# Distributed Memory Implementation Strategies for the kinetic Monte Carlo Algorithm



António Esteves Centro ALGORITMI

iPC

Alfredo Moura Institute of Polymers and Composites



University of Minho| Braga, PortugalSchool of Engineering

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- Objectives
- Al<sub>3</sub>Sc precipitation
- kinetic Monte Carlo method kMC
- synchronous parallel kinetic Monte Carlo method spkMC
- Parallelization of spkMC with MPI
- Results
- Conclusions

## **Objectives**

- Parallelize kMC with spkMC algorithm to speedup its execution
- Use distributed memory architecture and MPI
- Explore different computation vs. communication strategies
- Evaluate spkMC results by comparing them with kMC:
  - number of precipitates
  - dimension of precipitates
  - > precipitates normalized by lattice sites, etc.
- Compare and assess spkMC implementations performance and scalability







- Precipitation of Al<sub>3</sub>Sc in Aluminum is the formation of clusters of atoms with an Al<sub>3</sub>Sc structure
- **Precipitates** alter significantly the Al properties
- Precipitates have a Face-Centered Cubic crystalline structure
- <u>Sc atoms</u> on the vertices and <u>Al atoms</u> on the faces
- Atoms **move** in the lattice structure by means of:
  - > vacancy diffusion: jump to a neighbor vacant site
  - interstitial diffusion







• Used to model the **temporal evolution** of a system by stochastically exploring sequences of transitions

• Calculates the **transitions rates** for all trial **configurations**  $\rightarrow \Gamma_{i,j}$ 

• Selects a new configuration j with a probability proportional to  $\Gamma_{i,j}$ 



•  $\Gamma_{i,V}$  is called vacancy exchange frequency

$$\Gamma_{i,V} = v_i * e^{-\frac{\Delta E_{i,V}}{k_B T}}$$

•  $v_i \equiv$  attempt frequency for an *Al/Sc* atom

•  $\Delta E_{i,V} \equiv$  activation energy required to move an *Al/Sc* atom into a vacancy



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• Moving an *Al* atom through **vacancy diffusion** 



#### Selecting a move

- > A vacancy is surrounded by 12 first nearest neighbors
- > Calculate 12 jump frequencies  $\rightarrow \Gamma_1 \dots \Gamma_{12}$
- ➢ Generate a random number between 0 and 1
- $\triangleright$  Select the **n**-th jump frequency that verifies the relation:

$$\sum_{i=1}^{n} \Gamma_{i} \leq \text{random number} \leq \sum_{i=1}^{n+1} \Gamma_{i}$$



Perform a spatial decomposition into subdomains

• Obtain the **accumulated frequency** for each subdomain  $\rightarrow$ 

• Define the maximum frequency  $\rightarrow$ 

$$\Gamma_{\max} \geq \max_{k=1,\dots,K} \left\{ \Gamma_k \right\}$$

 $\Gamma_k = \sum_{i=1}^{k} \Gamma_{ik}$ 

• Assign a null event frequency to the subdomains  $\rightarrow \gamma_{0k} = \Gamma_{\max} - \Gamma_k$ 

■ Define the spkMC **time step** increment →





# spkMC implementation - spatial domain decomposition





- At simulation start, a **vacancy** is placed on every **sector**
- Vacancies are allowed to **migrate** out of their original sector
- Sprint is a sequence of MCS, performed on a sector, without communication
- At end of sprint, each process communicates boundary moves to its neighbors
- **Boundary region** is as large as possible and we keep track of the changes that occurred on it during the sprints
- Avoid conflicts with a **checker board** scheme

- MPI point-to-point communication:
  - Both processes participate actively
  - Complications:
    - when a process has multiples messages to be received
    - when the strong synchronization associated with blocking communication is unsuitable
    - possibility of deadlock
  - > <u>Alternatives</u>:
    - MPI nonblocking point-to-point pattern
    - MPI-2/3 one-sided communication (or RMA)

- \* 🗘
- Allows remote memory access (RMA) to a region called window
- Access epoch: RMA synchronization call on the window → 1+ RMA communication calls → RMA synchronization call
- Advantage: asynchronous, or at least, less synchronous
- Transfer routines: MPI\_Put, MPI\_Get, MPI\_Accumulate
- Synchronization mechanisms:
  - Fence, post-start-complete-wait, lock-unlock

※ 🗘

- Fence synchronization:
  - > It is collective over the entire communicator associated with the window
  - > It may result in communication overhead.
- Post-start-complete-wait (PSCW) synchronization:
  - Restricts synchronization to the minimum
  - Programmer selects the groups of processes that synchronize.
- Lock-unlock synchronization:
  - ➤ The origin process calls MPI\_Win\_lock to access the target window → calls transfer routines → calls MPI Win unlock
  - Emulates a shared memory model.

• PSCW and lock-unlock prototypes:

#### ≻ Use RMA

- ➤ A trimmed list of boundary and ghost moves is communicated to the adequate neighbor processes at the end of the sprints
- Lock-unlock proved to be 3x faster than PSCW
- ➢ Performance was not satisfactory → code profiling proved that a significant percentage of the execution time was spent in MPI barriers

※ 🗘

- Send-receive prototype:
  - Uses point-to-point communication
  - The communication pattern is simpler, regular and similar to the one used by SPPARKS/LAMMPS simulators
  - The communication runs in 3 steps: send and receive moves to/from nearest neighbor in +X (or -X) direction, in +Y (or -Y) direction, and in +Z (or -Z) direction
  - Due to checker board scheme we do not have to send and receive from both '+' and '-' directions in each step
  - > Send (or receive) the variable **number** of moves and the **moves**
  - ➢ Initiate a non-blocking receive (MPI\_Irecv) → do a blocking send
    (MPI\_Send) → wait for receiving to complete (MPI\_Wait).

# **spkMC** implementation - prototypes

- optimized send-receive prototype:
  - The tasks done during each MCS were optimized, mainly to simplify the analysis of the vacancy moves
  - > The data structures were simplified.



# spkMC implementation – opt-send-receive algorithm

Read simulation, lattice, energy, and parallelization parameters if (this is master process) then Send and extended subdomain to all processes Receive the extended subdomain from master process Compute the 1<sup>st</sup> and 2<sup>nd</sup> nearest neighbors for all subdomain for (each *sprint* of the simulation) do for (each sector in subdomain) do for (each MCS of a sprint) do kMC core for (each *vacancy* in current sector) do Calculate the activation energy associated with the 12 1<sup>st</sup> nearest neighbors of the vacancy Calculate vacancy exchange frequency and real time for this MCS Select randomly a 1<sup>st</sup> neighbor for new position of the vacancy Swap the vacancy with the selected neighbor Store the vacancy move in the array *moves* endFor endFor Eliminate false moves, convert coordinates, and generate movesX/Y/ZSend and receive *movesX* to/from the neighbor process in X direction Send and receive *movesY* to/from the neighbor process in Y direction Send and receive *movesZ* to/from the neighbor process in Z direction endFor

if (this sprint is a snapshot point) then

Master process gathers subdomains from all processes and writes configuration to file endFor



- The simulations were run on the University of Minho SeARCH cluster
- Cluster nodes run Linux x86\_64
- The code was compiled with gcc 4.9.0 and Open MPI 1.8.4
- Hardware configuration of each **node**:
  - ➢ 2 processors/sockets
  - Processors: Intel Xeon E5-2650 v2, with ivy bridge microarchitecture, 2.6 GHz, 8 physical cores, and 16 cores with hyper-threading
  - ➢ 64GB of RAM
  - ➢ 20MB of L3 cache
  - > 256KB of L2 cache per core
  - > 32KB of L1D and 32KB of L1I cache per core.
- Inter-node **communication**: Ethernet and Myrinet



T=673 K, 1% Sc



Mean radius (Å)



• Mean size (atoms)



Number of precipitates



Precipitates/Lattice sites







T=673 K, 1% Sc



Mean radius (Å)



Mean size (atoms)



Number of precipitates



Precipitates/Lattice sites







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 $T=673 \ K$ , 1% Sc



**Metrics**:



Mean size (atoms)



Number of precipitates



Precipitates/Lattice sites







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#### T=673 K, 1% Sc



**Metrics**:



Mean size (atoms)

+ -

Number of precipitates



Precipitates/Lattice sites

+ -













#### output from DBSCAN clustering









output from DBSCAN clustering









#### output from DBSCAN clustering









#### output from DBSCAN clustering









output from DBSCAN clustering





output from spkMC simulation



output from DBSCAN clustering









#### output from DBSCAN clustering









output from DBSCAN clustering









output from DBSCAN clustering









#### output from DBSCAN clustering





# **Performance of the different spkMC prototypes**





• Speedup of the best spkMC prototype in relation to sequential kMC is 4



spkMC presents a low parallel efficiency



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- spkMC reproduces accurately the statistical behavior of the sequential kMC
- The precipitation problem is not embarrassingly parallel → spkMC only presents a 4x speedup when compared to kMC
- Open MPI 1.8.4 does not support RMA natively → RMA did not disclosed its potential in the lock-unlock prototype

### **Future Work**



- Improve the parallel simulation performance and scalability
  - Prevent the migration of vacancies between sectors
    - eliminates the **iterations complexity** associated with multiple vacancies
    - improves the **load balancing** between processes
  - Overlap communication with computation
  - Use a hybrid MPI-OpenMP implementation to improve intra-node computation performance and still allow more parallelism than a single node
  - > Take advantage of the improved **RMA** support allowed by **Open MPI 2.0.0**