

### Molecular Dynamics - Assignment 3

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**TUM CIT** 

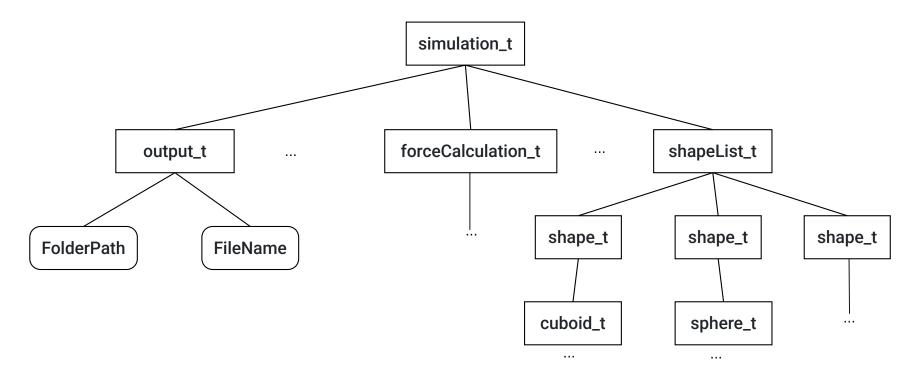
Lehrstuhl für wissenschaftliches Rechnen

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#### 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"



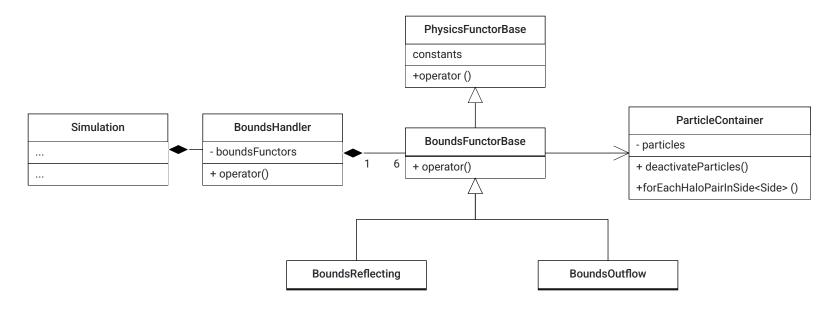


## **XSD- Code Snippet**





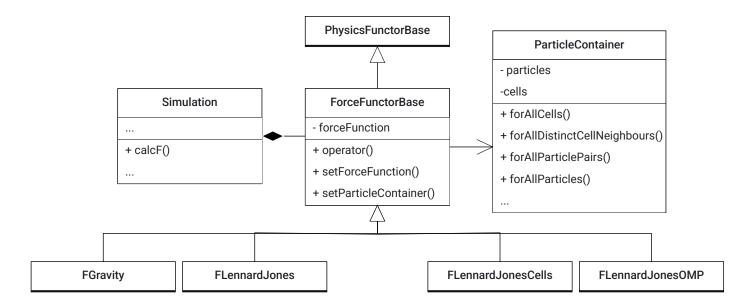
# **Bounds Handling**





#### **ForceFunctors**

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





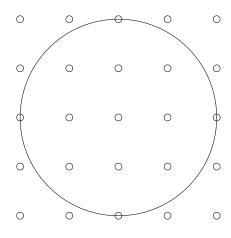


# **Spheres**

Expansion of the Body-struct utilized in Assignment 2

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

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

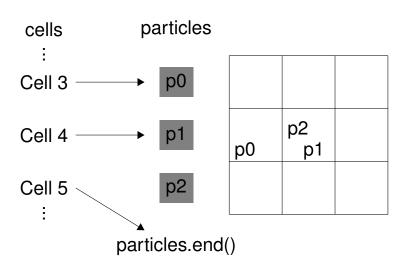




# The Cell Data-Structure - Approach 1

#### Idea:

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

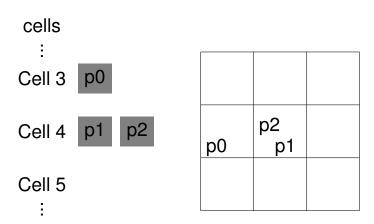




## The Cell Data Structure - Approach 2

Idea:

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

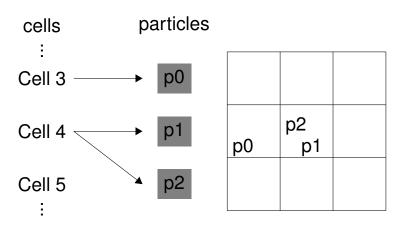




## The Cell Data Structure - Approach 3

#### 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
<ul><li>+ Interface for old Assignments remains unchanged</li></ul>	+ New Implementation of some methods needed	<ul> <li>Interface for old Assignments remains unchanged</li> </ul>
<ul> <li>Expensive struct swaps during sorting</li> </ul>	<ul> <li>Expensive struct copies with potential reallocs needed</li> </ul>	+ References are cheap
<ul> <li>Direct access to particles for calculations</li> </ul>	+ 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



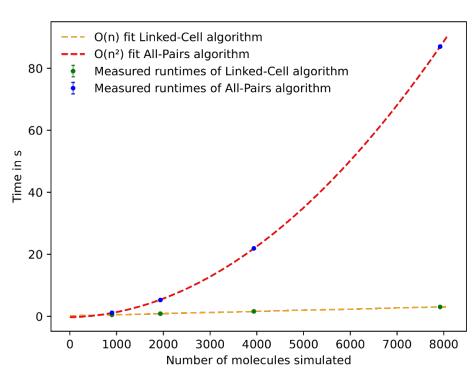
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- *r<sub>cutoff</sub>* =const. (aswell as other variables like eps, sig, brown, etc.)



- 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



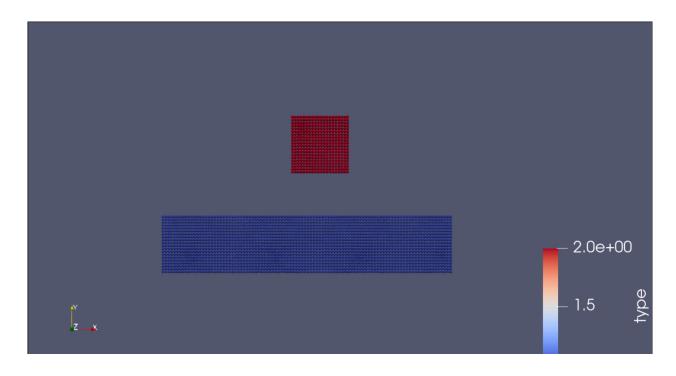
Hardware details:

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





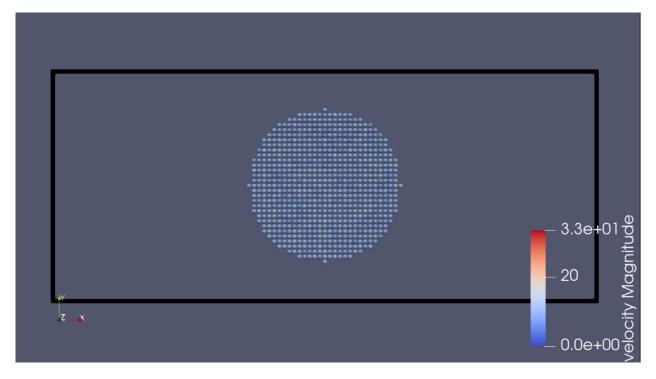
### Collision of two bodies







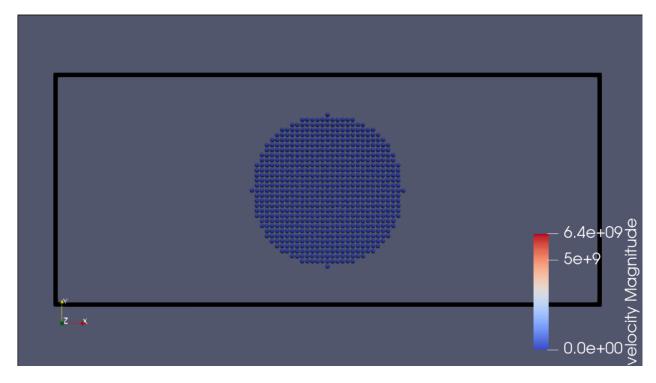
### Our Simulation







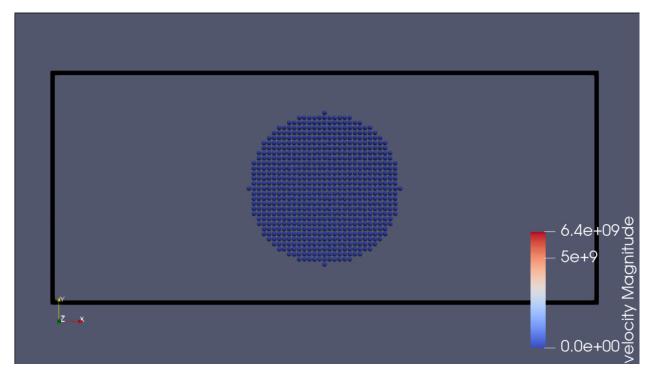
# Increasing $\Delta t$







# Further increasing $\Delta t$





#### 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)