

- 1 Discrete models for gene networks according to René Thomas
- 2 CTL
- 3 Techniques of software testing
- 4 General Schema for BRN
- 5 Genetically modified Hoare logic, and examples
  - Hoare Logic
  - Examples

$$\begin{array}{l}
 \{(x = x_0) \wedge (y = y_0)\} \\
 aux := x ; \\
 x := y ; \\
 y := aux \\
 \{(y = x_0) \wedge (x = y_0)\}
 \end{array}
 \quad
 \frac{}
 {
 \frac{}
 {
 \frac{}
 {
 \{Q[v \leftarrow expr]\} \quad v := expr \quad \{Q\}
 }
 }
 }
 }
 }
 \frac{}
 {
 \frac{}
 {
 \frac{}
 {
 \{P\}p_1\{Q'\} \quad \{Q'\}p_2\{Q\}
 }
 }
 }
 }
 }
 \frac{}
 {
 \{P\}p_1; p_2\{Q\}
 }
 }
 ;$$


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$$\frac{}
 {
 \frac{}
 {
 \frac{}
 {
 \{Q_3\}a_1\{Q_2\} \quad \{Q_2\}a_2\{Q_1\}
 }
 }
 }
 }
 \frac{}
 {
 \{P\}a_1; a_2\{Q_1\}
 }
 }
 ;
 \frac{}
 {
 \{Q_1\}a_3\{Q\}
 }
 }
 ;
 \frac{}
 {
 \{P\}a_1; a_2; a_3\{Q\}
 }
 }
 ;$$

$$\{(x = x_0)\} \leftarrow \left\{ \left( \frac{}
 {
 \frac{}
 {
 \frac{}
 {
 (x < 0) \wedge (-x \geq 0) \wedge ((-x)^2 = x_0^2)
 }
 }
 }
 }
 }
 \wedge \right) \vee \left( \frac{}
 {
 \frac{}
 {
 \frac{}
 {
 (x \geq 0) \wedge (x \geq 0) \wedge (x^2 = x_0^2)
 }
 }
 }
 }
 }
 \wedge \right) \right\}$$

```

if (x!=0) :
  r := -x
else :
  r := x
  
```

$$\{(r \geq 0) \wedge (r^2 = x_0^2)\}$$


---


$$\frac{}
 {
 \frac{}
 {
 \{Q_1\}p_1\{Q\} \quad \{Q_2\}p_2\{Q\}
 }
 }
 }
 \frac{}
 {
 \{(e \wedge Q_1) \vee (\neg e \wedge Q_2)\} \text{ if } e \text{ then } p_1 \text{ else } p_2 \{Q\}
 }
 }
 \text{if}$$

**Also :**

While loop :  $\frac{\{e \wedge I\}p\{I\} \quad (\neg e \wedge I) \Rightarrow Q}{\{I\} \text{ while } e \text{ with } I \text{ do } p\{Q\}}$

Empty program :  $\frac{P \Rightarrow Q}{\{P\} \varepsilon \{Q\}}$  use sparingly : loses weakest precondition !

$$\{(p = p_0) \wedge (q = q_0)\}$$

```

r := 0 ;
while p >= q with {q=q_0 & p >= 0 & p_0 = p + r.q_0} :
  p := p - q
  r := r + 1
  
```

$$\{0 \leq p_0 - r.q_0 < q_0\}$$


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$$\frac{}
 {
 \frac{}
 {
 \frac{}
 {
 \{Q[v \leftarrow expr]\} \quad v := expr \quad \{Q\}
 }
 }
 }
 }
 \frac{}
 {
 \frac{}
 {
 \frac{}
 {
 \{P\}p_1\{Q'\} \quad \{Q'\}p_2\{Q\}
 }
 }
 }
 }
 \frac{}
 {
 \{P\}p_1; p_2\{Q\}
 }
 }
 ;
 \frac{}
 {
 \frac{}
 {
 \{e \wedge I\}p\{I\} \quad \neg e \wedge I \Rightarrow Q
 }
 }
 }
 \frac{}
 {
 \{I\} \text{ while } e \text{ with } I \text{ do } p \{Q\}
 }
 }
 \text{loop}$$

Terms :  $v$  gene |  $n \in \mathbb{N}$  |  $K_{v,\{...\}}$  parameter symbols | + | -

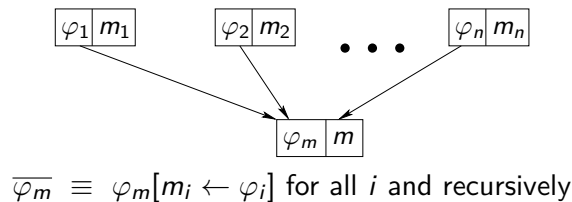
atoms :  $t \geq t'$  |  $t < t'$  |  $t = t'$  | ...

Connectives :  $\neg$  |  $\wedge$  |  $\vee$  |  $\implies$

Example :

$$(a \leq 3 \wedge d + 1 < K_{d,\{m,c\}}) \vee (K_{d,\{c\}} < K_{d,\{m,c\}} \wedge c \geq 3)$$

From multiplexes to assertions : flattening



$\omega$  is the set of ressources of  $v$  :

$$\Phi_v^\omega \equiv \left( \bigwedge_{m \in \omega} \overline{\varphi_m} \right) \wedge \left( \bigwedge_{m \in G^{-1}(v) \setminus \omega} \neg \overline{\varphi_m} \right)$$

$v$  can increase :

$$\Phi_v^+ \equiv \bigwedge_{\omega \subset G^{-1}(v)} (\Phi_v^\omega \implies K_{v,\omega} > v)$$

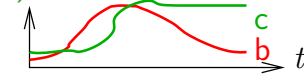
$v$  can decrease :

$$\Phi_v^- \equiv \bigwedge_{\omega \subset G^{-1}(v)} (\Phi_v^\omega \implies K_{v,\omega} < v)$$

- $x+ \mid x- \mid x := n \mid \text{assert}(\varphi)$
- $p_1; p_2; \dots; p_n$
- if  $\varphi$  then  $p_1$  else  $p_2$
- while  $\varphi$  with  $\psi$  do  $p$
- $\forall(p_1, p_2, \dots, p_n)$
- $\exists(p_1, p_2, \dots, p_n)$

Examples :

- $b+; c+; b-$
- $\exists(b+, b-, c+, c-, \varepsilon)$
- while  $(b < 2)$  with  $(c > 0)$  do  $\exists(b+, b-, \forall((c-; a-), c+))$  od;  $b-$



Incrementation rule :  $\frac{}{\{ \Phi_v^+ \wedge Q[v \leftarrow v + 1] \} v+ \{ Q \}}$

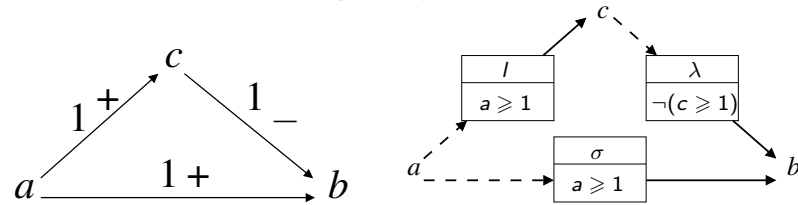
Decrementation rule :  $\frac{}{\{ \Phi_v^- \wedge Q[v \leftarrow v - 1] \} v- \{ Q \}}$

Assertion rule :  $\frac{}{\{ \varphi \wedge Q \} \text{assert}(\varphi) \{ Q \}}$

Universal quantifier rule :  $\frac{\{ P_1 \} p_1 \{ Q \} \quad \{ P_2 \} p_2 \{ Q \}}{\{ P_1 \wedge P_2 \} \forall(p_1, p_2) \{ Q \}}$

Existential quantifier rule :  $\frac{\{ P_1 \} p_1 \{ Q \} \quad \{ P_2 \} p_2 \{ Q \}}{\{ P_1 \vee P_2 \} \exists(p_1, p_2) \{ Q \}}$

Uri Alon most frequent regulatory network patterns



Behaviour of  $b$  after switching  $a$  from off to on ?

Simple off→on→off behaviour of  $b$  with the help of  $c$  :

$$\{(a = 1 \wedge b = 0 \wedge c = 0)\} b+ ; c+ ; b- \{b = 0\}$$

possible if and only if :

$$K_{b,\{\sigma,\lambda\}} = 1 \wedge K_{c,\{l\}} = 1 \wedge K_{b,\{a\}} = 0$$

off→on→off behaviour of  $b$  without the help of  $c$  :

$$\{(a = 1 \wedge b = 0 \wedge c = 0)\} b+ ; b- \{b = 0\}$$

$$\left. \begin{array}{l} b = 0 \\ ((c \geq 1) \wedge (a < 1)) \implies ((K_b = 1) \wedge (K_b = 0)) \\ ((c \geq 1) \wedge (a \geq 1)) \implies ((K_{b,\sigma} = 1) \wedge (K_{b,\sigma} = 0)) \\ ((c < 1) \wedge (a < 1)) \implies ((K_{b,\lambda} = 1) \wedge (K_{b,\lambda} = 0)) \\ ((c < 1) \wedge (a \geq 1)) \implies ((K_{b,\sigma\lambda} = 1) \wedge (K_{b,\sigma\lambda} = 0)) \end{array} \right\} \text{not satisfiable !}$$

Although  $b+ ; c+ ; b-$  is possible, if  $c$  becomes "on" before  $b$ , then  $b$  will never be able to get "on"

Proof by refutation :

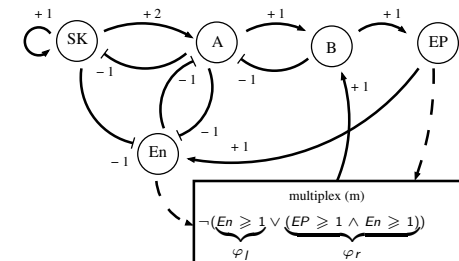
$$\left\{ \begin{array}{l} a = 1 \wedge b = 0 \wedge c = 1 \wedge \\ K_{b,\sigma\lambda} = 1 \wedge K_{c,l} = 1 \wedge K_{b,\sigma} = 0 \end{array} \right\}$$

while  $b < 1$  with  $l$  do  $\exists (b+, b-, c+, c-)$

$$\left\{ \begin{array}{l} b = 1 \end{array} \right\}$$

the triple is inconsistent, whatever the loop invariant  $l$  !

- A 22 gene model reduced to 5 variables using multiplexes



SK = Cyclin E/Cdk2, Cyclin H/Cdk7  
A = Cyclin A/Cdk1  
B = Cyclin B/Cdk1  
En = APC<sup>G1</sup>, CKI (p21, p27), Wee1  
EP = APC<sup>M</sup>, Phosphatases

- 48 states, 26 parameters, 339 738 624 possible valuations, 12 trace specifications and a few temporal properties

- 13 parameters have been entirely identified (50%) and only 8192 valuations remain possible according to the generated constraints (0.002%)
- Additional reachability constraints (e.g. endoreplication and quiescent phase) have been necessary, on an extended *hybrid* extension of the Thomas' framework, to identify (almost) all parameters
- This initial Hoare logic identification step was crucial : it gave us the sign of the derivatives in all the (reachable) states

- If there is a proof tree for  $\{P\}p\{Q\}$  then for each initial state satisfying  $P$ , there are traces in the regulatory network that realize the trace specification  $p$ , and for all of them, if terminating, they satisfy  $Q$  at the end.
- If for each initial state satisfying  $P$  there are traces that realize  $p$  in the regulatory network and if they all satisfy  $Q$  at the end, then there exists a proof tree for  $\{P\}p\{Q\}$ .
- There is a simple algorithm to compute, for each  $Q$ , the minimal loop invariant  $I$  such that  $\{I\}while\ e\ with\ I\ do\ p\{Q\}$ .  
(However well chosen slightly non minimal invariants can considerably simplify the proof tree...)

## Make explicit the hypotheses that motivate the biologist

A far as possible formalize them to get a computer aided approach  
Behavioural *properties* are as much important as *models*  
Mathematical models are not reality : let's use this freedom !  
(several views of a same biological object)  
Modelling is significant only with respect to the considered experimental *reachability* and *observability* (for refutability)

## Formal proofs can suggest wet experiments

“Kleenex” models help understanding main behaviours

Specialized qualitative approaches can make complex models simple

The more detailed models are not the more comprehensible ones