showdown

In *Proc. TACAS 2001*, LNCS 2031, pp. 1–22. Springer-Verlag, 2001.



David Janin and Igor Walukiewicz

On the expressive completeness of the propositional mu-calculus with respect to monadic second-order logic.

In *Proc. Concur '96*, LNCS 1119, pp. 263–277. Springer-Verlag, 1996.