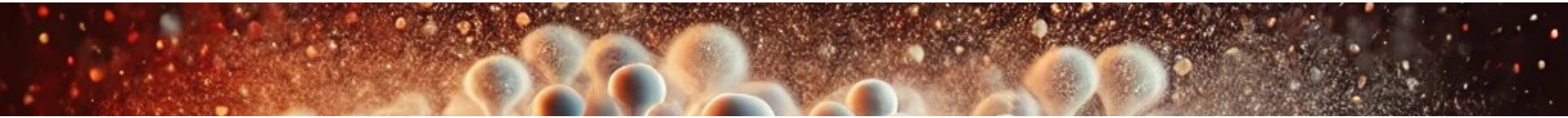
PSE Molekulardynamik
Sheet 4: Thermostats, Rayleigh-Taylor instability and “falling drops”



Group A: Daniel Schade, Ashutosh Solanki, Robin Cleve

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Thermostats

- Calculate temperature of the system for all particles:

$$T = \frac{\sum_i m_i |\mathbf{v}_i|^2}{\#\text{dimensions} \times \#\text{particles}}$$

- Rescale velocity for all particles:

$$\mathbf{v}_i \leftarrow b \cdot \mathbf{v}_i, \quad \text{where} \quad b := \sqrt{\frac{T_{\text{new}}}{T_{\text{current}}}}$$

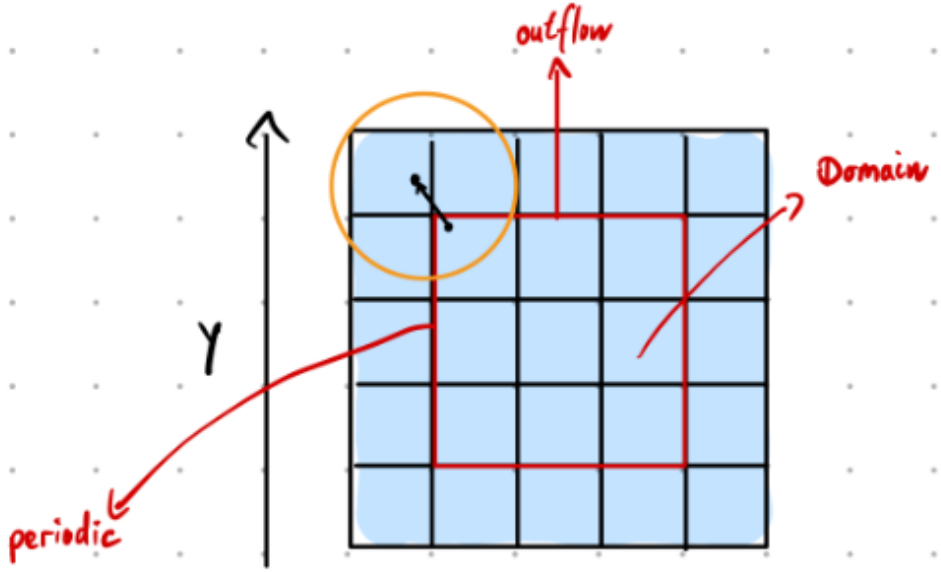
- Apply gradually, limit temperature change of the system by a certain value
- **Temperature of System with with static particles is not defined**

  ➡ Initialize system with Brownian motion!

Rayleigh-Taylor instability

Periodic boundaries

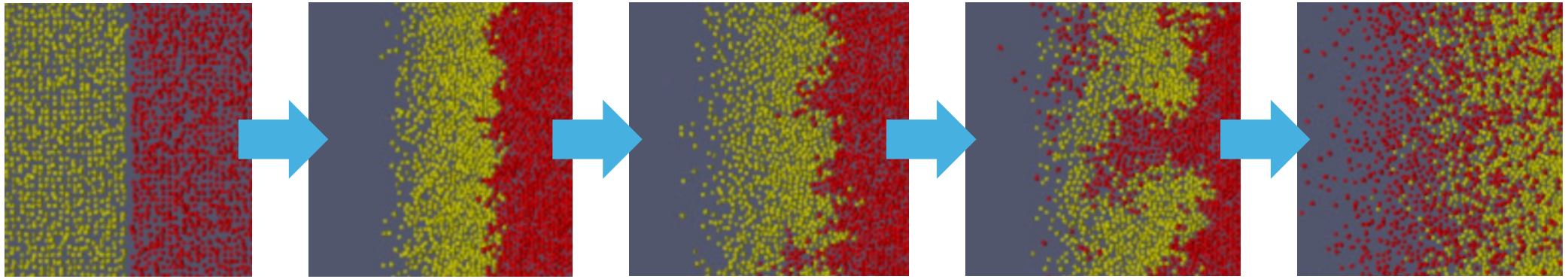
- **Main challenges:**
  1. Teleporting of particles that left the domain
     -> “easier part” of the implementation
  2. Calculate forces for particles close to the border of the domain
     -> Interpret cell of the opposite side as halo cell
- **General approach:**
  - split domain up in Sides, Edges, Corners
- **Edge case:**
  - Particle in the boundary cell corner
  - Which boundary did it cross
  - For now it depends on the processing order



Rayleigh-Taylor instability

Further steps

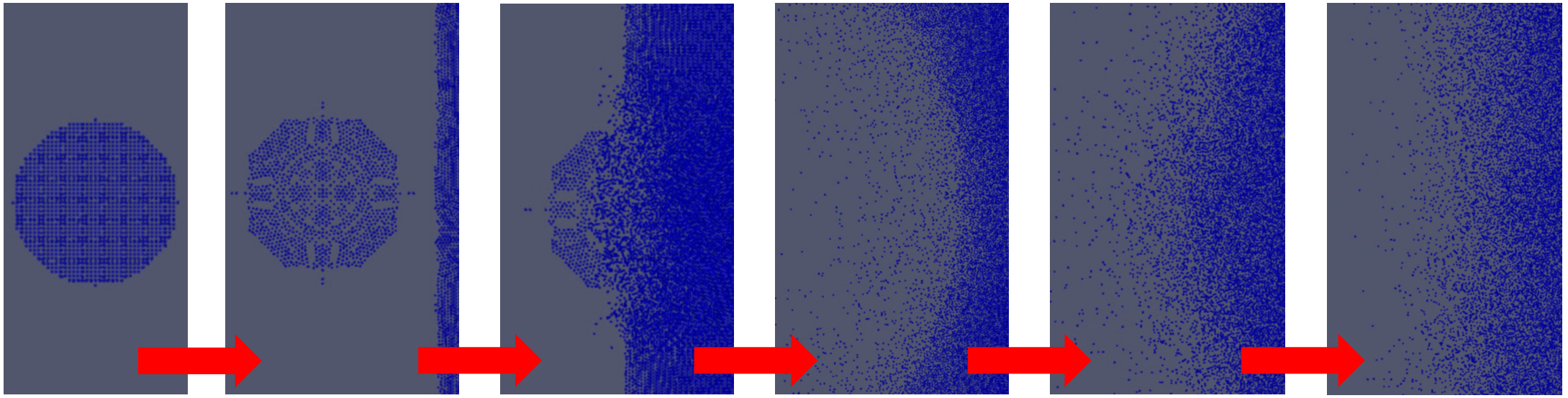
1. Gravity
   - Add force along the y-axis to each particle
2. Lorentz-Berthelot Mixing Rules
   - Epsilon and Delta can now be configured in each particle individually
   - Adapt Leonard Jones force
3. Simulation



Simulation of a falling drop - Liquid

Prepare liquid:

→ “checkpoint 1”



Performance Measurement and Profiling

- These methods consume the most runtime, 80.06% of total execution time

Method	s	
LeonardJonesForce::compute	2.68	39.59
LinkedCellsContainer::applyToAllUniquePairsInDomain	1.47	21.71
std::_Function_handler<void (Particle&, Particle&), Model::updateForces()>	0.77	11.37
Particle::getX()	0.50	7.39

- Observations:
  - Iterating over cells of the linked cells container consume a lot of time, as well as the
  - getter method for the position of molecule

- Linux cluster - Fist 100 iterations of the Rayleigh-Taylor instability of task 2

Running time	19.16 s
Molecule updates per second	522030 MUPS/s

Tuning the sequential performance

- Leonard-Jones force between two particles
  - Calculate difference in distance of the particles only once
  - **Speed up by 1.2s!**
- Reduce getter calls
  - establishing friend classes and adding multiple compiler flags (e.g. -Ofast)
  - **Speed up by 2.04s!**
- Conclusion
  - Only constant speed up, time complexity of program stays the same
  - More complex changes, like caching Leonard-Jones parameters, may be implemented in the following weeks

Tuning the sequential performance

Graph after second optimization step