Adapting Biochemical Kripke Structures for **Distributed Model Checking**

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Bounded Hamming Distance Kripke Structures



Distributed model checking



Fragmenting Kripke structures



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Model checking on chemical reactions

- Reactants → products
- Non-deterministic

$$\begin{array}{cccc} A+B & \rightarrow & B+C \\ A+B+\neg C & \rightarrow & \neg A+B+C \\ A+B+\neg C & \rightarrow & A+B+C \\ A+B+C & \rightarrow & \neg A+B+C \\ A+B+C & \rightarrow & A+B+C \end{array}$$

• Few entities on each side (maximum is 6 in all public databases)

The resulting Kripke Structure

 $M = (S, I, R, \mathcal{L})$ is a k-Bounded Hamming Distance Kripke Structure (k-BHDKS) when: $\forall s, s' \in S, \quad R(s, s') \Rightarrow (H(\mathcal{L}(s), \mathcal{L}(s')) \leq k)$

Properties:

- Relatively sparse: each state has at most $|\mathcal{AP}|^k$ neighbours
- Can be effectively distributed into fragments

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Assumption based model checking

A Kripke Structure is decomposed into fragments

Each distributed node stores only one fragment, thus allowing to process larger model checking problems

A fragment is created by extending a subset of the state-space to its immediate neighbors

Kripke structure fragments



Obrázek: Dividing a KS in two fragments



Obrázek: KS with no possible fragmentation



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Distributed fragment

A distributed fragment M' of M = (S, R) around a set of core states $T \subseteq S$:

$$egin{aligned} M' &= (S_T, R_T) \ S_T &= \{ s \in S | s \in T \lor \exists s' \in T \text{ such that } (s', s) \in R \} \ R_T &= \{ (s_1, s_2) \in R | s_1 \in T, s_2 \in S_T \} \end{aligned}$$

Separator V of a set T is the minimal set of states, satisfying that there is no path from T to $S \setminus T$ avoiding all states in V

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Existence of fragmentation in k-BHDKS

- Let |T| be the size of the core set
- Each state in a *k*-BHDKS has at most $|\mathcal{AP}|^k$ neighbors
- The size of the fragment is at most $|T| + |T| \cdot |AP|^k$ grows polynomially with the number of propositions

A hypercube

- A *k*-BHDKS with |AP| propositions can be partitioned to an *l*-dimensional hypercube as long as l < |AP|/k
- Construction of the partitioning:
 - Create 2^{*l*} symmetrically placed centers a_0 to a_{2^l-1} as states $0^{l,p}$, $0^{(l-1).p}1^p, \ldots, 1^{l,p}$ where p is $\left\lceil \frac{|\mathcal{AP}|}{l} \right\rceil$
 - Add a state *s* to fragment *n* if *a_n* is the nearest center from *s* with respect to Hamming distance
 - Include each fragment *n* include all immediate successors (the border)



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Properties of the partitioning

- The size of the core is approx. $\frac{1}{2^{l}}$. |KS|
- The size of the border associated with the core is below $\frac{1}{2^{l}}$. |KS|
- Transition can exist only between states in adjacent nodes of the hypercube

The End

Thank you for your attention Discussion



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