

NoPain – Meeting

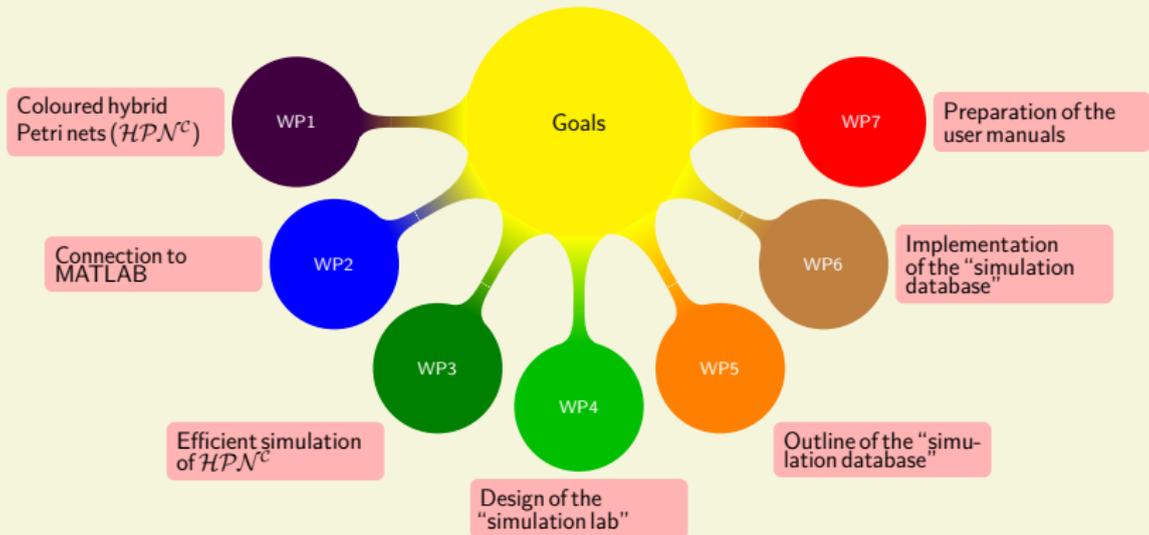
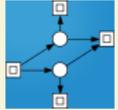
Monika Heiner Christian Rohr

Department of Computer Science
Brandenburg University of Technology Cottbus

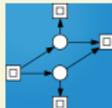
<http://www-dssz.informatik.tu-cottbus.de>

Jan 28, 2014

Work Packages



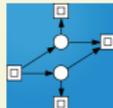
WP3



Efficient simulation of HPN^c

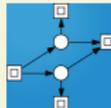
- a. Predecessor WPs: BTU-WP1, OvGUM-WP1
- b. Successor WPs: BTU-WP4
 - Examination of the Petri net models with regard to parallelisation potential
 - Investigation of optimisation possibilities and performance comparisons with alternative tools, i.e. StochKit2, Cain...

Parameter-free simulation



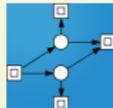
- Starting point: *The Signaling Petri Net-Based Simulator: A Non-Parametric Strategy for Characterizing the Dynamics of Cell-Specific Signaling Networks*
D. Ruths, M. Muller, Jen-Te Tseng, L. Nakhleh, P. T. Ram
Published: February 29, 2008; DOI: 10.1371/journal.pcbi.1000005
- „The key insight behind our approach is the assumption that, while all network parameters determine the actual signal propagation to some extent, the network connectivity is the most significant single determinant. While this is clearly a gross simplification, several researchers have observed that the connectivity of a biological network dictates, to a great extent, the network’s dynamics.”

Parameter-free simulation



- „Simulation of timed Petri nets with variable auto-concurrency”.
- The least possible time step is 1 time unit.
- All enabled transitions that are not mutually exclusive, are forced to fire within a time-step, something like *maximum step*.
- When the net is filled up with tokens, every transition will fire.

Parameter-free simulation



Transition firing

- Generate a random sequence of all transitions $t \in \mathcal{PN}$.
- No extra conflict resolution needed, because of serial firing.
- *Maximum step* \subseteq *random sequences*

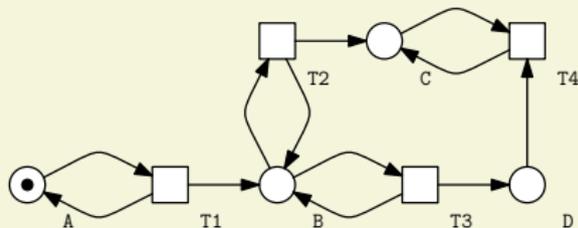
Example

Random sequences:

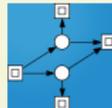
- 1 $(T_1, T_3, T_2, T_4) \rightarrow \{1, 1, 1, 0\}$
- 2 $(T_1, T_2, T_4, T_3) \rightarrow \{1, 1, 1, 1\}$
- 3 $(T_3, T_4, T_2, T_1) \rightarrow \{1, 1, 0, 0\}$

Maximum step:

- 1 $\{T_1\} \rightarrow \{1, 1, 0, 0\}$



Parameter-free simulation

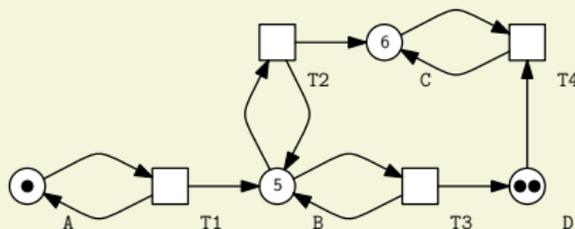


Transition firing

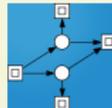
- A transition fires concurrently to itself, i.e. token flow increases.
- How often a transition concurrently fires depends on its enabledness degree and is randomly determined.
- $firing\ rate = random[0, enabledness\ degree]$
- This approximates the stochastic behaviour of mass-action kinetics.

Example

- $(T_1, T_2, T_3, T_4) \rightarrow \{1, 5, 10, 0\}$
- $T_1 = 1 \rightarrow 0 \rightarrow \{1, 5, 6, 2\}$
- $T_2 = 5 \rightarrow 4 \rightarrow \{1, 5, 10, 2\}$
- $T_3 = 5 \rightarrow 2 \rightarrow \{1, 5, 10, 4\}$
- $T_4 = 4 \rightarrow 4 \rightarrow \{1, 5, 10, 0\}$



Parameter-free simulation

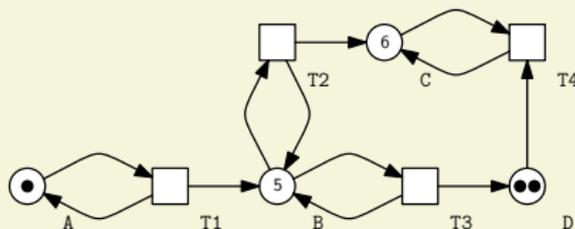


Transition firing

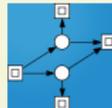
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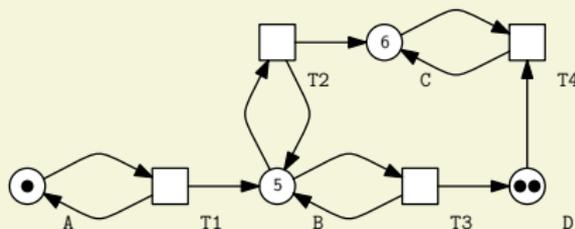


Transition firing

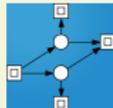
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Parameter-free simulation

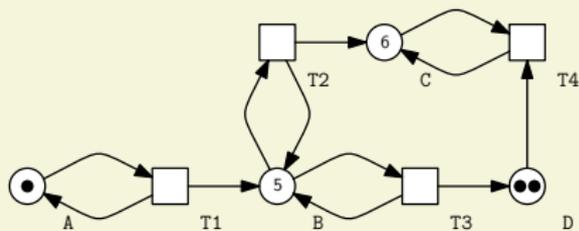


Transition firing

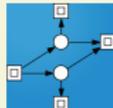
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Parameter-free simulation

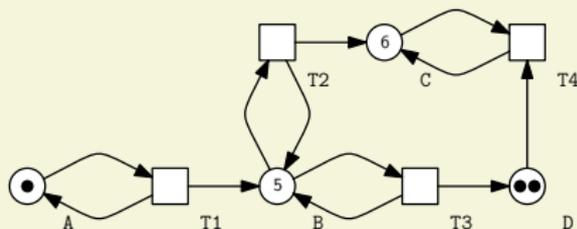


Transition firing

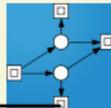
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Parameter-free simulation



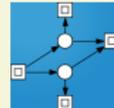
Algorithm 1 Parameter-free simulation algorithm

Require: \mathcal{PN} with initial marking m_0 , time interval $[\tau_0, \tau_{max}]$, runs r_{max}

Ensure: marking m at time point τ_{max}

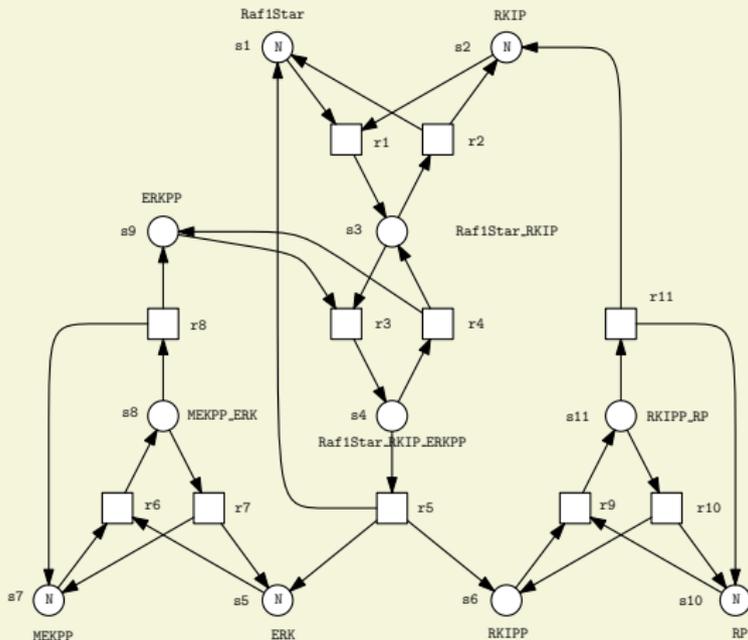
```
1: for  $r = 0$ ;  $r < r_{max}$ ;  $r \leftarrow r + 1$  do
2:    $initRand(seed)$ 
3:    $time \tau \leftarrow \tau_0$ ,  $marking m \leftarrow m_0$ ,  $T_r \leftarrow T$ 
4:   while  $\tau \leq \tau_{max}$  do
5:      $T_r \leftarrow random\_shuffle(T_r)$ 
6:     for all transitions  $t_j \in T_r$  do
7:        $e \leftarrow enablednessDegree(t_j, m)$ 
8:        $f \leftarrow random(0, e)$ 
9:        $m \leftarrow m + f * \Delta t_j$ 
10:    end for
11:     $generateResultPoint(\tau, m)$ 
12:     $\tau \leftarrow \tau + 1$ 
13:  end while
14: end for
```

Example

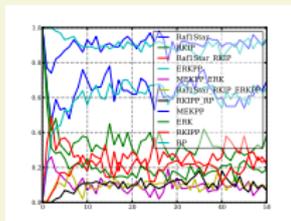
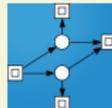


RKIP inhibited ERK Pathway [Gilbert et al. 2006]

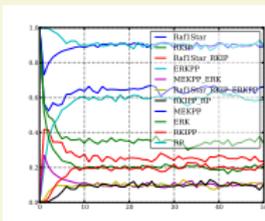
RKIP/MEK-ERK signalling pathway [wolkenhauer 2003], [Calder 2005]



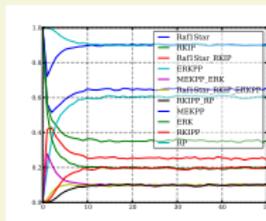
RKIP inhibited ERK Pathway



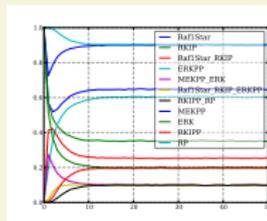
(a) SPN 100 runs (<1s)



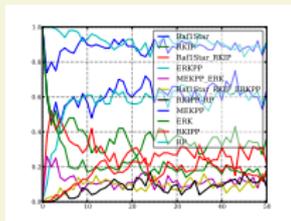
(b) SPN 1,000 runs (<1s)



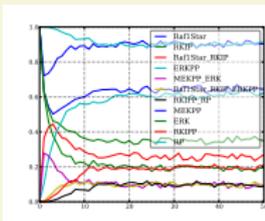
(c) SPN 10,000 runs (<1s)



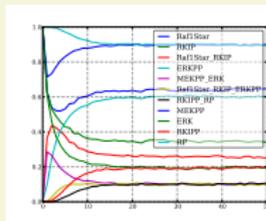
(d) SPN 100,000 runs (2s)



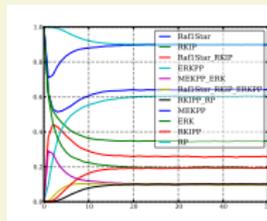
(e) QPN 100 runs (<1s)



(f) QPN 1,000 runs (<1s)



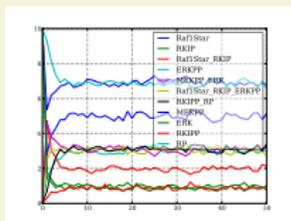
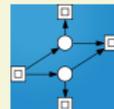
(g) QPN 10,000 runs (<1s)



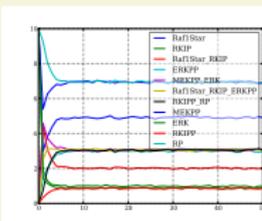
(h) QPN 100,000 runs (3s)

Figure: ERK, $N=1$

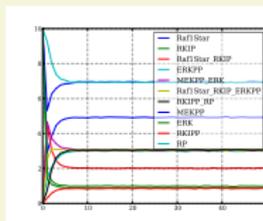
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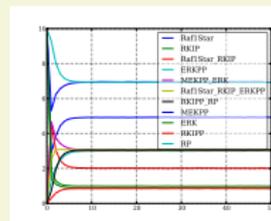
(a) SPN 100 runs (<1s)



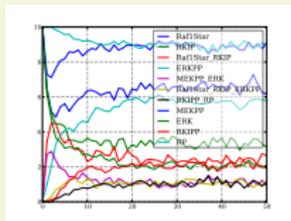
(b) SPN 1,000 runs (<1s)



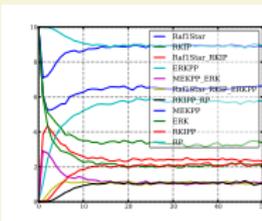
(c) SPN 10,000 runs (8s)



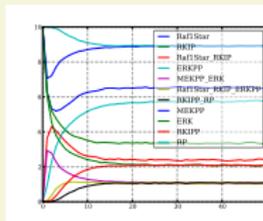
(d) SPN 100,000 runs (1m19s)



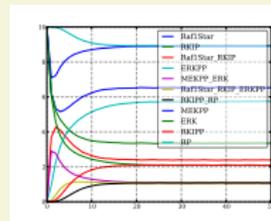
(e) QPN 100 runs (<1s)



(f) QPN 1,000 runs (<1s)



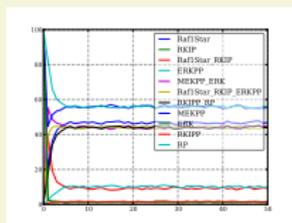
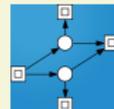
(g) QPN 10,000 runs (<1s)



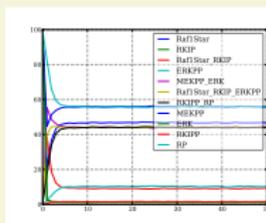
(h) QPN 100,000 runs (4s)

Figure: ERK, N=10

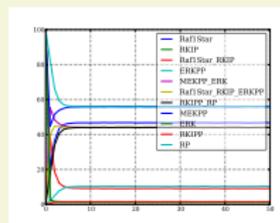
RKIP inhibited ERK Pathway



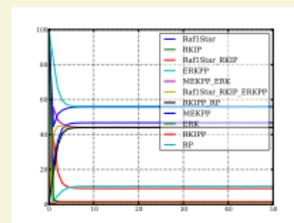
(a) SPN 100 runs (1s)



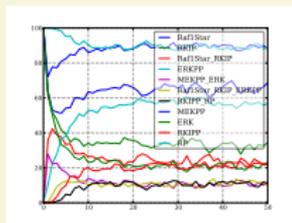
(b) SPN 1,000 runs (9s)



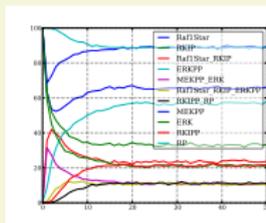
(c) SPN 10,000 runs (1m41s)



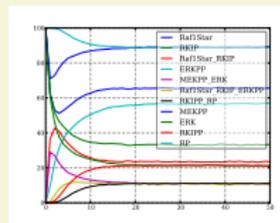
(d) SPN 100,000 runs (17m4s)



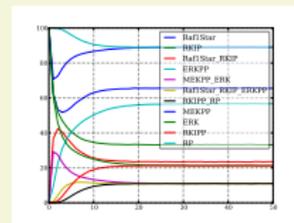
(e) QPN 100 runs (<1s)



(f) QPN 1,000 runs (<1s)



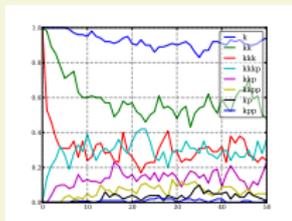
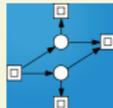
(g) QPN 10,000 runs (<1s)



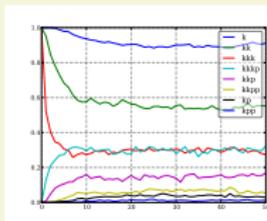
(h) QPN 100,000 runs (4s)

Figure: ERK, N=100

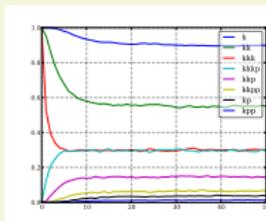
Mitogen-activated Protein Kinase



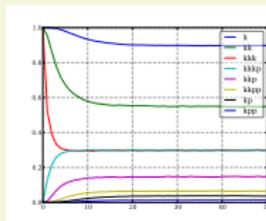
(a) SPN 100 runs (4s)



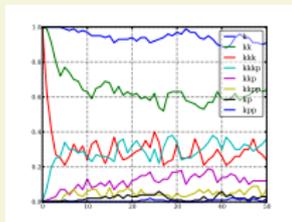
(b) SPN 1,000 runs (4s)



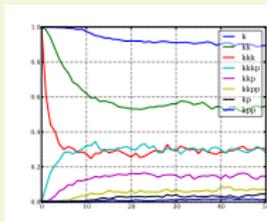
(c) SPN 10,000 runs (4s)



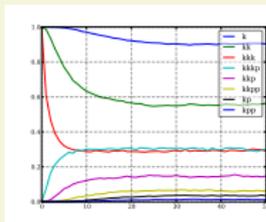
(d) SPN 100,000 runs (4s)



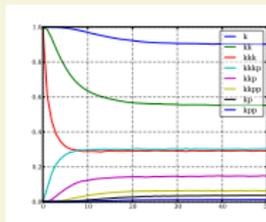
(e) QPN 100 runs (4s)



(f) QPN 1,000 runs (4s)



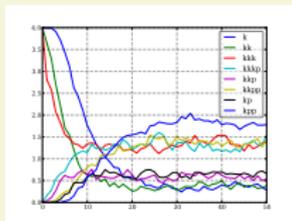
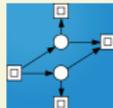
(g) QPN 10,000 runs (4s)



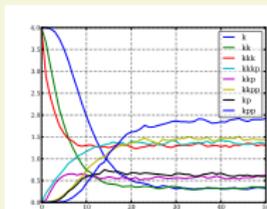
(h) QPN 100,000 runs (4s)

Figure: MAPK, $N=1$

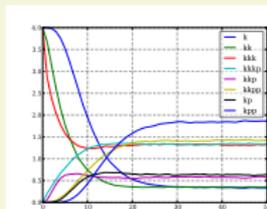
Mitogen-activated Protein Kinase



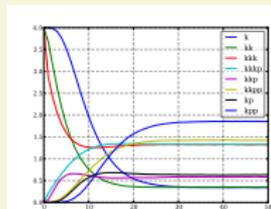
(a) SPN 100 runs (<1s)



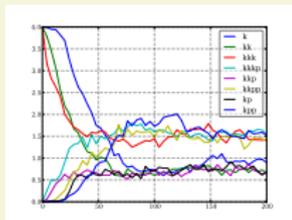
(b) SPN 1,000 runs (<1s)



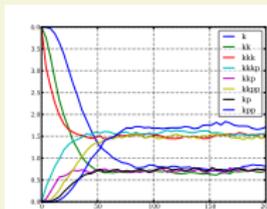
(c) SPN 10,000 runs (2s)



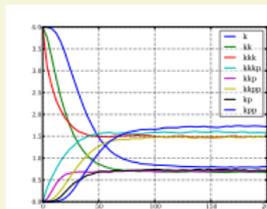
(d) SPN 100,000 runs (25s)



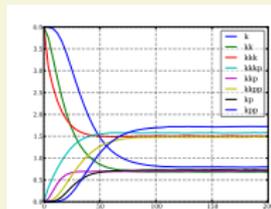
(e) QPN 100 runs (<1s)



(f) QPN 1,000 runs (<1s)



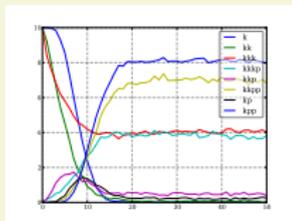
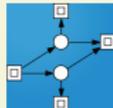
(g) QPN 10,000 runs (3s)



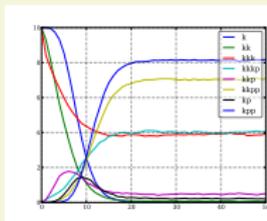
(h) QPN 100,000 runs (33s)

Figure: MAPK, $N=4$

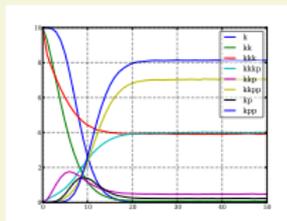
Mitogen-activated Protein Kinase



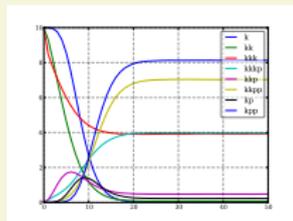
(a) SPN 100 runs (<1s)



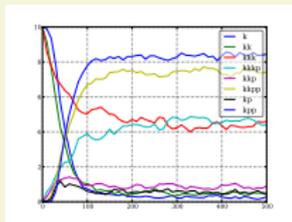
(b) SPN 1,000 runs (<1s)



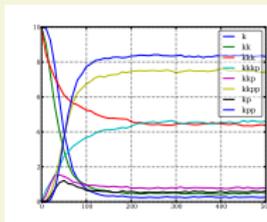
(c) SPN 10,000 runs (3s)



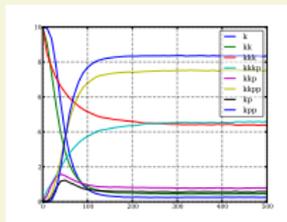
(d) SPN 100,000 runs (39s)



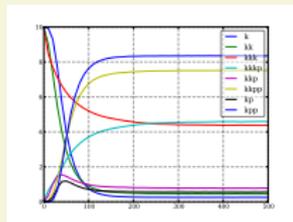
(e) QPN 100 runs (<1s)



(f) QPN 1,000 runs (<1s)



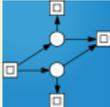
(g) QPN 10,000 runs (8s)



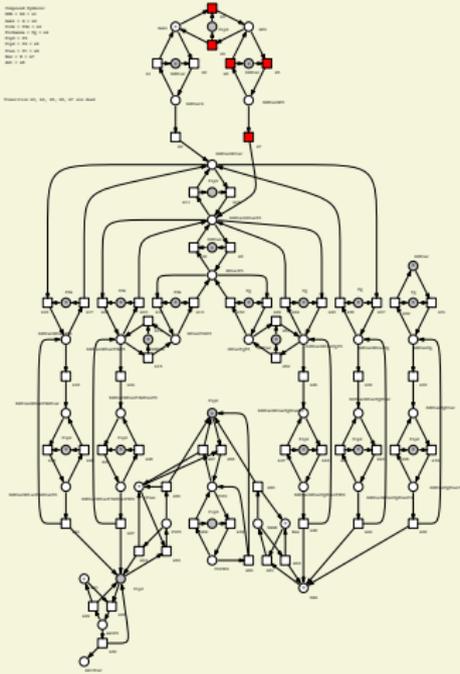
(h) QPN 100,000 runs (1m20s)

Figure: MAPK, N=10

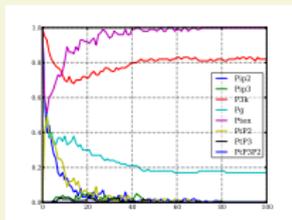
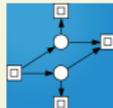
Example



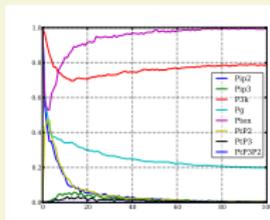
Angiogenesis [Napione et al. 2009]



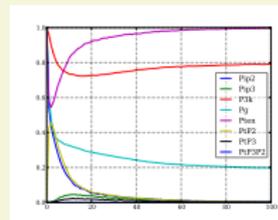
Angiogenesis



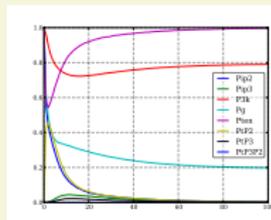
(a) SPN 100 runs (<1s)



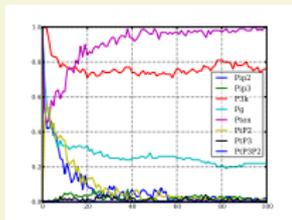
(b) SPN 1,000 runs (<1s)



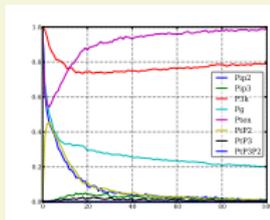
(c) SPN 10,000 runs (<1s)



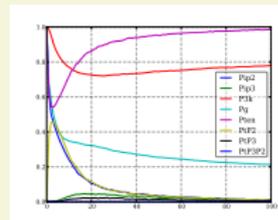
(d) SPN 100,000 runs (7s)



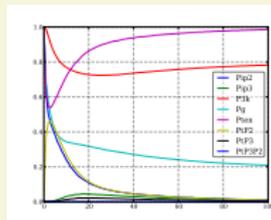
(e) QPN 100 runs (<1s)



(f) QPN 1,000 runs (<1s)



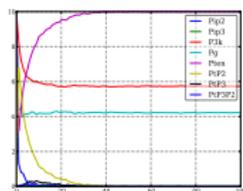
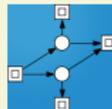
(g) QPN 10,000 runs (2s)



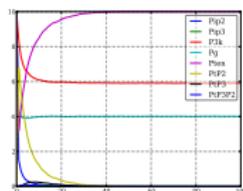
(h) QPN 100,000 runs (21s)

Figure: ANG, N=1

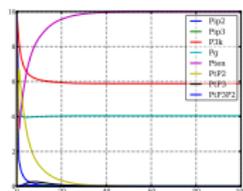
Angiogenesis



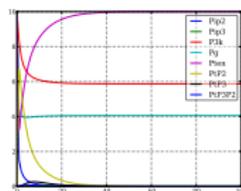
(a) SPN 100 runs (<1s)



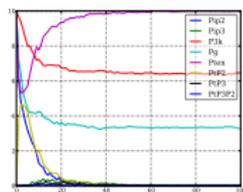
(b) SPN 1,000 runs (<1s)



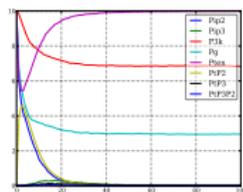
(c) SPN 10,000 runs (2s)



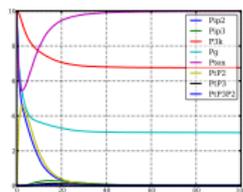
(d) SPN 100,000 runs (23s)



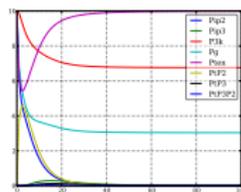
(e) QPN 100 runs (<1s)



(f) QPN 1,000 runs (<1s)



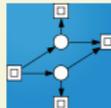
(g) QPN 10,000 runs (2s)



(h) QPN 100,000 runs (23s)

Figure: ANG, N=10

Parameter-free simulation



Conclusions

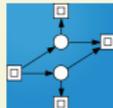
■ Mixed results:

- 1 performance comparable to stochastic simulation, some times better, some time worse
- 2 correct results for $N = 1$, contradictory for $N > 1$

■ Potential solutions:

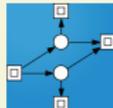
- 1 weighted shuffle of transitions
- 2 mass-action kinetics: $c_t \cdot \prod_{p \in \bullet t} \binom{m(p)}{f(p,t)}$
enabledness degree: $\min_{p \in \bullet t} \left(\left\lfloor \frac{m(p)}{f(p,t)} \right\rfloor \right)$

Milestones

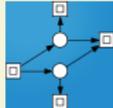


	2013				2014				2015			
	Q1	Q2	Q3	Q4	Q1	Q2	Q3	Q4	Q1	Q2	Q3	Q4
WP1	M1											
WP2		M2										
WP3			M3									
WP4					M4							
WP5						M4						
WP6							M5					
WP7											M6	

Next steps...



- Model compilation for simulation, i.e. a Petri net model and the simulation algorithm will be compiled into an executable file.
- Performance comparisons with alternative simulation tools, i.e. Stochkit2, Cain, Copasi, StochPy. . .



Thank you for your attention!