

Molecular Dynamics - Assignment 3

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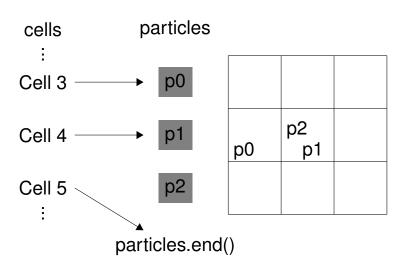




The Cell Data-Structure - Approach 1

Idea:

- Sort Particles in accordance to their Cell Position
- save which part of the particles-Vector corresponds to which cell

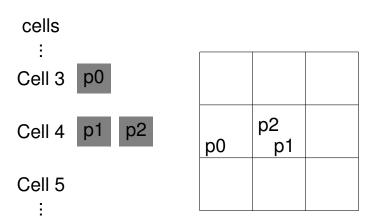




The Cell Data Structure - Approach 2

Idea:

Approach 1.1 stored multiple virtual vectors in one vector \rightarrow let's actually store the particles in vectors corresponding to their cell

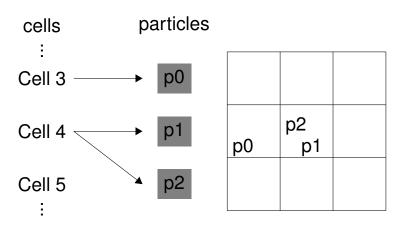




The Cell Data Structure - Approach 3

Idea:

- Each Cell only keeps references to their members
- No sorting or copying of entire particles required





Approach Comparison

Approach 1	Approach 2	Approach 3
+ Easy to implement	+ Easy to implement	+ Easy to implement
+ Interface for old Assignments remains unchanged	+ New Implementation of some methods needed	 Interface for old Assignments remains unchanged
 Expensive struct swaps during sorting 	 Expensive struct copies with potential reallocs needed 	+ References are cheap
 Direct access to particles for calculations 	+ Direct access to particles for calculations	+ Dereferencing needed



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In the end we decided to implement approach 3. ,



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input Loader gets chosen at compile time





Input parsing - Definition of Body

```
enum Shape {cuboid, sphere};

struct Body {
        Shape shape;
        Eigen::Vector3d fixpoint;
        Eigen::Vector3d dimensions;
        double distance;
        double mass;
        Eigen::Vector3d start_velocity;
} ;
```





• Protection of master branch





- Protection of master branch
- Deployment of CI/CD pipeline for all branches



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The pipeline consist of:

library installation



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- build process



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The pipeline consist of:

- library installation
- build process
- sanitizers
- unit tests for every major component







Logging

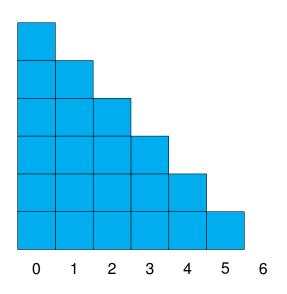
format:

[time][level::context] message

- 6 different log-levels available
- log-level can be chosen via command line input
- use of log functions since we don't log in performance critical areas
- no need to deactivate logging at compile time



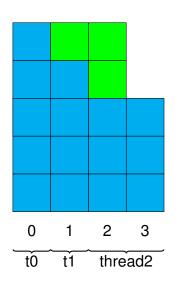
Performance optimization - "Gaussian multithreading"



- Idea: Force calculation can be multithreaded quite easily
- Evenly distribute Particle-pairs among multiple threads
- One rectangle represents one necessary force-calculation where min(p1,p2) is the number displayed below



Performance optimization - "Gaussian multithreading"



- Create "Gaussian rectangle" as good as possible and redistribute the resulting blocks
- Threads use personal accumulators
- Accumulators get summed up in the end

In the end outsourcing the problem to OpenMP turned out to be much easier