

This form is a summary description of the model entitled “MAPKbis” proposed for the Model Checking Contest @ Petri Nets. Models can be given in several instances parameterized by scaling parameters. Colored nets can be accompanied by one or many equivalent, unfolded P/T nets. Models are given together with property files (possibly, one per model instance) giving a set of properties to be checked on the model.

Description

This model is extracted from a set presented in [1] and modeling biological systems, obtained from influence graphs provided by biologists. This particular Petri net describes a Boolean model of Mitogen-Activated Protein Kinase network [2] (MAPK). It is a different model from the variant proposed in this contest in 2011.

References

1. Loïc Paulevé, “Reduction of Qualitative Models of Biological Networks for Transient Dynamics Analysis”, <https://hal.archives-ouvertes.fr/hal-01580765>
2. L. Grieco, L. Calzone, I. Bernard-Pierrot, F. Radvanyi, B. Kahn-Perls, and D. Thieffry, Integrative modelling of the influence of MAPK network on cancer cell fate decision, *PLoS Comput Biol*, vol. 9, no. 10, p. e1003286, oct 2013.

Scaling parameter

Parameter name	Parameter description	Chosen parameter values
$n1, n2, n3$	setting parameters	(53, 1, 0), (53, 2, 0)

Size of the model

Although the model is parameterized, its size does not depend on parameter values.

number of places: 106
 number of transitions: 173
 number of arcs: 986

Structural properties

ordinary — all arcs have multiplicity one ✓
simple free choice — all transitions sharing a common input place have no other input place ✗
extended free choice — all transitions sharing a common input place have the same input places ✗
state machine — every transition has exactly one input place and exactly one output place ✗
marked graph — every place has exactly one input transition and exactly one output transition ✗
connected — there is an undirected path between every two nodes (places or transitions) ?
strongly connected — there is a directed path between every two nodes (places or transitions) ?
source place(s) — one or more places have no input transitions ?
sink place(s) — one or more places have no output transitions ?
source transition(s) — one or more transitions have no input places ?
sink transitions(s) — one or more transitions have no output places ?
loop-free — no transition has an input place that is also an output place ✗
conservative — for each transition, the number of input arcs equals the number of output arcs ✓
subconservative — for each transition, the number of input arcs equals or exceeds the number of output arcs ✓
nested units — places are structured into hierarchically nested sequential units^(a) ?

^(a)the definition of Nested-Unit Petri Nets (NUPN) is available from <http://mcc.lip6.fr/nupn.php>

Behavioural properties

- safe** — *in every reachable marking, there is no more than one token on a place* ✓
- deadlock** — *there exists a reachable marking from which no transition can be fired* ?
- reversible** — *from every reachable marking, there is a transition path going back to the initial marking* ?
- quasi-live** — *for every transition t , there exists a reachable marking in which t can fire* ?
- live** — *for every transition t , from every reachable marking, one can reach a marking in which t can fire* ?