

Molecular Dynamics - Assignment 3

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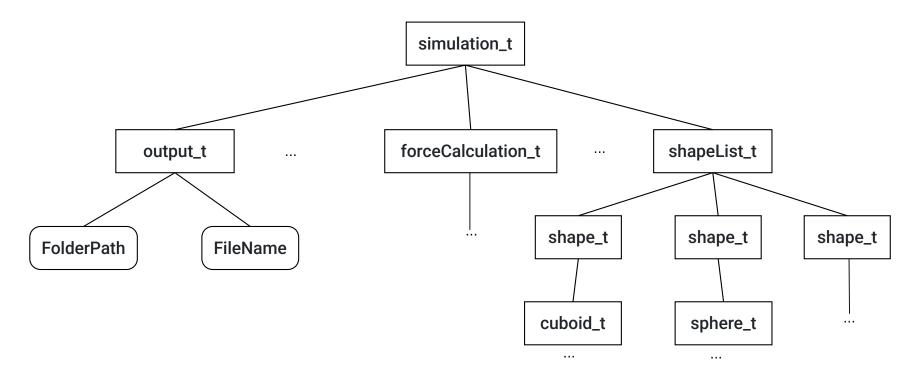
Lehrstuhl für wissenschaftliches Rechnen

9. Dezember 2022





XSD- tree-like definition of Datastructures







XSD- Code Snippet

- simulation_t stores all the necessary parameters
- utilizes defined other tree-like Datastructures
- "simulation_t is root"





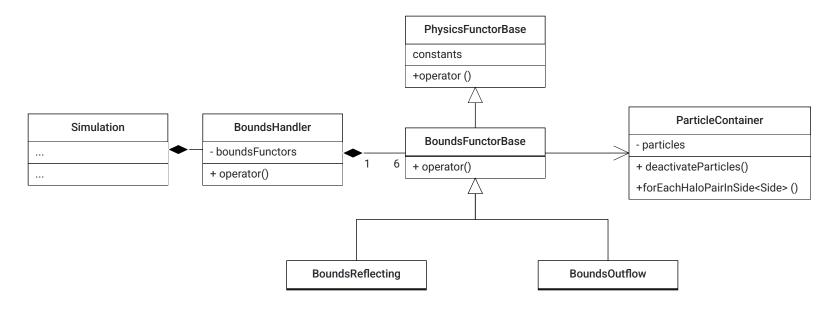
XSD- Code Snippet

Trade-off convenience \leftrightarrow complexity in action:





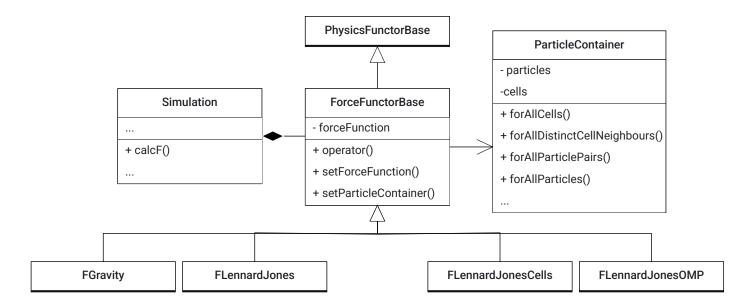
Bounds Handling





ForceFunctors

- Force function used gets determined on runtime via input parameters
- Force functor defines the algorithm (Linked-Cell algorithm/ All-Pairs algorithm) used





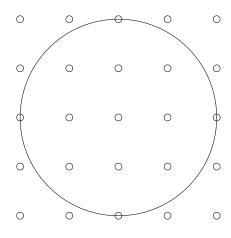


Spheres

Expansion of the Body-struct utilized in Assignment 2

$$\sqrt{x^2 + y^2 + z^2} <= r$$

$$\iff x^2 + y^2 + z^z <= r^2$$

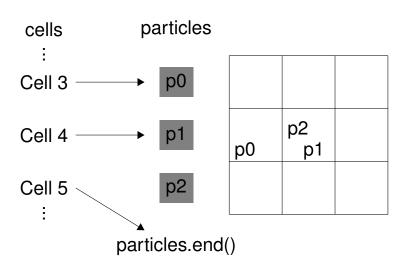




The Cell Data-Structure - Approach 1

Idea:

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

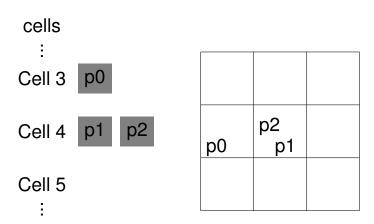




The Cell Data Structure - Approach 2

Idea:

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

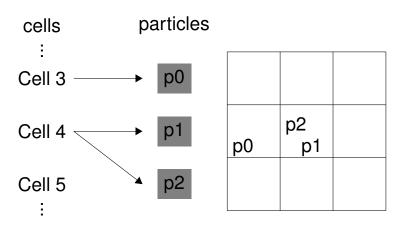




The Cell Data Structure - Approach 3

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.



ParticleContainer's new methods

- Functionality of first two methods is sufficient but hard to optimize
- Functionality of last two methods results in higher cohesion, but potential for runtime improvement



• Particles initiated in a square with varying dimensions



- Particles initiated in a square with varying dimensions
- Space between square and domain border kept at 10



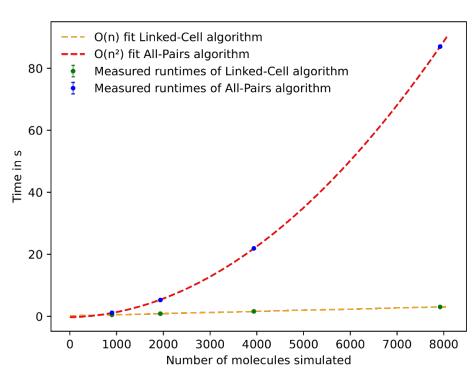
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- Space between square and domain border kept at 10
- *r_{cutoff}* =const. (aswell as other variables like eps, sig, brown, etc.)



- Particles initiated in a square with varying dimensions
- Space between square and domain border kept at 10
- r_{cutoff} =const. (aswell as other variables like eps, sig, brown, etc.)
- exact commands to recreate the results are in README and at the end of this presentation



Runtime Comparison of different algorithms



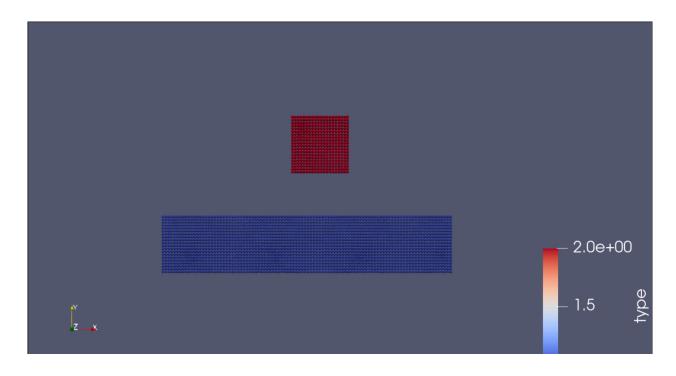
Hardware details:

Ubunutu 20.04 LTS i7-12700KF @5,0GHz 64GB RAM @ 3200MHz





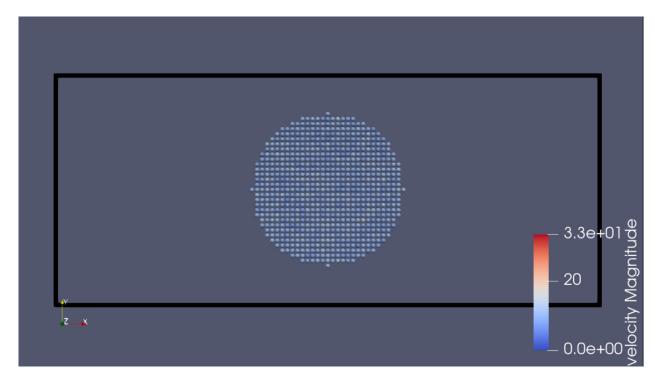
Collision of two bodies







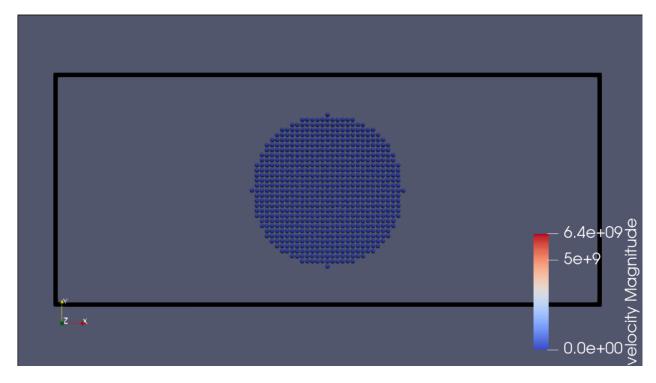
Our Simulation







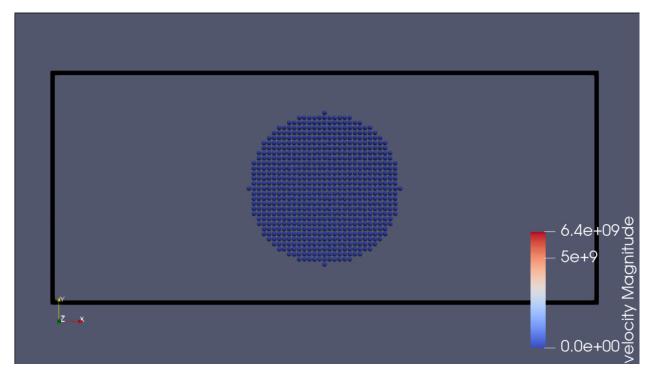
Increasing Δt







Further increasing Δt





Roadblocks and lessons learned

- Play the objective!
- Getting together and figuring out an elegant solution for one hour is an hour well spent
- Clearly define Interfaces





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Recreating benchmarks

Setup and compilation:

- 1. mkdir build output
- 2. cd build
- 3. cmake ...
- 4. make

Example of bechmark commands:

Linked-Cell algorithm:

./MolSim ../input/square1.txt -dt 0.0005 -et 0.5 -lc 1 -bbox0 50.0 -bbox1 50.0 -f lennardjonescell -rc 3.0 -bench file -i 10 > ../output/lc_square1.txt

All-Pairs algorithm:

./MolSim ../input/square1.txt -dt 0.0005 -et 0.5 -lc 0 -bbox0 50.0 -bbox1 50.0 -f lennardjones -rc 3.0 -bench file -i 10 > ../output/ap_square1.txt



Recreating benchmarks

- Results can be found in output-folder
- New benchmark files can be written and executed easily
- for other testsizes change input file, output file and bbox-size accordingly

bbox-sizes needed: (50,50), (64,64), (83,83), (109,109)