

Verifying Very Large Industrial Circuits Using 100 Processes and Beyond

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Printing remark: The graphs are nicer when printed in colors.

Abstract

Recent advances in scheduling and networking open the way to the efficient exploitation of large-scale distributed computing platforms, such as computational grids and huge clusters. Such infrastructure holds a great promise for the highly resource-demanding task of verifying and checking large models, given that model checkers would be designed with a high degree of scalability and flexibility in mind.

In this paper we focus on the mechanisms required to execute a high-performance, distributed, symbolic model checker, on top of a large-scale distributed environment. We develop a hybrid algorithm for slicing the state space and dynamically distribute the work among the worker processes. We show that the new approach is faster, more effective, and thus much more scalable than previous slicing algorithms. We then present a checkpoint-restart module that involves very low overhead. This module can be used to combat failures which become probable with the size of the computing platform. However, checkpoint-restart is even more handy for the scheduling system to avoid reservation of large numbers of workers, thus making the distributed computation work-efficient. Finally, we discuss for the first time the effect of reorder on the distributed model checker. We show how the distributed system performs more efficient reordering than the sequential one.

We implemented our contributions on a network of 200 processors, using a distributed scalable scheme that employs a high-performance industrial model checker from Intel. Our results in testing execution on real-life examples, show that the system was able to verify models much larger than was previously possible.

1 Introduction

This paper presents several novel techniques to enhance distributed reachability computation. The techniques enable effective use of a network of 100 computers for the verification of large industrial hardware designs that could not be verified by previously available tools.

For a long time the state explosion problem has been the showstopper of BDD-based (symbolic) model checking [3]: The BDD structures simply can not squeeze into the RAM available to a single computer. SAT-based model checking [2] can find errors in very large systems, but is limited when used for verification [9]. In fact, BDD-based model checking is usually superior, when verification is required. Larger systems usually have deeper diameter and therefore SAT-based bounded model checking can cover smaller parts of their state space.

In recent years, several distributed BDD-based reachability algorithms have been suggested [8, 7, 6], which run on a network of communicating computers with distributed memory. Reachability is an important problem since model checking of all temporal safety properties can be reduced to it [1]. Distributed reachability exploits the memory modules and the computation power of a heterogeneous cluster of computers, where more and more machines can be employed on demand. The collective storage offered by the cluster RAM is utilized in a memory- and work-efficient manner, essentially operating as a yet another layer in the memory hierarchy.

However, in order to make these algorithms scalable for very large models which require hundreds of computers, several enhancements are necessary.

First, fast and effective slicing is needed, in order to accommodate frequent splits of memory content of overflowed computers. Second, a checkpoint/restart mechanism is needed in order to recover from a single computer failure and in order to better utilize clusters of computers when memory requirements vary significantly during computation. Finally, dynamic BDD variable reordering should be adapted to work well with the distributed algorithm.

Our work provides solutions to all the requirements presented above. We developed a *hybrid* algorithm for slicing very large sets, quickly and effectively. The user provides the algorithm with measures for an effective slicer and the algorithm searches for an adequate one. The algorithm is aimed at spending as little time as possible in finding an adequate slicer, not necessarily the best one. It starts with a fast estimated computation. If no adequate slicer is found, it gradually applies more precise computations. We compare our hybrid algorithm with the fast estimating algorithm `Estimated` [5]. We show that our algorithm produces much less duplications. We also compare our algorithm with the exhaustive algorithm `Exhaustive` [8] which is better or equal to other exhaustive algorithms [4, 11, 10]. We show that it is faster than `Exhaustive`. In fact, the difference in run time increases when the size of the BDD or the size of its support increase.

We also propose a non-coordinated checkpoint/restart mechanism as part of the distributed reachability computation. In the distributed reachability analysis [8] each worker *owns* a subset of the state space, and iteratively computes the set of reachable states within its ownership. It may also find states owned by others, which it sends to their owners. Likewise, it receives owned states, found by others. In the checkpoint mechanism each worker occasionally freezes its configuration, including the set of states it owns, the set of states computed so far, the iteration number, and the BDD variable ordering. Restart is performed by finding a set of configurations, all taken from the same iteration, whose ownership covers the whole state space. A set of new free workers is then initialized with these configurations and resumes the computation.

The checkpoint/restart mechanism is particularly useful when running on a non-dedicated network. Two tasks running on such a network may reach their memory peak at the same time, thus blocking each other. It then might be necessary to freeze one of them, and enable the other to continue. When memory requirement of the active task reduces, the frozen one can be resumed. In addition, when memory requirements vary significantly during computation, an effective utilization will require clusters of varying sizes at different stages of the computation. This will be done by freezing the active workers, and restarting them on a different cluster with an appropriate size.

In order to maintain the effectiveness of the dynamic reorder we suggest a distributed paradigm to control the points at which dynamic variable reordering is performed. In sequential computation, reorder is invoked after garbage collection, if the BDD size exceeds a certain threshold. The distributed computation applies the same policy. In addition, for each worker, it uses two new controlling operations: enforcing

reorder when an overflow occurs; and updating the threshold following an action which reduces the BDD size. Reorder when overflow occurs may save unnecessary splits.

Another improvement to the BDD package enforces timeout on BDD operations that do not terminate within a reasonable time. Usually this is due to the size of their operands. We then split the BDDs and resume the operations on two smaller BDDs.

To show the usefulness of our scheme we have implemented it as a large-scale distributed engine with more than 100 computers, that uses a high-performance model checker. We ran our experiments on clusters composed of ordinary PCs. Our results show that the system can *verify* (apply full reachability to) much larger models than could previously be verified. In addition our results show that in some cases when the distributed algorithm needs more processes than available, it still reaches further step than SAT-base bounded model checking does.

In summary, the contribution of the paper are:

- A fast and effective slicing with small memory overhead.
- A checkpoint/restart mechanism.
- Enhanced BDD package: An adaptive dynamic variable reorder and timeout on BDD operations.
- Orthogonality to high-performance model checking: all features of the sequential model checking remain effective in the distributed framework.

All of the above allows to *verify* large industrial components.

The rest of the paper is organized as follows. Section 2 presents a new algorithm for fast and effective slicing of very large sets. Section 3 describes the checkpoint/restart mechanism. Section 4 presents our distributed reachability analysis including paradigm for dynamic variable reordering, and presents our experimental results on verification of large industrial designs. Finally, Section ?? gives our conclusions.

2 Hybrid Algorithm for Slicing Very Large Sets

In this section we present a new algorithm for slicing very large sets quickly and effectively. The approach makes use of user-supplied measures of effectiveness: the algorithm simply searches for a slicer that meets the measures. The algorithm attempts to reduce the time spent in finding a slicer which is sufficiently effective; it does not necessarily search for the best slicer. Rather than checking all variables in the support of the set to be sliced, which is the method previously used for slicing, the proposed algorithm makes use of the abundance of good slicers in the support to pick one from a randomly selected sample.

The algorithm gets as its input a set of states with characteristic function f and returns a variable v called slicer, which slices f into two subsets: $f \wedge v$ and $f \wedge \bar{v}$. Such slicing is *effective* if two requirements are fulfilled. First, the size of each of the subsets is smaller than the size of f itself: $\frac{\max\{|f \wedge v|, |f \wedge \bar{v}|\}}{|f|} < \delta_1$.

Second, the *duplication* is not too big: $\frac{|f \wedge v| + |f \wedge \bar{v}|}{|f|} < \delta_2$. The minimum reduction factor and the maximum duplication factor δ_1, δ_2 are provided by the user, or the higher-level procedure calling the algorithm.

The algorithm proceeds through a sequence of three phases. In each consecutive phase the algorithm spends more time than in the previous phase, trying to find an effective slicer. Once an effective slicer is found the algorithm declares success and terminates. After three unsuccessful phases the algorithm returns the best slicer it found so far.

In order to test the effectiveness of a candidate slicer the BDDs of $f \wedge v$ and $f \wedge \bar{v}$ must be built and their relative sizes must be measured. This is time and memory consuming. In contrast, one can estimate the sizes of the slices in a single scan of the BDD of f without creating a new BDD [12]. Estimation is a lot faster than precise calculation and requires much less resources.

In the first phase the algorithm employs the method `Estimated` [5] to search for an effective slicer. This method initially computes an estimate to the size of $f \wedge v$ and $f \wedge \bar{v}$, for each variable v in the support of f . Then it selects as a slicer the variable v for which the maximum of the estimates for $f \wedge v$ and $f \wedge \bar{v}$ is minimal among all other variables. Next, precise calculation is used to derive the effectiveness of v as a slicer. If v is found to be effective, the algorithm terminates, otherwise it proceeds to the next phase.

The second phase starts by randomly selecting a subset $varSet$ of variables out of the support of f . The size of $varSet$ depends on the required degree of confidence of finding at least one effective variable (See Subsection 2.1). $effectiveSet$ holds all the variables in $varSet$ that were first estimated as effective and only those that seem to be effective are checked precisely. If $effectiveSet$ is empty the second phase ends unsuccessfully. Otherwise, the best slicer from $effectiveSet$ is selected by `Exhaustive`. We remark that the `Exhaustive` procedure itself is not different from the slicing mechanisms described in [8]. Thus, in this paper, we use it as a black box.

The third phase is similar to the second. The difference is that $effectiveSet$ now holds all the variables from $varSet$ that slice effectively using precise check. Finally, if the third phase fails and none of the variables is effective, the `MostEffective` is selected among the variables that computed in the third phase and this variable is returned.

Figure 1 describes the algorithm `slice` for finding a slicer. Lines 1-2 describe the first phase that uses the `Estimated` method to select a slicer v . If v found to be effective the algorithm terminates. Lines 3-6 describe the second phase where $varSet$ is randomly selected from the variables in the support of f . Then $effectiveSet$ gets only the variables that are effective slicers. This computation is done by first applying a fast estimated check and only then a precise check. The precise check is applied only on slicers that are estimated to be effective. Finally, the algorithm `Exhaustive` is used to find the best slicer out of $effectiveSet$. Lines 7-9 describe the third phase where precise check is used on all the variables in $varSet$. Line 10 is the last phase where the most effective slicer found so far is returned.

```

function Hybrid(f)
1  v=Estimated(f)
2  if effective(v,precise) return v
3  varSet=randomselect(support(f))
4  effectiveSet={v | v ∈ varSet ∧ effective(v,fast) ∧ effective(v,precise)}
5  if effectiveSet ≠ ∅
6    return Exhaustive(f,effectiveSet)
7  effectiveSet={v | v ∈ varSet ∧ effective(v,precise)}
8  if effectiveSet ≠ ∅
9    return Exhaustive(f,effectiveSet)
10 return MostEffective(f,effectiveSet)

```

Figure 1: Pseudo-code for the slicing algorithm `Hybrid`

2.1 Size of the Random Selected Subset

In this section we discuss the relation between the confidence in finding at least one effective variable and the number of samples. Lemma 1 defines this relation.

Lemma 1 [*Sample size required*] *Let sup be the size of the support of a set. Let ef be the number of effective slicers in the support ($ef \leq sup$). Let s be the number of randomly selected variables ($s \leq sup$). Let pr be the confidence in finding at least one effective slicer out of s samples. Then,*

$$pr \geq 1 - \left(1 - \frac{ef}{sup}\right)^s$$

Proof: The probability to get ineffective variable in the first random sample is $1 - \frac{ef}{sup}$. The probability to get ineffective variable in the i th random sample is $1 - \frac{ef}{sup - (i-1)}$, assuming all previous samples were not effective. The confidence in finding at least one effective variable in s samples

$$pr = 1 - \prod_{0 \leq i < s} \left(1 - \frac{ef}{sup - i}\right).$$

Set	BDD sizes range	support sizes range	number of sets of states
Small	0.5 - 3 Million	70	25
Large	0.5 - 6 Million	239 - 255	46
Extra large	0.5 - 7 Million	687 - 712	18

Table 1: Benchmark suite characteristics. For each set of examples we give the BDD size of the sets of states and the support size.

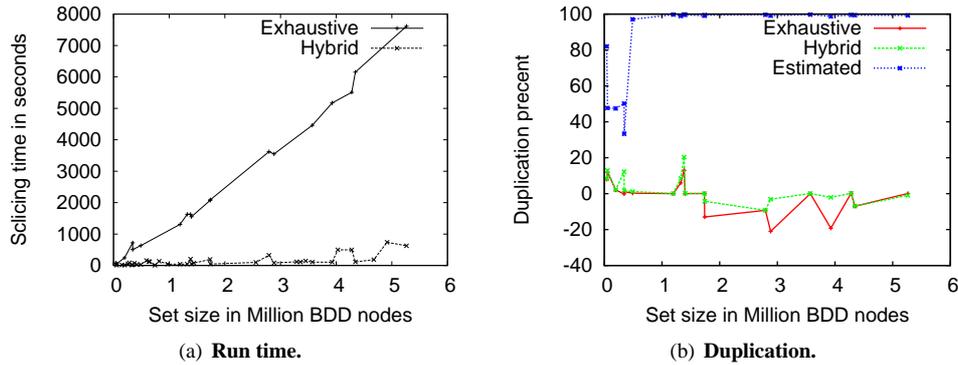


Figure 2: comparing the slicing algorithms, support size 687 - 712.

Since $1 - \frac{ef}{sup-i} \leq 1 - \frac{ef}{sup}$ for all i , $pr \geq 1 - \left(1 - \frac{ef}{sup}\right)^s$.

Q.E.D.

Our experimental results (Figures 6(a), 6(b), explained latter) show that the minimum percentage of effective slicers is 4%. Therefore, confidence in finding at least one effective variable converges to 100% exponentially fast in the number of samples. More importantly it does not depend on the number of variables. If for example we want 90% confidence that we will get at least one effective variable and 5% of the variables are effective we need only 45 samples.

2.2 Experimental Results

We compare three slicing algorithms. The new algorithm `Hybrid`, presented in Figure 1; The exhaustive algorithm `Exhaustive` when working on the entire set of support; And the fast estimation `Estimated` when working on the entire set of support.

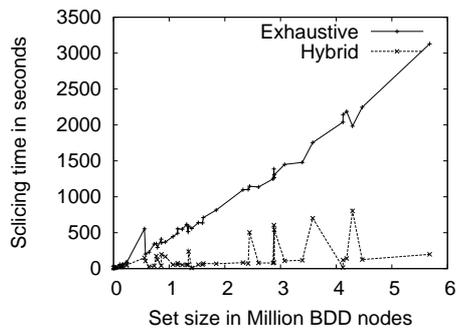
We use three sets of examples, each with different support size. Each set includes variable size of BDDs from half a million to 7 millions nodes. The characteristics of the three sets in Table 1.

2.2.1 Slicing Efficacy and Memory Overhead

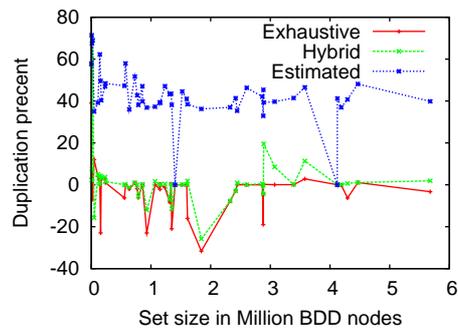
We now analyze the run time and the duplication by the different slicing algorithms. Figures 2(a), 3(a), 4 display run time of the slicing algorithms. In each graph the run time of `Hybrid`, and `Exhaustive` algorithms is given compared to the size of the set being sliced. Figure 4 shows the benefit of including the `Estimated` algorithm into `Hybrid`. For all the sets in Figure 4, `Hybrid` succeeds to find an effective slicer in the first phase using the `Estimated` algorithm. Therefore, `Hybrid` needs only a few seconds compared to up to two hours for the `Exhaustive` algorithm.

Figures 2(a), 3(a) show that the run time of the `Exhaustive` algorithm increases proportionally to the BDD size and increases proportionally to the support size, while the `Hybrid` algorithm run in constant time.

Figures 2(b), 3(b) compare the duplication obtained by the `Estimated` algorithm to that obtained by the `Hybrid` algorithm. In each graph, the percentage of duplication in the `Estimated` and in the `Hybrid` algorithms is given compared to the size of the set being sliced. The graphs show that when



(a) **Run time.**



(b) **Duplication.**

Figure 3: comparing the slicing algorithms, support size 239 - 255.

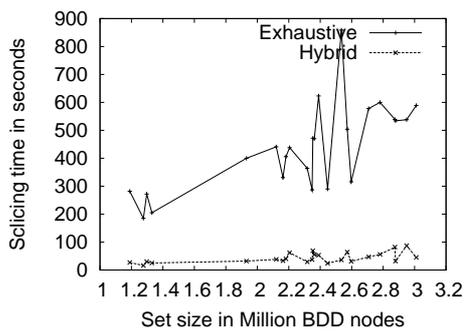


Figure 4: comparing the run time of the slicing algorithms, support size 70

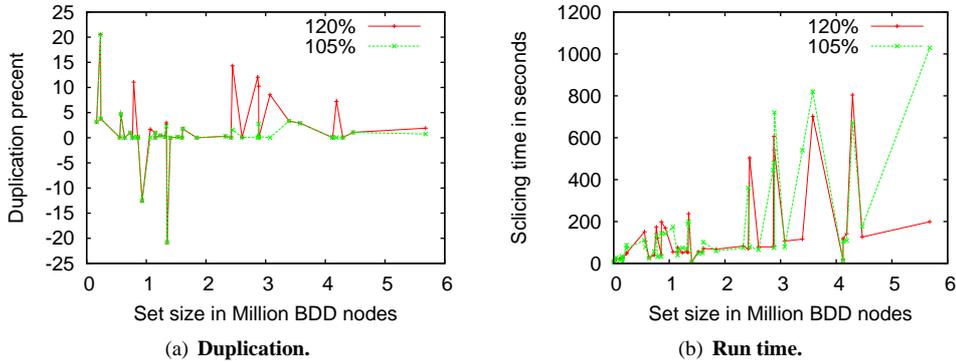


Figure 5: Support size 239 - 255. 105 means maximum duplication factor 105%. 120 means maximum duplication factor 120%.

the support size increases, the Estimated algorithm usually finds poorer slicing in terms of duplication. When the size of the support is 239 - 255 variables (Figure 3(b)), the average duplication of the Estimated algorithm is 50%, while the Hybrid algorithm creates 7% duplication on average. When the size of the support is 687 - 712 variables (Figure 2(b)), the average duplication of the Estimated algorithm is 89%, while the Hybrid algorithm creates 3% duplication on average.

Figures 2(b), 3(b) also compares the duplication obtained by the Exhaustive algorithm with the Hybrid algorithm. The percentage of duplication of the Exhaustive and the Hybrid algorithms is given compared to the size of the set being sliced. We set the maximum duplication factor δ_2 to be 120%. We set the minimum reduction factor δ_1 to be 85%. For all set sizes that are not too small (larger than 100K BDD nodes), the resulting slicer creates duplication which is smaller than the maximum duplication factor. When the set size is very small, no effective slicer was found by any of the three phases. Thus, the final phase finds a slicer with duplication of 150%. The small memory requirement by such small sets makes slicing of them not effective (with higher duplication than required).

In some cases the duplication may be negative. This means that the sum of the sizes of the two subsets is less than the original set size. The Exhaustive algorithm finds slicers with very small duplication down to 30% smaller than the original set size. The Hybrid algorithm may miss these slicers since it stops as soon as it finds an effective slicer.

2.2.2 Changing the Measures of Effectiveness

Figures 5(a), 5(b) present the effect of changing the maximum duplication factor from 120% to 105% on run time and on duplication. In each graph the duplication is given compared to the size of the set being sliced. Figure 5(a) presents the duplication for the maximum duplication factors of 105%, 120%. For all set sizes, the final slicer creates duplication which is smaller than the maximum duplication factors, hence, the duplication with 105% is less than or equal to the duplication with 120%.

Figure 5(b) presents the run time for duplication factors 105% and 120%. In most of the cases the run time of the algorithm is longer when the maximum duplication factor is 105%. In cases the algorithm needs to run more phases, the run time with 105% can take up to five times longer than that with 120%. Since the algorithm uses a random selection, different runs may terminate with different results. The random selection causes sometime deprecate results where the run time takes comparably longer when using larger maximum duplication

2.2.3 Percentage of Effective Slicers

The experiments presented in this Section demonstrate that for different set sizes, regardless of the BDD order, at least 4% of the variables are effective slicers. Figure 6(a) presents the percentage of effective slicers in different sets. The percentage of effective slicers is given compared to the size of the set being sliced. Figure 6(a) shows that regardless the set size, a minimum of 4% of the slicers are effective. This

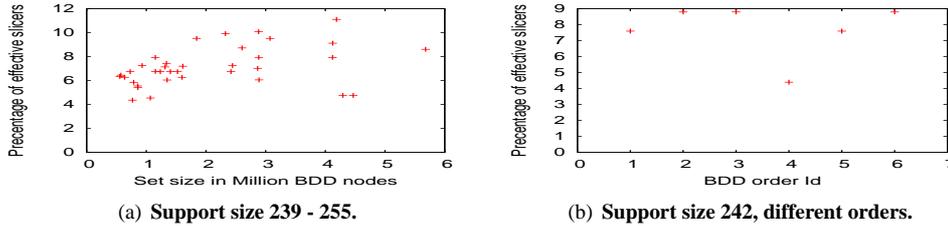


Figure 6: Percentage of effective slicers.

means that the confidence in finding at least one effective slicer converge to 100% exponentially fast in the number of samples (see Section 2.1).

Figure 6(b) presents the percentage of effective slicers in a single set with different BDD orders. This example has 242 variables in the support and the set size is 2.4 million BDD nodes with the best order. The percentage of effective slicers is given for each order. Figure 6(b) shows that even when we change the BDD order, as happen in the distributed reachability algorithm, a minimum of 4% of effective slicers is maintained.

3 The Checkpoint Restart Algorithm

In this section we briefly describe the iterative BDD-based distributed algorithm for reachability [7]. We explain how to extend this algorithm with checkpoints and how to exploit these checkpoints in order to restart the reachability algorithm when needed according to some scheduling policy. Finally we present experimental results which show that the associated overhead is negligible.

The basic paradigm followed by the algorithm is to compute the set of reachable states which are reachable from a given set of initial states. Starting from the set of initial states, at each iteration the set R is computed, consisting of reachable states found so far, and the set N of *undeveloped* states, that do not belong in R and are reachable from R in a single step, whose successors have not yet been found.

The distributed algorithm runs on a network of communicating workers with distributed memory. A set of *window functions* defines for each worker the subset of states it *owns*. This set is *complete*, meaning that it covers the whole state space. Worker id with window function W_{id} computes the sets R_{id} and N_{id} , both subsets of W_{id} .

Three coordinators control the distributed operation: the *pool manager* keeps track of the free processes; the *exchange coordinator* maintains the window functions of the active workers, and the *small coordinator* joins together the windows of workers whose memory utilization decreases below a certain threshold.

Figure 7 describes an extension of the distributed algorithm with checkpoint-restart capability, called **reach_checkpoint**. The pseudo code is described for a single worker. For brevity, we omit the worker subscript id from R_{id} , N_{id} , and W_{id} . We remark that the sets R and N , and the window function W , may change during the execution.

The algorithm follows the same lines of the distributed reachability algorithm, except at the end of each iteration workers sometimes store checkpoints. The data stored in a checkpoint consists of R , N , W , the iteration number $\#it$, and its current BDD order bdd_order (line 9). The checkpoint of a worker may be stored on a persistent storage system, e.g., a distributed file system such as NFS, or simply on the private disk of a peer worker (in which case it is assumed the peer worker does not crash in case the worker does).

Recall that the basic reachability paradigm is an iterative, synchronous process. Thus, the collection of all checkpoints from all workers at the end of an iteration forms a consistent view of the global reachability process at that point.

If a restart is needed because of a failure, or due to rescheduling of the reachability process on another distributed system, the collection of checkpoints may set a starting point for pursuing the computation. The restart algorithm searches for a set of checkpoints taken from the same iteration, which forms a complete set of window functions. If an incomplete set is found, indicating that some but not all the workers succeeded

to store checkpoints for the corresponding iteration previous to the abort, then the algorithm searches for a complete set that was stored at the end of a previous iteration. Such a set is guaranteed to exist since the workers follow the same policy at the end of which iterations checkpoints are stored, and since a previous checkpoint is never removed before the current global checkpoint is known to be complete (e.g., at the end of the next iteration).

Every active worker in the restarted process is restored using its local checkpoint data, and is replaced by a worker from the free pool in the new distributed system. The new worker restores R and N according to the BDD order bdd_order and assumes W as its window function.

```

function reach_checkpoint( $R, W, N$ )
1  Loop until termination()
2   $N = \text{Image}(N)$ , split if needed
3  send non-owned states ( $N \setminus W$ ) to their owners
4   $N = N \cup$  (received states in  $W$  from others),
   split if needed
5   $N = N \setminus R$ 
6   $R = R \cup N$ 
7  Collect_small( $R, W, N$ )
8  if ( $W = \emptyset$ ) return to pool
9  Check_point( $R, W, N, \#it, bdd\_order$ )

```

Figure 7: Pseudo-code for a worker in the distributed reachability computation with checkpoints.

3.1 Experimental Results

The resources consumed by the distributed algorithm are evaluated by considering the following two figures. The *reservation* is the size of the cluster of machines carrying the computation. These machines are either actively taking part in the computation, or otherwise they are part of the free pool, in which case they might not be carrying any useful computation as they are reserved as potential additional resources for the reachability computation. The *utilization* is the number of active non-free workers that are actually taking part in the reachability computation. Of course, at any point during computation the utilization is less than the reservation.

Figure 8 presents the utilization and reservation of workers during the distributed reachability computation. The graph shows how the checkpointing mechanism is used in order to reduce reservation to a minimum. Checkpointing is used to vary the reservation, starting from a small cluster with only 10 machines. When more than 10 machines are required the run temporarily halts, the last checkpoint is moved to a larger reservation with more machines, and the computation is resumed on that cluster. In this way the free pool (of idle machines) is never too large relative to the utilization.

Clearly, yet another contribution of the checkpoint restart mechanism is in the case of termination as a result of failure in one of the resources. In this case, in the absence of fail over mechanism, the amount of resources that would have been wasted grows together with the integral over the reservation from the start till the iteration where the failure appeared. Furthermore, the more machines take part in the computation, and the more iterations are involved in the computation, the higher the chance of a failure. Thus, the importance of the checkpoint restart mechanism increases with the number of iterations to fixpoint, and with the scale of the model checked (as indicated by the reservation).

Figure 9 compares the run time of image computation and the time to store the checkpoint data. The graph shows for each set size the run time required for image computation and the run time required to store the checkpoint data. The graph shows that for all set sizes the checkpoint run time takes less than 20 seconds. Moreover, if there is no job failure, there is almost no overhead for storing checkpoints.

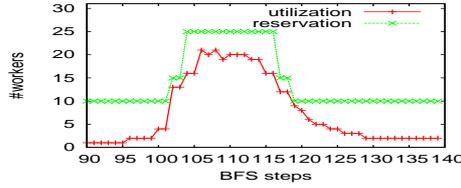


Figure 8: *reservation* is the size of the cluster of machines carrying the computation. *utilization* is the number of active non-free workers.

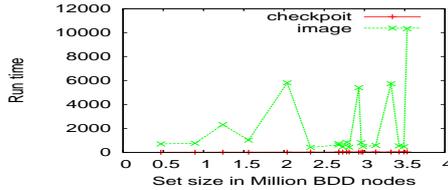


Figure 9: Run time of image computation and the time to store the checkpoint data.

4 Distributed Reachability Analysis for Very Large Circuits Using 100 PCs

In the previous sections two enhancements to distributed reachability analysis were discussed. This section describes the extensions to the algorithm **reach_checkpoint** which enable a high performance distributed reachability analysis for very large circuits using 100 PCs. With these extensions, the algorithm verifies circuits that could not be verified by any other tool. Furthermore, although an overflow in the required number of workers occurs in several cases, the distributed scheme still reaches an iteration much farther than that reached by the sequential BDD based model checker.

In order to have our distributed scheme scale out, two additional extensions to **reach_checkpoint** are needed: *distributed reorder* and *BDD operations timeout*. We discuss the two extensions and then give experimental results.

4.1 Distributed Dynamic Variable Reordering

Dynamic variable reorder suggested by Rudell [13], works well for the sequential algorithm. Here we show how to use it also with our distributed approach.

Rudell's algorithm is called by the BDD package according to the growth in the number of BDD nodes. A dynamic reorder threshold *dr_this* is maintained, where the next threshold is set after each invocation of reorder, according to the number of nodes in the new order. The threshold is examined after each cycle of the garbage collection, and variable reordering is triggered if the number of nodes allocated after the garbage collection is greater than *dr_th*.

In the distributed scheme the BDD package uses Rudell's algorithm in the same way. However, since in the distributed scheme there are events such as splits and joins which affect the size of the BDD package, the distributed algorithm also controls the value of *dr_th* externally. The worker forces the BDD packages to adjust the value of *dr_th* after splitting a worker, in which the number of nodes reduces dramatically, and after exchanging nonowned states, in which the number of nodes may decrease or increase due to exchange of non-owned states.

In addition, in case of overflow, triggering reorder may reduce the size of the BDD and thus avoid the costly splitting. Therefore, when an overflow occurs after a long computation of a micro step but before the micro step is completed, the worker invokes reorder and then tries to complete the micro step computation. However, in case the BDD package triggered reorder just before the micro step overflowed, the worker avoids the additional reorder since it is unlikely to prevent the splitting.

Circuit	#vars	Overflow step	Overflow	R
H21	274	55	3,203,064	
H20	276	44	3,922,742	
I1	147	98	8,006,120	
H11	300	44	5,211,955	
I3	793	46	5,557,672	
I3s	439	54	7,076,762	
s1423	88	14	9,705,214	

Table 2: Benchmark suite characteristics. In each example we give the step in which the memory requirements by the sequential model checker overflow and the size of the BDD representing the set of reachable states R at that step.

4.2 Escape from BDD Operation Livelock Using Timeouts

BDD engines use cache for previously executed BDD operations. Using this cache, the run time commonly becomes linear in the sizes of the BDD operators, rather than exponential. Since the size of the cache cannot hold all the BDD operations, the engine replaces old results with new ones. Sometime the result of a BDD operation that has been replaced is needed, and therefore recalculated. Recalculation increases the run time, and in some cases, can cause the execution of a single BDD operation to proceed for hours.

In the distributed scheme a split can help a single worker if it stuck on a single BDD operation, as the size of the cache is effectively doubled as a result of the split. To this end, if a single micro step turns out too long, it is stopped as if a memory overflow occurred, and split is invoked. Our experiments show that cases in which a large number of recalculations take hours can be efficiently avoided in this way.

4.3 Experimental results

Our parallel testbed consists of 100 PC machines, each a two way 2.4GHz Pentium 4 processors with 1GB memory. For optimal utilization of this configuration we let two workers execute on the same machine. The communication between the nodes uses a Fast Ethernet. The sequential runs uses a PC machine consisting of four way 3.1GHz Pentium 4 with 4GB memory.

The distributed algorithm that we have tested follows **reach_checkpoint** enhanced with the algorithm **Hybrid**, the distributed dynamic reordering and the micro steps timeout. The external model checker used by the distributed algorithm is a high-performance industrial tool from Intel.

We conducted our experiments using examples for which the fixpoint was never reached before, such as the s1423 design from the ISCAS89 benchmarks. We remark that other examples from this benchmark suit, such as s3330, s1269 and s5378, require only a single process when using the high-performance model checker of Intel. Thus, they are not suitable as benchmarks for the distributed system. In addition to s1423, we experimented with six large examples which are components in Intel’s designs.

The characteristics of the six test cases are given in Table 2.

The distributed reachability analysis results are given in Table 3. Four examples reached fixpoint and the verification completed. Three examples required more workers than was available to us at this point (we did not always have all 100 machines at our disposal), therefore worker overflow occurred at some step, always much further than the step reached by the sequential model checker.

We next compare the results in Table 3 to the results of the high performance industrial SAT model checker tool of Intel. The SAT model checker could not complete the verification of any of the examples. Computing bounded model checking with timeout of 10,000 seconds SAT reached the bounds of 85 and 94 on I3s and I3, respectively.

We also compare the results in Table 3 to the results obtained in previous distributed symbolic model checking [7] and [8]. In [8] a high performance model checker was used, still, s1423 reached only step 17, while the distributed algorithm reached step 19. Other examples from ISCAS89 are so small they were completed by the sequential model checker. Other examples from [8] are not published to the public. In [7] a non-sophisticated model checker was used, therefore a relatively small example such as s3330 required

Circuit	Fixpoint	Max workers	Time	$\max_{ite} \sum_i R_i $	at Seq Overflow	
					$\sum_i R_i $	#workers
H21	85	3	23h			
H20	85	9	11h			
I1	139	25	70h	15.5M	6.6M	3
H11	98	7	28.5h	4.4M	1.3M	4
I3	WOvf(60)	>50		47.2M	7.1M	5
I3s	WOvf(118)	>150		358.8M	7.1M	4
s1423	WOvf(19)	>200		208.3M	8.8M	8

Table 3: Distributed reachability on the benchmark suite. Four examples reached fixpoint and verification completed. Three examples required more workers than was available to us, therefore worker overflow occurred. The Max workers column describes the maximum number of active workers during the computation. The run time in case the verification is completed is given in hours. Two measures are given for the iteration the sequential algorithm overflows: The sum of sizes of the BDDs representing the subsets of reachable states, and the number of active workers at this iteration

54 workers to complete. The high performance model checker used in this work can complete this example using a single worker.

It is especially interesting to compare the Tables 2 and 3. It turns out that at the point where the sequential algorithm overflows, the aggregate space requirement by the distributed algorithm (given in the tables as the size of R in BDD nodes) is *smaller* than the corresponding size in the sequential algorithm! This means that the distributed algorithm is more efficient in maintaining its data structures, sometimes to a factor of two or more. This comes as a surprise, since the common wisdom would have expected some overhead and duplication of work, rather than increased efficiency.

The explanation, however, is straightforward. Recall that with the distributed scheme reorder is optimized individually at every worker, taking into account the worker data only. In this way, the distributed algorithm becomes much more efficient in its reorder activity, as order at every worker better fits the worker data. The overall effect is an aggregate reduction in the number of BDD nodes, which implies improved overall efficiency. Furthermore, as a side-effect it also implies a better sequential algorithm that follows by simulating the distributed parallel scheme.

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