# P systems simulations on massively parallel architectures

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Molivation

The (SAT)isfiability problem

Parallel design of the simulation

#### Performance Evaluation

Performance on shared memory platform

Performance on distributed memory platform

Performance on a GPU-based cluster

#### Conclusions

**Appendix** 

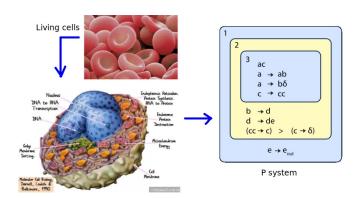




- Membrane computing is an emergent branch in Natural Computing.
- It is based on the behaviour of living cells to define bio-inspired computation devices, called **P systems**.
- P systems computational features include:
  - Polynomial time solutions to NP-Complete problems by trading time for space.
  - Modeling of biological phenomena.
- We focus on the family of recognizer P systems with active membranes.



#### P system apperance



#### P system simulation

- ▶ No P systems implementations neither *in vivo* nor *in vitro*.
- Only P systems simulations on silicon.
- Simulation of P systems poses challenges in three different aspects:
  - The intrinsic massively parallelism (like cells).
  - The exponential memory requirements (trading time for space).
  - The non-intensive floating point nature (for recognizer P systems).

**Work Motivation**: Which parallel platform is best suited for the P system simulation?



The (SAT)isfiability problem



# The SAT problem

- ► The first-known NP-complete problem (Stephen Cook, 1971 [3]).
- Formula in CNF with a conjuction of clauses formed by a disjuntion of literals.
- Literal is either a variable or its negation.
- ➤ To determine whether exists a truth assignment to its n variables.
- Paramount importance in many computer science areas:
   Hardware design, algorithmc, etc.

(A or  $\neg$ B) and ( $\neg$ A or B)



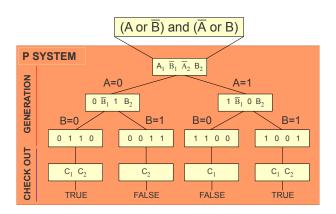
# Solving the SAT problem through P systems

- The P system computation for SAT is described in [4].
- It can be summarized in four stages:
  - Generation. All possible truth assignments to the variables are generated by using P systems rules. 2<sup>n</sup> internal membranes are created and each one encodes a truth assignment to the variables of the formula.
  - 2. **Synchronization.** The objects encoding a true clause (a partial solution to the CNF formula) are unified in the membrane.
  - Check out. The goal here is to determine how many (and which) clauses are *true* in every internal membrane (that is, by the assignment that represents).
  - Output. Internal membranes encoding a solution send an object to the skin. If the skin has such object from some membrane, the object Yes is sent to the environment. Otherwise, the object No is sent.



# Solving the SAT problem through P systems (II)

A small example



## Outline

Parallel design of the simulation



- 1. P systems are massively parallel, having a double level of parallelism:
  - The first level of parallelism is among membranes (coarse-grained).
  - ► The second level of parallelism is within each membrane (fine-grained).
- P systems create an exponential workspace (for the SAT equation 1).

$$Size = 2^{n} (membranes) * k(objects) * 4(uint) Bytes.$$
 (1)



# Parallelism among membranes

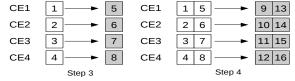
#### Sequential Membranes layout

	1 2 Step 1 Gener															ration	
	1	2	3	4 Step 2													
	1	2	3	4	5	6	7	8	St	tep 3	3						
Γ	1	2	3	1	5	6	7	g	a	10	11	12	13	1/1	15	16	Step 4

#### **Parallel Membranes layout**

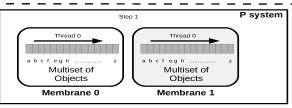


#### Generation

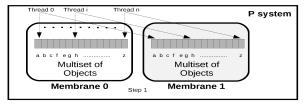


#### Parallelism within each membrane

#### Sequential Membrane Generation

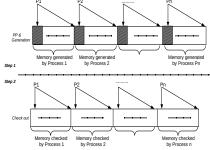


#### Parallel Membrane Generation



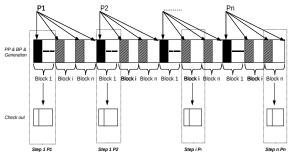


- ► The shared memory space is equally distributed among processes.
- ► Master process creates as many membranes as OpenMP processes.
- Initial membranes are located at the beginning of each memory space for each process.
- Each process performs generation stage in parallel, and then Check out.



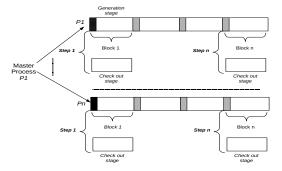
# Data locality on the Shared memory implementation (OpenMP)

- Many cache misses with the previous (first writes and then reads).
- Block-based data layout to improve locality.



# Distributed memory implementation(MPI)

- Similar to the previous one but in a distributed memory.
- Both implementations: Blocking and non-Blocking.

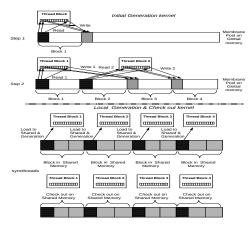


- A thread block per membrane and a thread per object (or set of objects).
- CPU creates as many membranes as GPUs in the system.
- First attempt with two kernels: First Generation and then Check out.
- Need of global synchronization among blocks.
- Performance issues: Only global memory accesses



# Blocking on GPUs (CUDA)

Taking advantage of the shared memory.



Performance Evaluation

# Outline

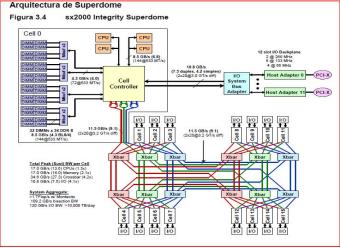
#### Performance Evaluation



### Hardware environment

- Shared memory platform is a HP Integrity Superdome SX2000 endowed with 64 CPUs
  - Intel Itanium 2 dual-core Montvale.
  - 16 Kbytes L1, 256 Kbytes L2, 18 Mbytes L3.
  - Total DRAM memory available is 1.5 Tbytes.
  - Interconnection network is a 4x DDR Infinihand.
- Distributed memory system is a HP BladeSystem with 102 nodes
  - Dual socket containing a quad-core Intel Xeon E5450.
  - 12 MBytes L2 cache.
  - DRAM memory capacity for the whole system is 1072 Gbytes.
  - Interconnection network is also a 4x DDR Infiniband.
- GPU-based platform includes
  - Four-socket, guad-core Intel Xeon E5530 socket.
  - 8 MBytes L2 cache.
  - DRAM memory capacity 16 Gbytes
  - 4 Tesla C1060 with up to 16Gbytes of video memory.





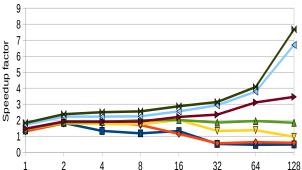
Performance Evaluation Cond

Performance on shared memory platform

# Blocking Vs Non-blocking algorithm

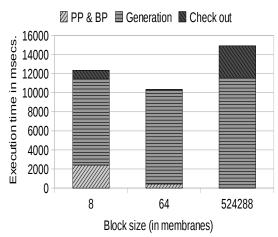
Performance increases with both: Problem size and number of processes

13 variables → 15 variables ▼ 17 variables ★ 19 variables
12 variables → 23 variables ★ 25 variables

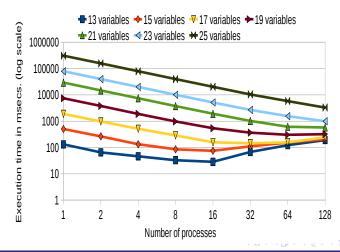




# Effects of the blocking size



# Execution time depending on the problem size



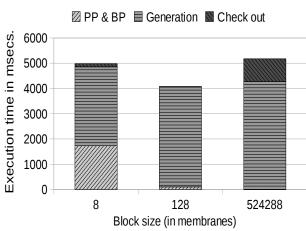
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# Blocking Vs non-Blocking

- The blocking technique reaches up to 2x versus non-blocking alternative (25 variables)
- Memory banks are independent on this platform.
- Blocking algorithm ONLY takes advantage of data locality to improve memory bandwidth.

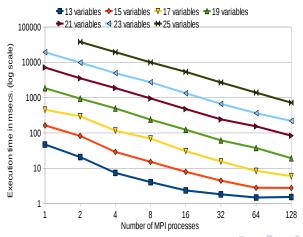


# Effects of the blocking size



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# Execution time depending on the problem size





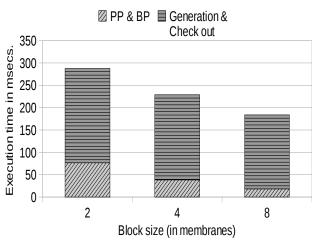
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# Blocking Vs non-Blocking

► The tiling alternative obtains up to 1.75x speed up versus non-tiling version.



# Effects of the blocking size



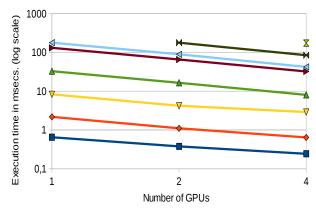
Performance Evaluation

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Performance on a GPU-based cluster

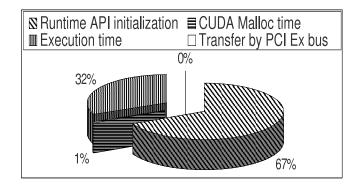
# Execution time depending on the problem size

13 variables → 15 variables
 17 variables → 19 variables
 21 variables
 22 variables
 23 variables
 24 variables





## GPUs associated overheads

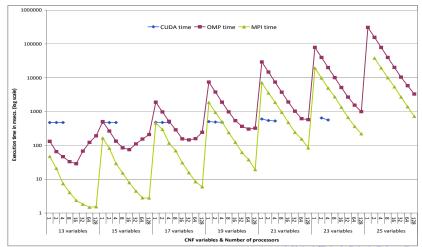


Performance Evaluation

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Performance on a GPU-based cluster

# Overall comparison





# Outline

#### Conclusions



#### Conclusions

- Simulation of a recognizer P systems with active membranes for the SAT.
- Three parallel architectures: Shared Memory, Distributed Memory and GPUs.
- Blocking increases the bandwidth in all targeted systems (data locality).
- OpenMP sim is the lowest performance, increasing overheads along with the number of processors.
- OpenMP sim is the only one able to execute all our benchmarks due to the memory requirements.
- MPI sim exhibits good scalability with the number of processors.
- GPU sim is the best platform in terms of execution time, reaching up to 10x speed up versus MPI and 40x speed up versus OpenMP.

#### Future work

- Newest generation of GPUs, such as Fermi, increase performance and flexibility.
- Cloud computing and Heterogeneous computing increase memory space without sacrifying performance.
- Other P system models for modeling ecosystems can be simulated on HPC solution.



# Thanks for your attention Questions?



- J. M. Čecilia, J. M. Ġarcía, G. D. Guerrero, M. A. M. del Amor, I. Pérez-Hurtado and M. J. Pérez-Jiménez. Simulating a P system based efficient solution to SAT by using GPUs. *Journal of Logic and Algebraic Programming*, 79(6):317–325, 2010.
- J. M. Cecilia, J. M. García, G. D. Guerrero, M. A. M. del Amor, I. Pérez-Hurtado and M. J. Pérez-Jiménez. Simulation of P systems with active membranes on CUDA. *Briefings in Bioinformatics*, 11(3):313–322, 2010.
- S. A. Cook. The complexity of theorem-proving procedures. In STOC '71: Proceedings of the third annual ACM symposium on Theory of computing, pages 151–158, New York, NY, USA, 1971. ACM.
- M. J. Pérez-Jiménez, Á. Romero-Jiménez and F. Sancho-Caparrini. Complexity classes in models of cellular computing with membranes. *J. Natural Computing*, 2(3):265–285, 2003.



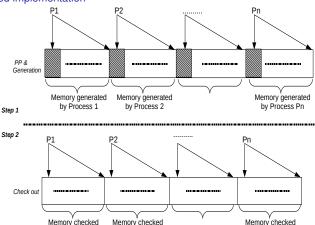
# Outline

**Appendix** 



# General data policy description

#### Non-block based implementation



by Process 2

by Process 1

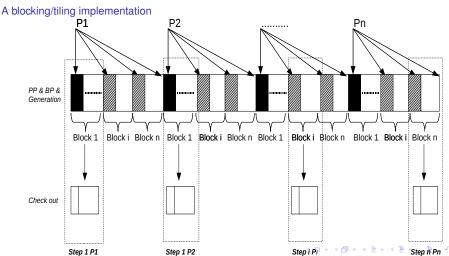
by Process n

# General data policy description

Reduced bandwidth

- Many cache misses (first writes and then reads).
- Block-based data layout to improve locality.
- Intel Vtune profiler to see memory behaviour, and Cuda Visual profiler for GPUs.
- Needs efficent handling of the exponential workspace.







V. Nguyen, D. Kearney and G. Gioiosa. An extensible, maintainable and elegant approach to hardware source code generation in reconfig-p. *J. Logic and Algebraic Programming*, 79(6):383–396, 2010.



G. Paun. Membrane computing. An introduction. *Springer-Verlag*, pages 9–419, 2002.

