

A Linear Parallel Algorithm to Compute Bisimulation and Relational Coarsest Partitions

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Abstract. The most efficient way to calculate strong bisimilarity is by finding the relational coarsest partition of a transition system. We provide the first linear-time algorithm to calculate strong bisimulation using parallel random access machines (PRAMs). More precisely, with n states, m transitions and $|Act| \leq m$ action labels, we provide an algorithm for $\max(n,m)$ processors that calculates strong bisimulation in time $\mathcal{O}(n+|Act|)$ and space $\mathcal{O}(n+m)$. The best-known PRAM algorithm has time complexity $\mathcal{O}(n\log n)$ on a smaller number of processors making it less suitable for massive parallel devices such as GPUs. An implementation on a GPU shows that the linear time-bound is achievable on contemporary hardware.

1 Introduction

The notion of bisimilarity for Kripke structures and Labelled Transition Systems (LTSs) is commonly used to define behavioural equivalence. Deciding this behavioural equivalence is important in the field of modelling and verifying concurrent and multi-component systems [4,15]. Kanellakis and Smolka proposed a partition refinement algorithm for obtaining the bisimilarity relation for Kripke structures [11]. The proposed algorithm has a run time complexity of $\mathcal{O}(nm)$ where n is the number of states and m is the number of transitions of the input. Later, a more sophisticated refinement algorithm running in $\mathcal{O}(m \log n)$ steps was proposed by Paige and Tarjan [16].

In recent years the increase in the speed of sequential chip designs has stagnated due to a multitude of factors such as energy consumption and heat generation. In contrast, parallel devices such as graphics processing units (GPUs) keep increasing rapidly in computational power. In order to profit from the acceleration of these devices, we require algorithms with massive parallelism. The article "There's plenty of room at the Top: What will drive computer performance after

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Moore's law" by Leierson et al. [13] indicates that the advance in computational performance will come from software and algorithms that can employ hardware structures with a massive number of simple, parallel processors, such as GPUs. In this paper, we propose such an algorithm to decide bisimilarity.

Deciding bisimilarity is P-complete [1], which suggests that bisimilarity is an inherently sequential problem. This fact has not withheld the community from searching for efficient parallel algorithms for deciding bisimilarity of Kripke structures. In particular, Lee and Rajasekaran [12,17] proposed a parallel algorithm based on the Paige Tarjan algorithm that works in $\mathcal{O}(n \log n)$ time complexity using $\frac{m}{\log n}\log\log n$ Concurrent, Read Concurrent Write (CRCW) processors, and one using only $\frac{m}{n}log n$ Concurrent Read Exclusive Write (CREW) processors. Jeong et al. [10] presented a linear time parallel algorithm, but it is probabilistic in the sense that it has a non-zero chance to output the wrong result. Furthermore, Wijs [22] presented a GPU implementation of an algorithm to solve the strong and branching bisimulation partition refinement problems. In a distributed setting, Blom and Orzan studied algorithms for refinement [2]. Those algorithms use message passing as a way of communication between different workers in a network and rely on a small number of processors. Therefore, they are very different in nature than our algorithm. Those algorithms were extended and optimized for branching bisimulation [3].

In this work, we improve on the best known theoretical bound for PRAM algorithms using a higher degree of parallelism. The proposed algorithm improves the run time complexity to $\mathcal{O}(n)$ on $\max(n,m)$ processors and is based on the sequential algorithm of Kanellakis and Smolka [11]. The larger number of processors used in this algorithm favours the increasingly parallel design of contemporary and future hardware. In addition, the algorithm is *optimal* w.r.t. the sequential Kanellakis-Smolka algorithm, meaning that overall, it does not perform more work than its sequential counterpart.

We first present our algorithm on Kripke structures where transitions are unlabelled. However, as labelled transition systems (LTSs) are commonly used, and labels are not straightforward to incorporate in an efficient way (cf. for instance [21]), we discuss how our algorithm can be extended to take action labels into account. This leads to an algorithm with a run time complexity of $\mathcal{O}(n + |Act|)$, with Act the set of action labels.

Our algorithm has been designed for and can be analyzed with the CRCW PRAM model, following notations from [20]. This model is an extension of the normal RAM model, allowing multiple processors to work with shared memory. In the CRCW PRAM model, parallel algorithms can be described in a straightforward and elegant way. In reality, no device exists that completely adheres to this PRAM model, but with recent advancements, hardware gets better and better at approximating the model since the number of parallel threads keeps growing. We demonstrate this by translating the PRAM algorithm to GPU code. We straightforwardly implemented our algorithm in CUDA and experimented with an NVIDIA Titan RTX, showing that our algorithm performs mostly in line with what our PRAM algorithm predicts.

The paper is structured as follows: In Sect. 2, we recall the necessary preliminaries on the CRCW PRAM model and state the partition refinement problems this paper focuses on. In Sect. 3, we propose a parallel algorithm to compute bisimulation for Kripke structures, which is also called the Relational Coarsest Partition Problem (RCPP). In this section, we also prove the correctness of the algorithm and provide a complexity analysis. In Sect. 4, we discuss the details for an implementation with multiple action labels, thereby supporting LTSs, which forms the Bisimulation Coarsest Refinement Problem (BCRP). In Sect. 5 we discuss the results of the implementation and in Sect. 6 we draw conclusions.

2 Preliminaries

2.1 The PRAM Model

The Parallel Random Access Machine (PRAM) is a natural extension of the normal Random Access Machine (RAM), where an arbitrary number of parallel processors can access the memory. Following the definitions of [20] we use a version of PRAM that is able to Concurrently Read and Concurrently Write (CRCW PRAM). It differs from the model introduced in [6] in which the PRAM model was only allowed to concurrently read from the same memory address, but concurrent writes (to the same address) could not happen.

A CRCW PRAM consists of a sequence of numbered processors P_0, P_1, \ldots These processors have all the natural instructions of a normal RAM such as addition, subtraction, and conditional branching based on the equality and less-than operators. There is an infinite amount of common memory the processors have access to. The processors have instructions to read from and write to the common memory. In addition, a processor P_i has an instruction to obtain its unique index i. A PRAM also has a function $\mathcal{P}: \mathbb{N} \to \mathbb{N}$ which defines a bound on the number of processors given the size of the input.

All the processors have the same program and run synchronized in a single instruction, multiple data (SIMD) fashion. In other words, all processors execute the program in lock-step. Parallelism is achieved by distributing the data elements over the processors and having the processors apply the program instructions on 'their' data elements.

Initially, given input consisting of n data elements, the CRCW PRAM assumes that the input is stored in the first n registers of the common memory, and starts the first $\mathcal{P}(n)$ processors $P_0, P_1, \ldots, P_{\mathcal{P}(n)-1}$.

Whenever a concurrent write happens to the same memory cell, we assume that one arbitrary write will succeed. This is called the *arbitrary CRCW PRAM*.

A parallel program for a PRAM is called *optimal* w.r.t. a sequential algorithm if the total work done by the program does not exceed the work done by the sequential algorithm. More precisely, if T is the parallel run time and P the number of processors used, then the algorithm is optimal w.r.t. a sequential algorithm running in S steps if $P \cdot T \in \mathcal{O}(S)$.

2.2 Strong Bisimulation

To formalise concurrent system behaviour, we use LTSs.

Definition 1 (Labeled Transition System). A Labeled Transition System (LTS) is a three-tuple $A = (S, Act, \rightarrow)$ where S is a finite set of states, Act a finite set of action labels, and $\rightarrow \subseteq S \times Act \times S$ the transition relation.

Let $A = (S, Act, \rightarrow)$ be an LTS. Then, for any two states $s, t \in S$ and $a \in Act$, we write $s \xrightarrow{a} t$ iff $(s, a, t) \in \rightarrow$.

Kripke structures differ from LTSs in the fact that the states are labelled as opposed to the transitions. In the current paper, for convenience, instead of using Kripke structures where appropriate, we reason about LTSs with a single action label, i.e., |Act| = 1. Computing the coarsest partition of such an LTS can be done in the same way as for Kripke structures, apart from the fact that in the latter case, a different initial partition is computed that is based on the state labels (see, for instance, [8,9]).

Definition 2 (Strong bisimulation). On an LTS $A = (S, Act, \rightarrow)$ a relation $R \subseteq S \times S$ is called a strong bisimulation relation if and only if it is symmetric and for all $s, t \in S$ with sRt and for all $a \in Act$ with $s \stackrel{a}{\longrightarrow} s'$, we have:

$$\exists t' \in S.t \xrightarrow{a} t' \land s'Rt'$$

Whenever we refer to bisimulation we mean strong bisimulation. Two states $s, t \in S$ in an LTS A are called *bisimilar*, denoted by $s \rightleftharpoons t$, iff there is some bisimulation relation R for A that relates s and t.

A partition π of a finite set of states S is a set of subsets that are pairwise disjoint and whose union is equal to S, i.e., $\bigcup_{B \in \pi} B = S$. Every element $B \in \pi$ of this partition π is called a *block*.

We call partition π' a refinement of π iff for every block $B' \in \pi'$ there is a block $B \in \pi$ such that $B' \subseteq B$. We say a partition π of a finite set S induces the relation $R_{\pi} = \{(s,t) \mid \exists B \in \pi.s \in B \land t \in B\}$. This is an equivalence relation of which the blocks of π are the equivalence classes.

Given an LTS $A = (S, Act, \rightarrow)$ and two states $s, t \in S$ we say that s reaches t with action $a \in Act$ iff $s \xrightarrow{a} t$. A state s reaches a set $U \subseteq S$ with an action a iff there is a state $t \in U$ such that s reaches t with action a. A set of states $V \subseteq S$ is called stable under a set of states $U \subseteq S$ iff for all actions a either all states in V reach U with a, or no state in V reaches U with a. A partition π is stable under a set of states U iff each block $B \in \pi$ is stable under U. The partition π is called stable iff it is stable under all its own blocks $B \in \pi$.

Fact 1. [16] Stability is inherited under refinement, i.e. given a partition π of S and a refinement π' of π , then if π is stable under $U \subseteq S$, then π' is also stable under U.

The main problem we focus on in this work is called the bisimulation refinement problem (**BCRP**). It is defined as follows:

Input: An LTS $M = (S, Act, \rightarrow)$.

Output: The partition π of S which is the coarsest partition, i.e., has the smallest number of blocks, that forms a bisimulation relation.

In a Kripke structure, the transition relation forms a single binary relation, since the transitions are unlabelled. This is also the case when an LTS has a single action label. In that case, the problem is called the Relational Coarsest Partition Problem (**RCPP**) [11,12,16]. This problem is defined as follows:

Input: A set S, a binary relation \rightarrow : $S \times S$ and an initial partition π_0

Output: The partition π which is the coarsest refinement of π_0 and which is a bisimulation relation.

It is known that BCRP is not significantly harder than RCPP as there are intuitive translations from LTSs to Kripke structures [5, Dfn. 4.1]. However, some non-trivial modifications can speed-up the algorithm for some cases, hence we discuss both problems separately. In Sect. 3, we discuss the basic parallel algorithm for RCPP, and in Sect. 4, we discuss the modifications required to efficiently solve the BCRP problem for LTSs with multiple action labels.

3 Relational Coarsest Partition Problem

In this section, we discuss a sequential algorithm based on the one of Kanellakis and Smolka [11] for RCPP (Sect. 3.1). This is the basic algorithm that we adapt to the parallel PRAM algorithm (Sect. 3.2). The algorithm starts with an input partition π_0 and refines all blocks until a stable partition is reached. This stable partition will be the coarsest refinement that defines a bisimulation relation.

3.1 The Sequential Algorithm

The sequential algorithm, Algorithm 1, works as follows. Given are a set S, a transition relation $\rightarrow \subseteq S \times S$, and an initial partition π_0 of S. Initially, we mark the partition as not necessarily stable under all blocks by putting these blocks in a set Unstable. In any iteration of the algorithm, if a block B of the current partition is not in Unstable, then the current partition is stable under B. If Unstable is empty, the partition is stable under all its blocks, and the partition represents the required bisimulation.

As long as some blocks are in *Unstable* (line 3), a single block $B \in \pi$ is taken from this set (line 4) and we split the current partition such that it becomes stable under B. Therefore, we refer to this block as the *splitter*. The set $S' = \{s \in S \mid \exists t \in B.s \to t\}$ is the reverse image of B (line 6). This set consists of all states that can reach B, and we use S' to define our new blocks. All blocks B' that have a non-empty intersection with S', i.e., $B' \cap S' \neq \emptyset$, and are not a subset of S', i.e., $B' \cap S' \neq B'$ (line 7), are split in the subset of states in S' and the subset of states that are not in S' (lines 8–9). These two new blocks are added to the set of *Unstable* blocks (line 10). The number of states is finite, and blocks can be split only a finite number of times. Hence, blocks are only finitely often put in *Unstable*, and so the algorithm terminates.

Algorithm 1: Sequential algorithm based on Kanellakis-Smolka

```
1 \pi := \pi_0;
 2 Unstable := \pi;
 3 while Unstable \neq \emptyset do
          foreach B \in Unstable do
 4
                Unstable := Unstable \setminus \{B\};
 5
               S' := \{ s \in S \mid \exists t \in B.s \to t \};
 6
               foreach B' \in \pi with \emptyset \subset B' \cap S' \subset B' do
 7
                    // Split B' into B' \cap S' and B' \setminus S'
                    \pi := \pi \setminus \{B\};
 8
                    \pi := \pi \cup \{B' \cap S', B' \setminus S'\};
 9
                     Unstable := Unstable \cup \{B' \cap S', B' \setminus S'\};
10
11
               end
          end
12
13 end
```

3.2 The PRAM Algorithm

Next, we describe a PRAM algorithm to solve RCPP that is based on the sequential algorithm given in Algorithm 1.

Block Representation. Given an LTS $A = (S, Act, \rightarrow)$ with |A| = 1 and |S| = n states, we assume that the states are labeled with unique indices $0, \ldots, n-1$. A partition π in the PRAM algorithm is represented by assigning a block label from a set of block labels L_B to every state. The number of blocks can never be larger than the number of states, hence, we use the indices of the states as block labels: $L_B = S$. We exploit this in the PRAM algorithm to efficiently select a new block label whenever a new block is created. We select the block label of a new block by electing one of its states to be the leader of that block and using the index of that state as the block label. By doing so, we maintain an invariant that the leader of a block is also a member of the block.

In a partition π , whenever a block $B \in \pi$ is split into two blocks B' and B'', the leader s of B which is part of B' becomes the leader of B', and for B'', a new state $t \in B''$ is elected to be the leader of this new block. Since the new leader is not part of any other block, the label of t is fresh with respect to the block labels that are used for the other blocks. This method of using state leaders to represent subsets was first proposed in [22, 23].

Data Structures. The common memory contains the following information:

- 1. $n: \mathbb{N}$, the number of states of the input.
- 2. $m:\mathbb{N}$, the number of transitions of the input relation.
- 3. The input, a fixed numbered list of transitions. For every index $0 \le i < m$ of a transition, a source $source_i \in S$ and target $target_i \in S$ are given, representing the transition $source_i \to target_i$.

- 4. $C: L_B \cup \{\bot\}$, the label of the current block that is used as a splitter; \bot indicates that no splitter has been selected.
- 5. The following is stored in lists of size n, for each state with index i:
 - (a) $mark_i : \mathbb{B}$, a mark indicating whether state i is able to reach the splitter.
 - (b) $block_i : L_B$, the block of which state i is a member.
- 6. The following is stored in lists of size n, for each potential block with block label i:
 - (a) $new_leader_i : L_B$ the leader of the new block when a split is performed.
 - (b) $unstable_i : \mathbb{B}$ indicating whether π is possibly unstable w.r.t. the block.

As input, we assume that each state with index i has an input variable $I_i \in L_B$ that is the initial block label. In other words, the values of the I_i variables together encode π_0 . Using this input, the initial values of the block label $block_i$ variables are calculated to conform to our block representation with leaders. Furthermore in the initialization, $unstable_i = false$ for all i that are not used as block label, and true otherwise.

The Algorithm. We provide our first PRAM algorithm in Algorithm 2. The PRAM is started with $\max(n, m)$ processors. These processors are dually used for transitions and states.

The algorithm performs initialisation (lines 1–6), after which each block has selected a new leader (lines 3–4), ensuring that the leader is one of its own states, and the initial blocks are set to unstable. Subsequently, the algorithm enters a single loop that can be explained in three separate parts.

- Splitter selection (lines 8–14), executed by n processors. Every variable $mark_i$ is set to false. After this, every processor with index i will check $unstable_i$. If block i is marked unstable the processor tries to write i in the variable C. If multiple write accesses to C happen concurrently in this iteration, then according to the arbitrary PRAM model (see Sect. 2), only one process j will succeed in writing, setting C := j as splitter in this iteration.
- Mark states (lines 15–17), executed by m processors. Every processor i is responsible for the transition $s_i \to t_i$ and checks if t_i (target_i) is in the current block C (line 15). If this is the case the processor writes true to $mark_{source_i}$ where $source_i$ is s_i . This mark now indicates that s_i reaches block C.
- Performing splits (lines 18–26), executed by n processors. Every processor i compares the mark of state i, i.e., $mark_i$, with the mark of the leader of the block in which state i resides, i.e., $mark_{block_i}$ (line 20). If the marking is different, state i has to be split from $block_i$ into a new block. At line 21, a new leader is elected among the states that form the newly created block. The index of this leader is stored in $new_leader_{block_i}$. The instability of block $block_i$ is set to true (line 22). After that, all involved processors update the block index for their state (line 23) and update the stability of the new block (line 24).

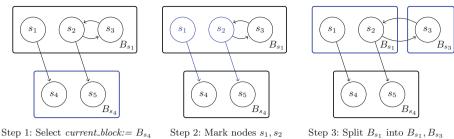


Fig. 1. One iteration of Algorithm 2

The steps of the program are illustrated in Fig. 1. The notation B_{s_i} refers to a block containing all states that have state s_i as their block leader. In the figure on the left, we have two blocks B_{s_1} and B_{s_4} , of which at least B_{s_4} is marked unstable. Block B_{s_4} is selected to be splitter, i.e., $C = B_{s_4}$ at line 12 of Algorithm 2. In the figure in the middle, $mark_i$ is set to true for each state i that can reach B_{s_4} (line 16). Finally, block B_{s_4} is set to stable (line 19), all states compare their mark with the leader's mark, and the processor working on state s_3 discovers that the mark of s_3 is different from the mark of s_1 , so s_3 is elected as leader of the new block B_{s_3} at line 21 of Algorithm 2. Both B_{s_1} and B_{s_3} are set to unstable (lines 22 and 24).

The algorithm repeats execution of the **while**-loop until all blocks are marked stable.

3.3 Correctness

The $block_i$ list in the common memory at the start of iteration k defines a partition π_k where states $s \in S$ with equal block labels $block_i$ form the blocks:

$$\pi_k = \{ \{ s \in S \mid block_s = s' \} \mid s' \in S \} \setminus \{\emptyset\}$$

A run of the program produces a sequence π_0, π_1, \ldots of partitions. Partition π_k is a refinement of every partition $\pi_0, \pi_1, \ldots, \pi_{k-1}$, since blocks are only split and never merged.

A partition π induces a relation of which the blocks are the equivalence classes. For an input partition π_0 we call the relation induced by the coarsest refinement of π_0 that is a bisimulation relation $\rightleftharpoons_{\pi_0}$.

We now prove that Algorithm 2 indeed solves RCPP. We first introduce Lemma 1 which is invariant throughout the execution and expresses that states which are related by \cong_{π_0} are never split into different blocks. This lemma implies that if a refinement forms a bisimulation relation, it is the coarsest.

Lemma 1. Let S be the input set of states, \rightarrow : $S \times S$ the input relation and π_0 the input partition. Let π_1, π_2, \ldots be the sequence of partitions produced by

Algorithm 2: The algorithm for RCPP for each processor P_i in the PRAM

```
1 if i < n then
        unstable_i := false;
 2
 3
        new\_leader_{I_i} := i;
 4
        block_i := new\_leader_{I_i};
 5
        unstable_{block_i} := true;
 6 end
 7 do
        C := \bot;
 8
        if i < n then
 9
             mark_i := false;
10
            if unstable_i then
11
12
                C := i;
13
            end
        end
14
        if i < m and block_{target_i} = C then
15
            mark_{source_i} := true;
16
17
        end
        if i < n and C \neq \bot then
18
             unstable_C := false;
19
             if mark_i \neq mark_{block_i} then
20
                 new\_leader_{block_i} := i;
21
                 unstable_{block_i} := true;
                 block_i := new\_leader_{block_i};
23
                 unstable_{block_i} := true;
24
            end
25
        end
26
27 while C \neq \bot;
```

Algorithm 2, then for all initial blocks $B \in \pi_0$, states $s, t \in B$ and iterations $k \in \mathbb{N}$:

$$s \bowtie_{\pi_0} t \implies \exists B \in \pi_k.s, t \in B$$

Proof. This is proven by induction on k. In the base case, π_0 , this is true by default. Now assume for a particular $k \in \mathbb{N}$ that the property holds. We know that the partition π_{k+1} is obtained by splitting with respect to a block $C \in \pi_k$. For two states $s, t \in S$ with $s \cong_{\pi_0} t$ we know that s and t are in the same block in π_k . In the case that both s and t do not reach C, then $mark_s = mark_t = false$. Since they were in the same block, they will be in the same block in π_{k+1} .

Now consider the case that at least one of the states is able to reach C. Without loss of generality say that s is able to reach C. Then there is a transition $s \to s'$ with $s' \in C$. By Definition 2, there exists a $t' \in S$ such that $t \to t'$ and $s' \rightleftharpoons_{\pi_0} t'$. By the induction hypothesis we know that since $s' \rightleftharpoons_{\pi_0} t'$, s' and t' must be in the same block in π_k , i.e., t' is in C. This witnesses that t is also able to reach C and we must have $mark_s = mark_t = true$. Since the states s and t

are both marked and are in the same block in π_k , they will remain in the same block in π_{k+1} .

Lemma 2. Let S be the input set of states with \rightarrow : $S \times S$, $L_B = S$ the block labels, and π_n the partition stored in the memory after the termination of Algorithm 2. Then the relation induced by π_n is a bisimulation relation.

Proof. Since the program finished, we know that for all block indices $i \in L_B$ we have $unstable_i$ = false. For a block index $i \in L_B$, $unstable_i$ is set to false if the partition π_k , after iteration k, is stable under the block with index i and set to true if it is split. So, by Fact 1, we know π_n is stable under every block B, hence stable. Next, we prove that a stable partition is a bisimulation relation.

We show that the relation R induced by π_n is a bisimulation relation. Assume states $s, t \in S$ with sRt are in block $B \in \pi_n$. Consider a transition $s \to s'$ with $s' \in S$. State s' is in some block $B' \in \pi_n$, and since the partition is stable under block B', and s is able to reach B', by the definition of stability, we know that t is also able to reach B'. Therefore, there must be a state $t' \in B'$ such that $t \to t'$ and s'Rt'. Finally, by the fact that R is an equivalence relation we know that R is also symmetric, therefore it is a bisimulation relation.

Theorem 1. The partition resulting from executing Algorithm 2 forms the coarsest relational partition for a set of states S and a transition relation $\rightarrow: S \times S$, solving RCPP.

Proof. By Lemma 2, the resulting partition is a bisimulation relation. Lemma 1 implies that it is the coarsest refinement which is a bisimulation.

3.4 Complexity Analysis

Every step in the body of the **while**-loop can be executed in constant time. So the asymptotic complexity of this algorithm is given by the number of iterations.

Theorem 2. RCPP on an input with m transitions and n states is solved by Algorithm 2 in $\mathcal{O}(n)$ time using $\max(n,m)$ CRCW PRAM processors.

Proof. In iteration $k \in \mathbb{N}$ of the algorithm, let us call the total number of blocks $N_k \in \mathbb{N}$ and the number of unstable blocks $U_k \in \mathbb{N}$. Initially, $N_0 = U_0 = |\pi_0|$. In every iteration k, a number of blocks $l_k \in \mathbb{N}$ is split, resulting in l_k new blocks, so the new total number of blocks at the end of iteration k is $N_{k+1} = N_k + l_k$.

First the current block C in iteration k which was unstable is set to stable which causes the number of unstable blocks to decrease by one. In this iteration k the l_k blocks B_1, \ldots, B_{l_k} are split, resulting in l_k newly created blocks. These l_k blocks are all unstable. A number of blocks $l'_k \leq l_k$ of the blocks $B_1, \ldots B_{l_k}$, were stable and are set to unstable again. The block C which was set to stable is possibly one of these l'_k blocks which were stable and set to unstable again. The total number of unstable blocks at the end of iteration k is $U_{k+1} = U_k + l_k + l'_k - 1$.

For all $k \in \mathbb{N}$, in iteration k we calculate the total number of blocks $N_k = |\pi_0| + \sum_{i=0}^{k-1} (l_i)$ and unstable blocks $U_k = |\pi_0| - k + \sum_{i=0}^{k-1} (l_i + l_i')$. The number

of iterations is given by $k = \sum_{i=0}^{k-1} (l_i + l_i') - U_k + |\pi_0|$. By definition, $l_i' \leq l_i$, and the total number of newly created blocks is $\sum_{i=0}^{k-1} (l_i) = N_k - |\pi_0|$, hence $\sum_{i=0}^{k-1} (l_i + l_i') \leq 2 \sum_{i=0}^{k-1} (l_i) \leq 2N_k - 2|\pi_0|$. The number of unstable blocks is always positive, i.e., $U_k \geq 0$, and the total number of blocks can never be larger than the number of states, i.e., $N_k \leq n$, so the total number of iterations z is bounded by $z \leq 2N_z - |\pi_0| \leq 2n - |\pi_0|$.

4 Bisimulation Coarsest Refinement Problem

In this section we extend our algorithm to the Bisimulation Coarsest Refinement Problem (BCRP), i.e., to LTSs with multiple action labels.

Solving BCRP can in principle be done by translating an LTS to a Kripke structure, for instance by using the method described in [18]. This translation introduces a new state for every transition, resulting in a Kripke structure with n+m states. If the number of transitions is significantly larger than the number of states, then the number of iterations of our algorithm increases undesirably.

4.1 The PRAM Algorithm

Instead of introducing more states, we introduce multiple marks per state, but in total we have no more than m marks. For each state s, we use a mark variable for each different outgoing action label relevant for s, i.e., for each a for which there is a transition $s \stackrel{a}{\to} t$ to some state t. Each state may have a different set of outgoing action labels and thus a different set of marks. Therefore, we first perform a preprocessing procedure in which we group together states that have the same set of outgoing action labels. This is valid, since two bisimilar states must have the same outgoing actions. That two states of the same block have the same set of action labels is then an invariant of the algorithm, since in the sequence of produced partitions, each partition is a refinement of the previous one. For the extended algorithm, we need to maintain extra information in addition to the information needed for Algorithm 2. For an input LTS $A = (S, Act, \rightarrow)$ with n states and m transitions the extra information is:

- 1. Each action label has an index $a \in \{0, \dots, |Act| 1\}$.
- 2. The following is stored in lists of size m, for each transition $s \xrightarrow{a} t$ with transition index $i \in \{0, ..., m-1\}$:
 - (a) $a_i := a$
 - (b) $order_i : \mathbb{N}$, the order of this action label, with respect to the source state s. E.g., if a state s has the list [1,3,6] of outgoing action labels, and transition i has label 3, then $order_i$ is 1 (we start counting from 0).
- 3. mark: [B], a list of up to m marks, in which there is a mark for every state, action pair for which it holds that the state has at least one outgoing transition labelled with that action. This list can be interpreted as the concatenation of sublists, where each sublist contains all the marks for one state. For each state $s \in S$ we have:

- (a) $off(s): \mathbb{N}$, the offset to access the beginning of the sublist of the marks of the state s in mark. The positions $mark_{off(s)}$ up to $mark_{off(s+1)}$ contain the sublist of marks for state s. E.g., if state s has outgoing transitions with 3 distinct action labels, we know that $off(s+1) \equiv off(s)+3$, and we have 3 marks for state s. We write $mark_{off(s)+order_i}$ to access the mark for transition i which has source state s.
- 4. $mark_length$: The length of the mark list. This allows us to reset all marks in constant time using $mark_length$ processors. This number is not larger than the number of transitions $(mark_length \le m)$.
- 5. In a list of size n, we store for each state $s \in S$ a variable $split_s : \mathbb{B}$. This indicates if the state will be split off from its block.

With this extra information, we can alter Algorithm 2 to work with labels. The new version is given in Algorithm 3. The changes involve the following:

- 1. Lines 7–9: Reset the mark list.
- 2. Line 11: Reset the split list.
- 3. Line 17: When marking the transitions, we do this for the correct action label, using $order_i$. Note the indexing into mark. It involves the offset for the state $source_i$, and $order_i$.
- 4. Lines 19–21: We tag a state to be split when it differs for any action from the block leader.
- 5. Line 24: If a state was tagged to be split in the previous step, it should split from its leader.
- 6. Line 29: If any block was split, the partition may not be stable w.r.t. the splitter.

To use Algorithm 3, we need to do two preprocessing steps. First, we need to partition the states w.r.t. their set of outgoing action labels. This can be done with an altered version of Algorithm 2, which does one iteration for each action label. For the second preprocessing step, we need to gather the extra information that is needed in Algorithm 3. This is done via sorting the action labels and subsequently performing some parallel segmented (prefix) sums [19]. In total the preprocessing takes $\mathcal{O}(|Act| + \log m)$ time. For details how this is implemented see the full version of this paper [14].

4.2 Complexity and Correctness

For Algorithm 3, we need to prove why it takes a linear number of steps to construct the final partition. This is subtle, as an iteration of the algorithm does not necessarily produce a stable block.

Theorem 3. Algorithm 3 on an input LTS with n states and m transitions will terminate in $\mathcal{O}(n + |Act|)$ steps.

Proof. The total preprocessing takes $\mathcal{O}(|Act| + \log m)$ steps, after which the while-loop will be executed on a partitioning π_0 which was the result of the

Algorithm 3: The algorithm for BCRP, the highlighted lines differ from Algorithm 2.

```
1 if i < n then
        unstable_i := false;
        unstable_{block_i} := true;
 3
 4 end
 5 do
        C := \bot;
 6
        if i < mark\_length then
 7
            mark_i := false;
        end
 9
        if i < n then
10
             split_i := false;
11
            if unstable_i then
12
              C := i;
13
            end
14
15
        end
        if i < m and block_{target_i} = C then
16
            mark_{off(source_i)+order_i} := true;
17
18
        end
        if i < m and mark_{off(source_i) + order_i} \neq mark_{off(block_{source_i}) + order_i} then
19
            split_{source_s} := true;
20
21
        end
22
        if i < n \& C \neq \bot then
23
             unstable_C := false;
            if split_i then
24
25
                 new\_leader_{block_i} := i;
                 unstable_{block_i} := true;
26
                 block_i := new\_leader_{block_i};
27
                 unstable_{block_i} := true;
28
                 unstable_C := true;
29
30
            end
        end
31
32 while C \neq \bot;
```

preprocessing on the partition $\{S\}$. Every iteration of the **while**-loop is still executed in constant time. Using the structure of the proof of Theorem 2, we derive a bound on the number of iterations.

At the start of iteration $k \in \mathbb{N}$ the total number of blocks and unstable blocks are $N_k, U_k \in \mathbb{N}$, initially $U_0 = N_0 = |\pi_0|$. In iteration k, a number l_k of blocks is split in two blocks, resulting in l_k new blocks, meaning that $N_{k+1} = N_k + l_k$. All new l_k blocks are unstable and a number $l'_k \leq l_k$ of the old blocks that are split were stable at the start of iteration k and are now unstable. If $l_k = l'_k = 0$ there are no blocks split and the current block C becomes stable. We indicate this with a variable c_k : $c_k = 1$ if $l_k = 0$, and $c_k = 0$, otherwise. The total number

of iterations up to iteration k in which no block is split is given by $\sum_{i=0}^{k-1} c_i$. The number of iterations in which at least one block is split is given by $k - \sum_{i=0}^{k-1} c_i$.

If in an iteration k at least one block is split, the total number of blocks at the end of iteration k is strictly higher than at the beginning, hence for all $k \in \mathbb{N}$, $N_k \geq k - \sum_{i=0}^{k-1} c_i$. Hence, $N_k + \sum_{i=0}^{k-1} c_i$ is an upper bound for k. We derive an upper bound for the number of iterations in which no blocks

We derive an upper bound for the number of iterations in which no blocks are split using the total number of unstable blocks. In iteration k there are $U_k = \sum_{i=0}^{k-1} (l_i + l_i') - \sum_{i=0}^{k-1} c_i + |\pi_0|$ unstable blocks. Since the sum of newly created blocks $\sum_{i=0}^{k-1} (l_i) = N_k - |\pi_0|$ and $l_i' \leq l_i$, the number of unstable blocks U_k is bounded by $2N_k - \sum_{i=0}^{k-1} c_i - |\pi_0|$. Since $U_k \geq 0$ we have the bound $\sum_{i=0}^{k-1} c_i \leq 2N_k - |\pi_0|$. This gives the bound on the total number of iterations $z \leq 3N_z - |\pi_0| \leq 3n - |\pi_0|$.

With the time for preprocessing this makes the run time complexity $\mathcal{O}(n + |Act| + \log m)$. Since the number of transitions m is bounded by $|Act| \times n^2$, this simplifies to $\mathcal{O}(n + |Act|)$.

5 Experimental Results

In this section we discuss the results of our implementation of Algorithm 3 from Sect. 4. Note that this implementation is not optimized for the specific hardware it runs on, since the goal of this paper is to provide a generic parallel algorithm. This implementation is purely a proof of concept, to show that our algorithm can be mapped to contemporary hardware and to understand how the algorithm scales with the size of the input.

The implementation targets GPUs since a GPU closely resembles a PRAM and supports a large amount of parallelism. The algorithm is implemented in CUDA version 11.1 with use of the Thrust library. As input, we chose all benchmarks of the VLTS benchmark suite for which the implementation produced a result within 10 min. The VLTS benchmarks are LTSs that have been derived from real concurrent system models.

The experiments were run on an NVIDIA Titan RTX with 24 GB memory and 72 Streaming Multiprocessors, each supporting up to 1,024 threads in flight. Although this GPU supports 73,728 threads in flight, it is very common to launch a GPU program with one or even several orders of magnitude more threads, in particular to achieve load balancing between the Streaming Multiprocessors and to hide memory latencies. In fact, the performance of a GPU program usually relies on that many threads being launched.

The left-hand side of Table 1 shows the results of the experiments we conducted. The |Act| column corresponds to the number of different action labels. The |Blocks| column indicates the number of different blocks at the end of the algorithm, where each block contains only bisimilar states. With #It we refer to the number of **while**-loop iterations that were executed (see Algorithm 3),

The source code can be found at https://github.com/sakehl/gpu-bisimulation.

² https://cadp.inria.fr/resources/vlts/.

	I		I	_							1_		I_
Benchmark name	1	Blocks		T_{pre}	-			$T_{Par-BCRP}/n$	_	$T_{Par-BCRP}$	T_{LR}	T_{Wss}	T_{Wms}
Vasy_0_1	2	9	16	0.50		0.06	1.78	0.003	0.023	0.87	2.29	0.49	0.45
Cwi_1_2	26	1,132	2,786	0.63		1.43	2.46	0.029		57.1	17	18.8	21.8
Vasy_1_4	6	28	45	0.56	1.01	0.04	1.61	0.001	0.022	1.58	4.78	1.68	0.62
Cwi_3_14	2	62	122	0.63	2.68	0.03	1.97	0.001	0.022	3.30	60	3.80	3.72
Vasy_5_9	31	145	193	0.84	4.22	0.04	1.33	0.001	0.022	5.06	134	35.3	3.45
Vasy_8_24	11	416	664	0.70	13.9	0.07	1.59	0.002	0.021	15	277	31.5	3.03
Vasy_8_38	81	219	319	1.12	6.64	0.04	1.46	0.001	0.021	7.76	127	35.1	5.94
Vasy_10_56	12	2,112	3,970	0.73	82.0	0.37	1.88	0.008	0.021	82.7	860	40.9	4.6(0.2)
Vasy_18_73	17	4,087	6,882	1.01	142	0.37	1.68	0.008	0.021	143	1,354	211	21.7
Vasy_25_25	25,216	25,217	25,218	159	519	1.00	1.00	0.027	0.021	678	21,960	t.o	416
Vasy_40_60	3	40,006	87,823	0.87	1,810	2.20	2.20	0.045	0.021	1,811	17,710	1,290	1,230
Vasy_52_318	17	8,142	15,985	2.52	338	0.31	1.96	0.007	0.021	340	11,855	368	152(20)
Vasy_65_2621	72	65,536	98,730	12.2	10,050	1.51	1.51	0.154	0.102	10,060	t.o	27,000	1,230
Vasy_66_1302	81	66,929	91,120	6.70	5,745	1.36	1.36	0.086	0.063	5,752	480,600	20,450	240(20)
Vasy_69_520	135	69,754	113,246	4.13	3,780	1.62	1.62	0.054	0.033	3,780	94,800	16,090	35.4
Vasy_83_325	211	83,436	148,012	4.41	3,093	1.77	1.77	0.037	0.021	3,097	57,190	21,500	5,880
Vasy_116_368	21	116,456	210,537	2.50	5,900	1.81	1.81	0.051	0.028	5,900	80,900	6,360	2,930
Cwi_142_925	7	3,410	5,118	4.85	238	0.04	1.50	0.002	0.047	243	3,363	220(30)	140(20)
Vasy_157_297	235	4,289	9,682	4.58	201	0.06	2.26	0.001	0.021	206	1,058	1,240	579
Vasy_164_1619	37	1,136	1,630	8.34	125	0.01	1.43	0.001	0.077	134	8,173	470(30)	46.8
Vasy_166_651	211	83,436	145,029	6.13	5,710	0.87	1.74	0.034	0.039	5,720	80,210	29,660	9,560
Cwi_214_684	5	77,292	149,198	3.58	6,948	0.70	1.93	0.032	0.047	6,952	19,250	440(30)	450(50)
Cwi_371_641	61	33,994	85,858	4.72	4,050	0.23	2.53	0.011	0.047	4,050	26,940	6,970	1,548
Vasy_386_1171	73	113	199	7.38	14.0	0.00	1.76	0.000	0.070	21	334	30.6	34.8
Cwi_566_3984	11	15,518	23,774	16.0	3,707	0.04	1.53	0.007	0.156	3,723	98,200	6,700	2,200(200)
Vasy_574_13561	141	3,577	5,860	71.5	3,770	0.01	1.64	0.007	0.643	3,841	144,810	11,700	1,853
Vasy_720_390	49	3,292	3,782	3.97	143	0.01	1.15	0.0002	0.038	147	2,454	1,633	183
Vasy_1112_5290	23	265	365	24.0	99.3	0.0003	1.38	0.0001	0.272	123	4,570	293	36.8
Cwi_2165_8723	26	31,906	66,132	37.0	23,660	0.03	2.07	0.011	0.358	23,700	140,170	9,700	1,965
Cwi_2416_17605	15	95,610	152,099	64.1	96,400	0.06	1.59	0.040	0.634	96,500	257,200	16,300(1100)	15,300
Vasy_6020_19353	511	7,168	12,262	221	11,690	0.002	1.71	0.002	0.954	11,910	107,900	34,000(2000)	19,230
Vasy_6120_11031	125	5,199	10,014	74.0	6,763	0.002	1.93	0.001	0.675	6,837	55,750	7,010	1,280
Vasy_8082_42933	211	408	660	281	1,149	0.0001	1.62	0.0002	1.739	1,429	17,272	5,530	2,030

Table 1. Benchmark results for Par-BCRP (Algorithm 3) on a GPU, times (T) are in ms. The right-hand side compares the total times from the different algorithms.

before all blocks became stable. The number of states and transitions can be derived from the benchmark name. In the benchmark ' X_N_M ', N*1000 is the number states and M*1000 is the number of transitions. The T_{pre} give the preprocessing times in milliseconds, which includes doing the memory transfers to the GPU, sorting the transitions and partitioning. The T_{alg} give the times of the core algorithm, in milliseconds. The $T_{Par-BCRP}$ is the sum of the preprocessing and the algorithm, in milliseconds. We have not included the loading times for the files and the first CUDA API call that initializes the device. We ran each benchmark 10 times and took the averages. The standard deviation of the total times varied between 0% and 3% of the average, thus 10 runs are sufficient. All the times are rounded with respect to the standard error of the mean.

We see that the bound as proven in Sect. 4.2 ($k \leq 3n$) is indeed respected, #It/n is at most 2.20, and most of the time below that. The number of iterations is tightly related to the number of blocks that the final partition has, the #It/|Blocks| column varies between 1.00 and 2.53. This can be understood by the fact that each iteration either splits one or more blocks or marks a block as stable, and all blocks must be checked on stability at least once. This also means that for certain LTSs the algorithm scales better than linearly in n. The preprocessing often takes the same amount of time (about a few milliseconds). Exceptions are those cases with a large number of actions and/or transitions.

Concerning the run times, it is not true that each iteration takes the same amount of time. A GPU is not a perfect PRAM machine. There are two key differences. Firstly, we suspect that the algorithm is memory bound since it is performing a limited amount of computations. The memory accesses are irregular, i.e., random, which caches can partially compensate, but for sufficiently large n and m, the caches cannot contain all the data. This means that as the LTSs become larger, memory accesses become relatively slower. Secondly, at a certain moment, the maximum number of threads that a GPU can run in parallel is achieved, and adding more threads will mean more run time. These two effects can best be seen in the $T_{alg}/\#It$ column, which corresponds to the time per iteration. The values are around 0.02 up to 300,000 transitions, but for a larger number of states and transitions, the amount of time per iteration increases.

5.1 Experimental Comparison

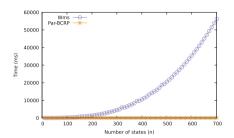
We compared our implementation (Par-BCRP) with an implementation of the algorithm by Lee and Rajasekaran (LR) [12] on GPUs, and the optimized GPU implementation by Wijs based on signature-based bisimilarity checking [2], with multi-way splitting (Wms) and with single-way splitting (Wss) [22]. Multi-way splitting indicates that a block is split in multiple blocks at once, which is achieved by computing a signature for each state in every partition refinement iteration, and splitting each block off into sets of states, each containing all the states with the same signature. The signature of a state is derived from the labels of the blocks that this state can reach in the current partition. Note that we are not including comparisons with CPU bisimulation checking tools; the fact that those tools run on completely different hardware makes a comparison problematic, and such a comparison does not serve the purpose of evaluating the feasibility of implementing Algorithm 3. Optimising our implementation to make it competitive with CPU tools is planned for future work.

The running times of the different algorithms can be found in the right-hand side of Table 1. Similarly to our previous benchmarks, the algorithms were run 10 times on the same machine using the same VLTS benchmark suite with a time-out of 10 min. In some cases, the non-deterministic behaviour of the algorithms Wms and Wss led to high variations in the runs. In cases where the standard error of the mean was more than 5% of the mean value, we have added the standard error in Table 1 in between parentheses. Furthermore, all the results are rounded with respect to the standard error of the mean. As a pre-processing step for the LR, Wms and Wss algorithms the input LTSs need to be sorted. We did not include this in the times, nor the reading of files and the first CUDA API call (which initializes the GPU).

This comparison confirms the expectation that our algorithm in all cases (except one small LTS) out-performs LR. This confirms our expectation that LR is not suitable for massive parallel devices such as GPUs.

Furthermore, the comparison demonstrates that in most cases our algorithm (Par-BCRP) outperforms Wss. In some benchmarks (Cwi_1_2, Cwi_214_684, Cwi_2165_8723 and Cwi_2416_17605) Wss is more than twice as fast, but in 16

other cases our algorithm is more than twice as fast. The last comparison shows us that our algorithm does not out-perform Wms. Wms employs multi-way splitting which is known to be very effective in practice. Furthermore, contrary to our implementation, Wms is optimized for GPUs while the focus of the current work is to improve the theoretical bounds and describe a general algorithm.



States	Run time	Wms (ms) Par-BCRP (ms)
10	1	1
100	182	5
200	1350	9
300	4463	13
400	10519	18
500	20508	22
600	35392	26
700	56183	30

Fig. 2. Run times of Par-BCRP and Wms on the LTS Fan_out_n .

In order to understand the difference between Wms and our algorithm better, we analysed the complexity of Wms [22]. In general this algorithm is quadratic in time, and the linearity claim in [22] depends on the assumption that the fan-out of 'practical' transition systems is bounded, i.e., every state has no more than c outgoing transitions for c a (low) constant. We designed the transition systems Fan_out_n for $n \in \mathbb{N}^+$ to illustrate the difference. The LTS $Fan_out_n = (S, \{a, b\}, \rightarrow)$ has n states: $S = \{0, \dots, n-1\}$. The transition function contains $i \xrightarrow{a} i + 1$ for all states 1 < i < n-1. Additionally, from state 0 and 1 there are transitions to every state: $0 \xrightarrow{b} i, 1 \xrightarrow{b} i$ for all $i \in S$. This LTS has n states, 3n-3 transitions and a maximum out degree of n transitions.

Figure 2 shows the results of calculating the bisimulation equivalence classes for Fan_out_n , with Wms and Par-BCRP. It is clear that the run time for Wms increases quadratically as the number of states grows linearly, already becoming untenable for a small amount of states. On the other hand, in conformance with our analysis, our algorithm scales linearly.

6 Conclusion

We proposed and implemented an algorithm for RCPP and BCRP. We proved that the algorithm stops in $\mathcal{O}(n + |Act|)$ steps on $\max(n, m)$ CRCW PRAM processors. We implemented the algorithm for BCRP in CUDA, and conducted experiments that show the potential to compute bisimulation in practice in linear time. Further advances in parallel hardware will make this more feasible.

For future work, it is interesting to investigate whether RCPP can be solved in sublinear time, that is $\mathcal{O}(n^{\epsilon})$ for a $\epsilon < 1$, as requested in [12]. It is also intriguing whether the practical effectiveness of the algorithm in [22] by splitting blocks

simultaneously can be combined with our algorithm, while preserving the linear time upperbound. Furthermore, it remains an open question whether our algorithm can be generalised for weaker bisimulations, such as weak and branching bisimulation [7,9]. The main challenge here is that the transitive closure of so-called internal steps needs to be taken into account.

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