

# Tropical Equilibration by Constraint Solving

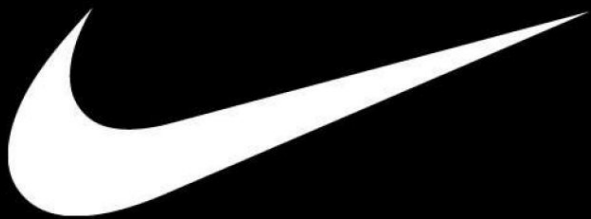
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# ODE System Reduction by Tropical Equilibration

- 1 rescale parameters using a small  $\epsilon$   
(round values)
- 2 search for a rescaling of variables  $x_i = \epsilon^{a_i} \bar{x}_i$   
 $a_i \in \mathbb{Z}$  is an unknown
- 3 equilibrate the degrees of *at least* the biggest positive and the biggest negative term in some/all right sides of the ODEs
- 4 take care of conservation laws
- 5 profit!



## CSP over integers

Express the problem as constraints over the  $a_i$

Note: degrees are linear combinations of the  $a_i$

We will need constraints of the form  $E_1 < E_2$

And **reified constraints** of the form  $B \Leftrightarrow (E_1 = E_2)$

Use a *Constraint Solver* to get solutions (reduce domains by **propagation**, if necessary **enumerate** over domains)

$x_1$  is Cdc2,  $x_2$  Cdc2~{p},  $x_3$  MPF and  $x_4$  preMPF

$$\frac{\partial x_1}{\partial t} = k_9 x_2 - k_8 x_1 + k_6 x_3$$

$\Rightarrow$

$$\frac{\partial \bar{x}_1}{\partial t} = \epsilon^{-3+a_2-a_1} \bar{k}_9 \bar{x}_2 - \epsilon^{-6} \bar{k}_8 \bar{x}_1 + \epsilon^{a_3-a_1} \bar{k}_6 \bar{x}_3$$

**Do not enumerate!**

```
equilibrate(LPos, LNeg) :-  
    get_min(LPos, Min),  
    get_min(LNeg, Min).
```

# Conservations

$$x_1 + x_2 + x_3 + x_4 = 1$$

⇒

$$\epsilon^{a_1} \bar{x}_1 + \epsilon^{a_2} \bar{x}_2 + \epsilon^{a_3} \bar{x}_3 + \epsilon^{a_4} \bar{x}_4 = \epsilon^0$$

Ensure proper limit when  $\epsilon \rightarrow 0$

conserve(L, D) :-

get\_min\_and\_count(L, D, C),

C #>= 2.

## Actually...

```
get_min(L, M) :-  
    get_min_and_count(L, M, C),  
    C #>= 1.
```

```
get_min_and_count([H | T], M, C) :-  
    M #=< H,  
    B #<==> M #= H,  
    C #= B + CC,  
    get_min_and_count(T, M, CC).
```

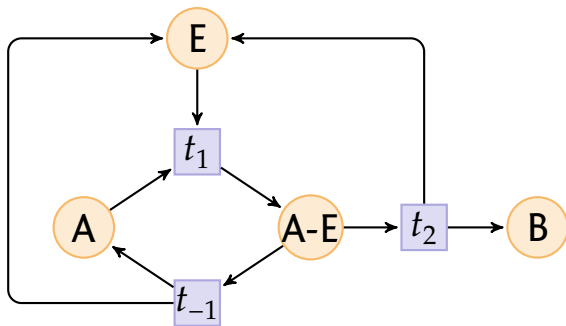
```
get_min_and_count([], _, 0).
```

But how do we get conservation laws?



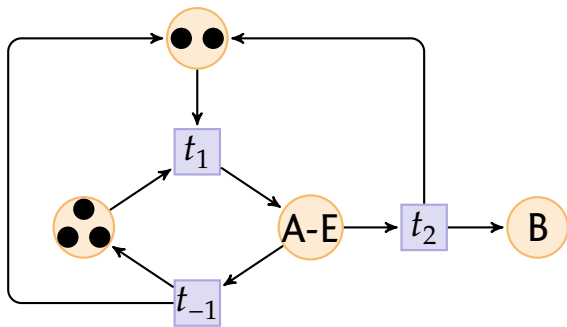
## Petri-net view of a reaction model

Species are places and reactions are transitions, thus  $A + E \rightleftharpoons A-E \Rightarrow B + E$  corresponds to the following Petri-net:



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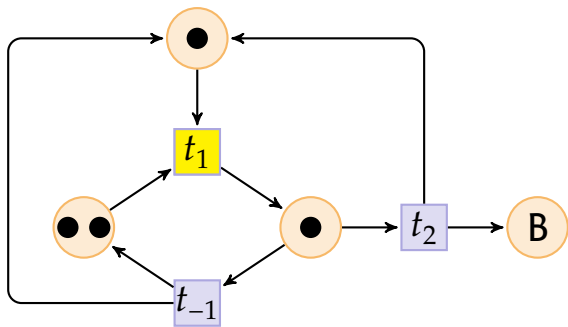
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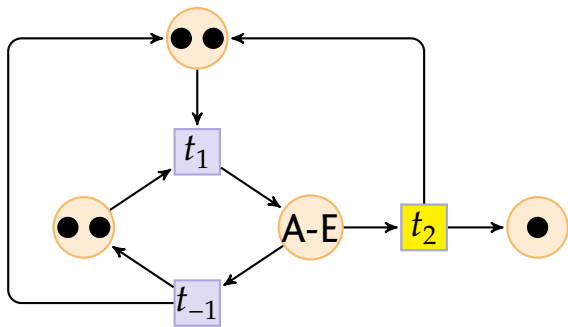
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## Petri-net view of a reaction model

Species are places and reactions are transitions, thus  $A + E \rightleftharpoons A-E \Rightarrow B + E$  corresponds to the following Petri-net:



Consume a token in A and E to produce one in A-E ( $t_1$ ), then do the exact opposite ( $t_{-1}$ ) or remove the A-E token to add one B and one E ( $t_2$ ).

## P-invariants

A (minimal) P-invariant is a marking (with minimal support) of a Petri-net, representing a **weighted sum invariant by all transitions**.

$$V \cdot M = 0$$

A P-invariant defines a **conservation law**, whatever the semantics and the dynamics. There are other ones, e.g.:

**MA** ( $k_1$ ) **for**  $\_ = [A] \Rightarrow B$ .  
**MA** ( $k_2$ ) **for**  $\bar{A} \Rightarrow \_$ .

when  $k_1 = k_2$

## Finding (minimal) P-invariants as a CSP

For a Petri net with  $p$  places and  $t$  transitions, a P-invariant is a vector  $V \in \mathbb{N}^p$  s.t.

$$\forall 1 \leq i \leq t \quad V \cdot L_i = V \cdot R_i$$

$\Rightarrow t$  (linear) equality constraints on  $p$  Finite Domain variables

Non trivial iff  $V \cdot \mathbf{1} > 0$

Ensure minimality by (labelling from small to big and) **branch and bound** on a partial base  $\mathcal{B}$  of vectors:  $\forall B \in \mathcal{B} \quad \prod_{B_i \neq 0} V_i = 0$

Removing some subsumed P-invariants remains necessary.

## Example

Using the Petri-net of our example we have:

$A + E \Rightarrow A - E$	$A + E = AE$	(1)
$A - E \Rightarrow A + E$	$AE = A + E$	(2)
$A - E \Rightarrow B + E$	$AE = B + E$	(3)

where obviously equation (2) is redundant.

We add  $A + E + AE + B > 0$ .

And find **two minimal semi-positive P-invariants**:

- $E = AE = 1$  and  $A = B = 0$
- $A = B = AE = 1$  and  $E = 0$

## Exponential blow-up

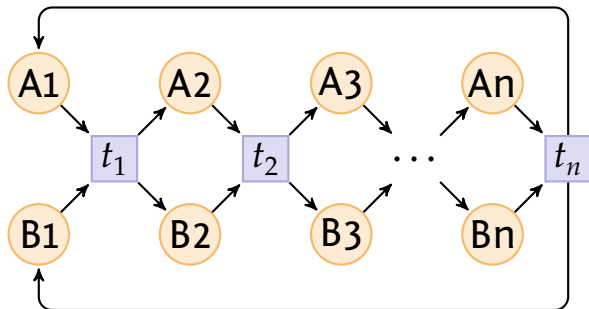
Bad news, the problem actually has an **exponential** number of solutions

$$A_1 + B_1 \Rightarrow A_2 + B_2$$

$$A_2 + B_2 \Rightarrow A_3 + B_3$$

...

$$A_n + B_n \Rightarrow A_1 + B_1$$



Use CSPs symmetry breaking  
(similar to *Equality classes* optimization for FM)



## Evaluation

Minimal semi-positive P-invariant computation on big models of biochemical reaction networks is quite efficient: **all < 1s**

Model	$t$	$p$	#	Invariant size
Schoeberl's MAPK [N. Biotech 02]	125	105	14	from 2 to 44 ODE model!
Curie's E2F/Rb [MSB 08]	~500	~400	79	from 1 (EP300) to ~230 (E2F1 box)
Kohn's map [MBC 99]	~800	~500	65	from 1 (Myt1) to ~200 (e.g., cdk2)

All curated models of `biomodels.net` also < 1s

## Conclusion

- + Conservation law computation is a CSP
- + Tropical Equilibration is a CSP
- + Reasonable performance
- Not symbolic
- Rounding errors

“If debugging is the process of removing software bugs, then programming must be the process of putting them in.”

– Edsger Dijkstra