

01

Scientific Computing (PSE) Molecular Dynamics  
Group D

# Worksheet 5



# SIMULATION TASKS

Task 1 to 3 -Worksheet 5

## Simulation of a membrane

Storing neighboring particles and avoiding self-penetration to simulate a flexible membrane

## Parallelization

Using the OpenMP framework to allow multithreading and parallel computation

## Rayleigh-Taylor instability in 3D

Fluid cuboid with higher density above fluid cuboid with lower density subjected to gravitational forces in three dimensional space

## Parallelization & Rayleigh Taylor 3D

Parallel processing at the beginning to be able to have better performance (at least in some cases) in further tasks and simulations

## TIMELINE

### Adaptation of Thermostat

### Implementation of Nanotubes

Nanotubes with different fixed walls can be created with XML input. Extended thermostat to ignore total velocity of the fluid

We decided to do the membrane at the end and the "enhancing", so feature addition or extension as well as performance enhancing through parallelization, at the beginning to use and test it in the different scenarios after that.

### Membrane

Kept the membrane as last task as it can work independently and seemed to be more complex to implement to us

04

# NANOSCALE FLOW SIMULATION

Task 4 - Option A

## Fixed walls

Position-fixed cuboids that represent walls but still interact (with Lennard-Jones forces) with the fluid inside the tube

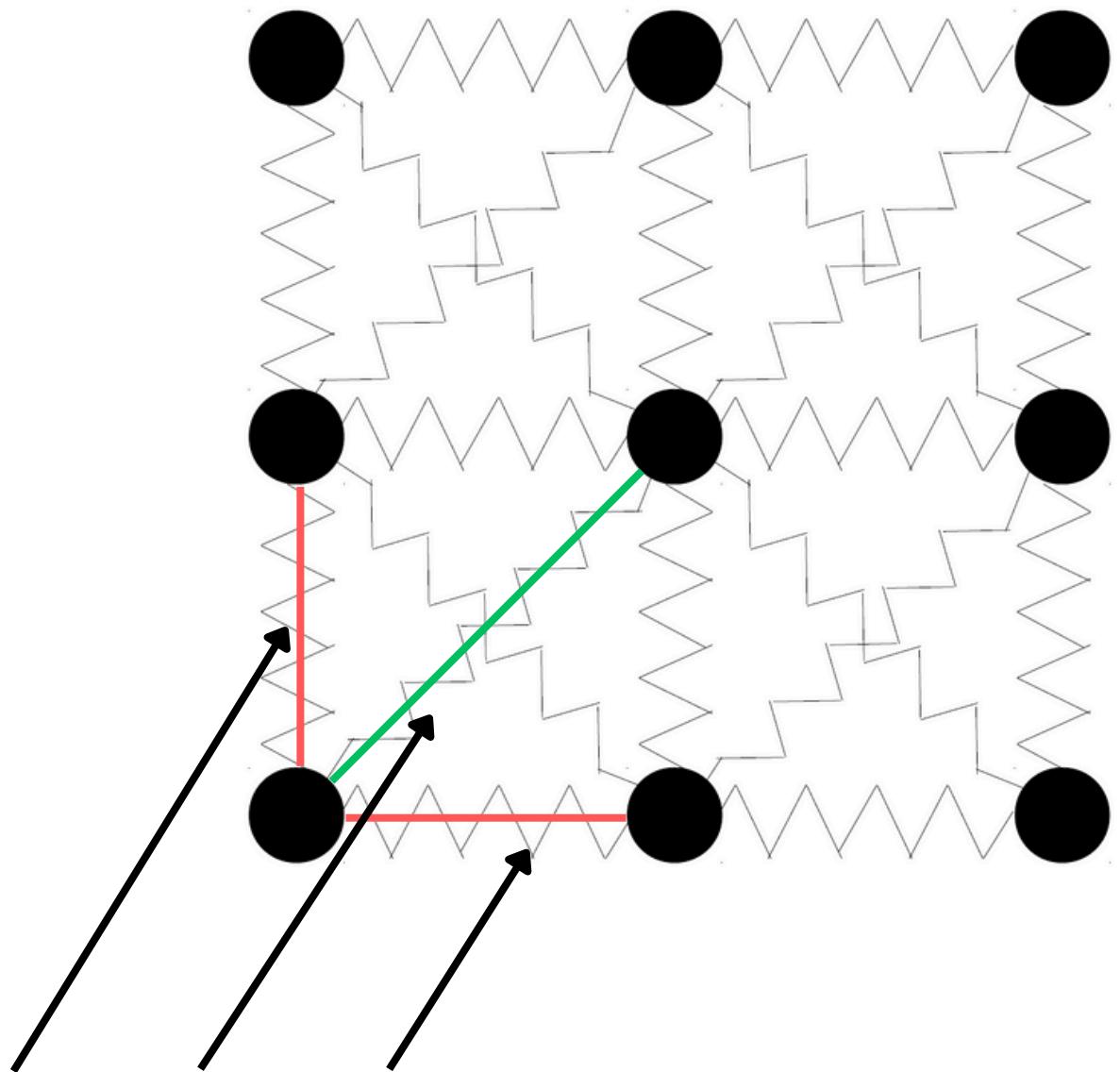
## Adapted thermostat

Extension of the existing thermostat from last sheet, ignoring the total velocity of the fluid

## Density & Velocity-Profile

Output of .csv files that give information about the density and velocity of molecules in each bin

# Particle Class



Extended the Particle class to store neighbor molecules

Initialized on a rectangular grid, setting **direct** and  
**diagonal** neighbors

# MEMBRANE

## Features

Neighboring molecules with different properties than just two molecules next to each other

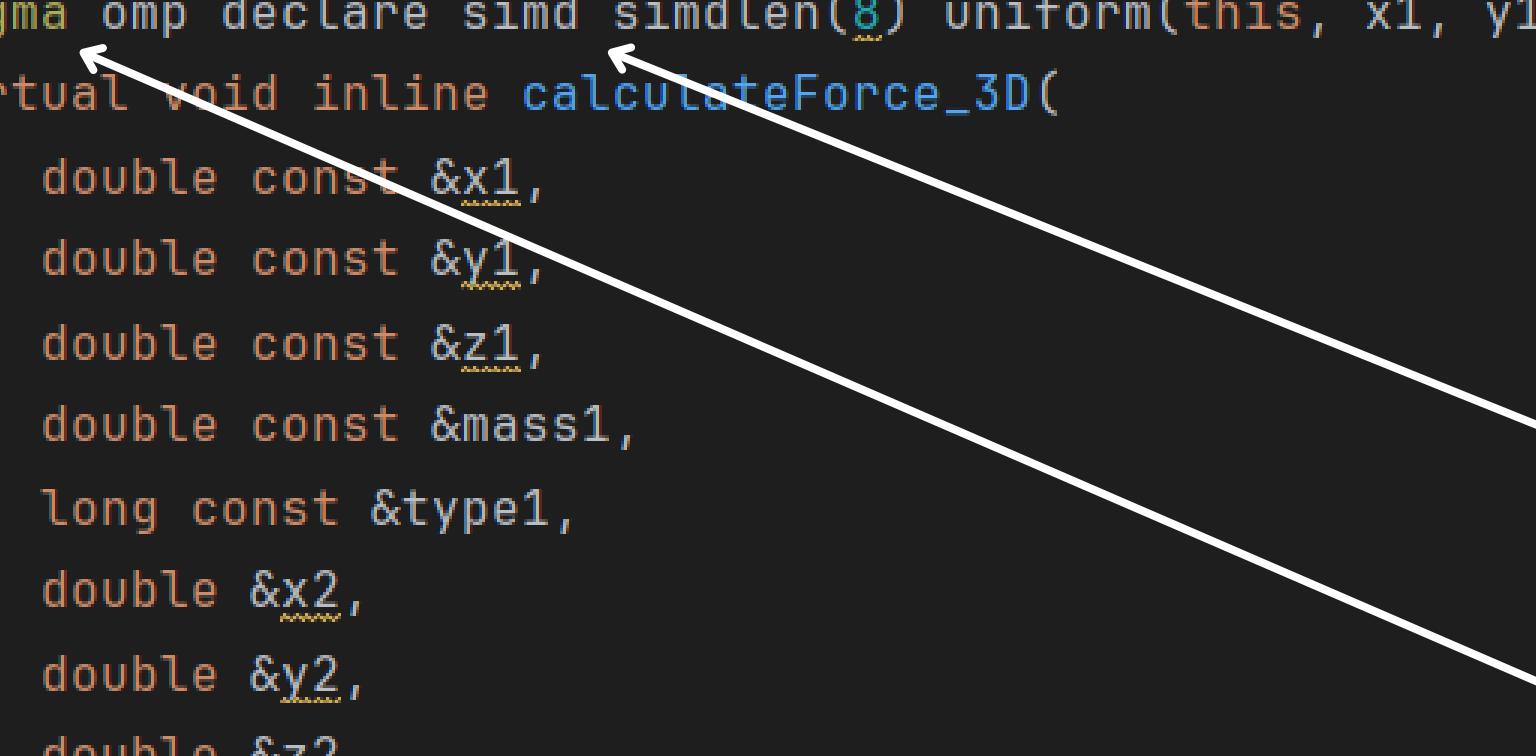
Avoidance of self penetration by repulsive Lennard-Jones

Neighboring particles interact in a harmonic potential (new force)

# OpenMP

Parallelization

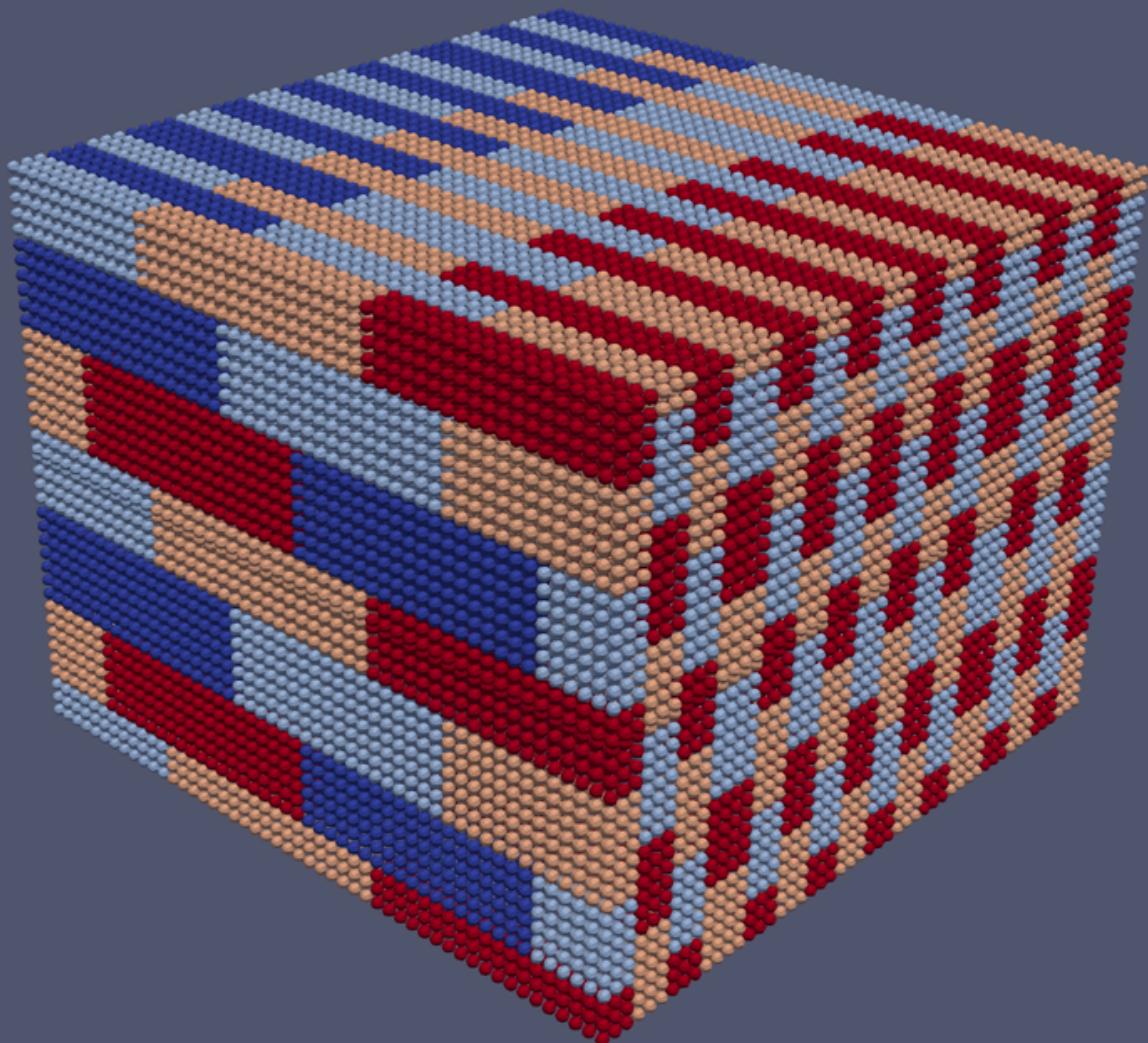
```
#pragma omp declare simd simdlen(4) uniform(this, x1, y1, z1, mass1, type1, correction) linear(ref(x2, y2, z2, mass2, type2))
#pragma omp declare simd simdlen(8) uniform(this, x1, y1, z1, mass1, type1, correction) linear(ref(x2, y2, z2, mass2, type2))
virtual void inline calculateForce_3D(
    double const &x1,
    double const &y1,
    double const &z1,
    double const &mass1,
    long const &type1,
    double &x2,
    double &y2,
    double &z2,
    double &mass2,
    long &type2,
    double &result_x,
    double &result_y,
    double &result_z,
    std::array<double, 3> &correction) = 0;
```



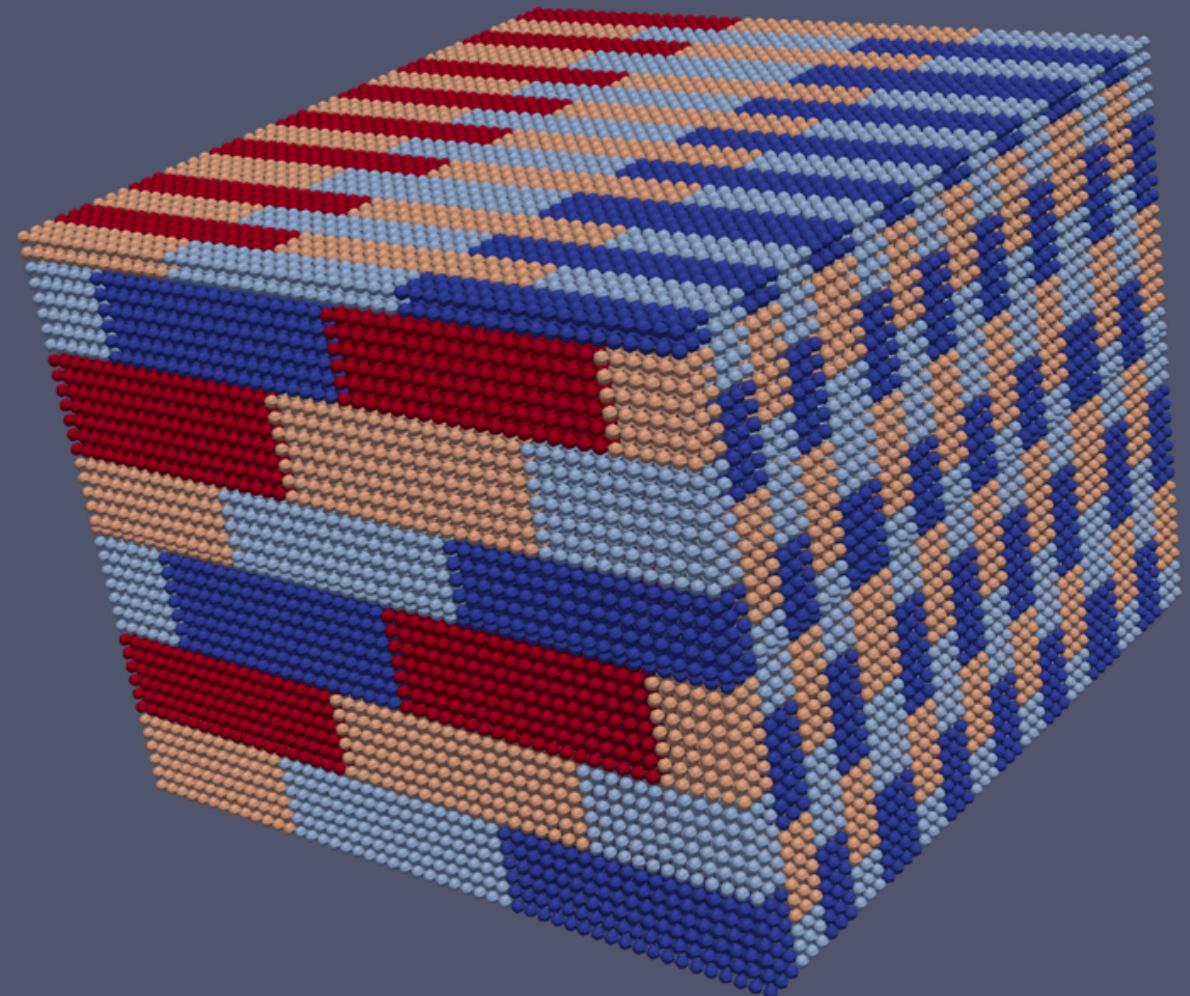
simd  
declarations of #pragma omp

# Domain Coloring

Parallelization



Rayleigh-Taylor  
3D cuboid  
by color



# TASK 3

Rayleigh-Taylor instability in 3D

```
scenario xmlns:xs="http://www.w3.org/2001/XMLSchema-instance"
  <header base_name="Rayleigh-Taylor instability"
    output_frequency="100"
    t_end="100" seed="1337"
    delta_t="0.0005"
    dimensions="3"
    vectorized="false"
    parallelized="true"
  />
  <thermostat t_init="40" frequency="1000" brownian_motion="0"/>
  <container>
    <linked_cells cutoff_radius="3.6">
      <domain_size x="60" y="60" z="60"/>
      <boundary_conditions>
        <boundary_condition type="periodic"/>
        <boundary_condition type="reflecting"/>
        <boundary_condition type="periodic"/>
        <boundary_condition type="periodic"/>
        <boundary_condition type="reflecting"/>
        <boundary_condition type="periodic"/>
      </boundary_conditions>
    </linked_cells>
    <!--      <vector/>-->
  </container>
  <forces>
    <lennard_jones>
      <gravity>-12.44</gravity>
      <particleTypes>
        <particleType id="0" sigma="1.2" epsilon="1" mass="1"/>
        <particleType id="1" sigma="1.1" epsilon="1" mass="2"/>
      </particleTypes>
      <particles>
        <cuboid spacing="1.2" particleTypeId="0">
          <coordinate x="0.6" y="0.6" z="0.6"/>
          <dimensions x="50" y="20" z="50"/>
          <velocity x="0" y="0" z="0"/>
        </cuboid>
        <cuboid spacing="1.2" particleTypeId="1">
          <coordinate x="0.6" y="24" z="0.6"/>
          <dimensions x="50" y="20" z="50"/>
          <velocity x="0" y="0" z="0"/>
        </cuboid>
      </particles>
    </lennard_jones>
  </forces>
```

## Parallelization upfront

Implemented parallelization task beforehand, so the runtimes are manageable. Local runtime of around **42 seconds** for contest file. The large experiment took about **8 hours** on a laptop with 12 cores.

## Simulation

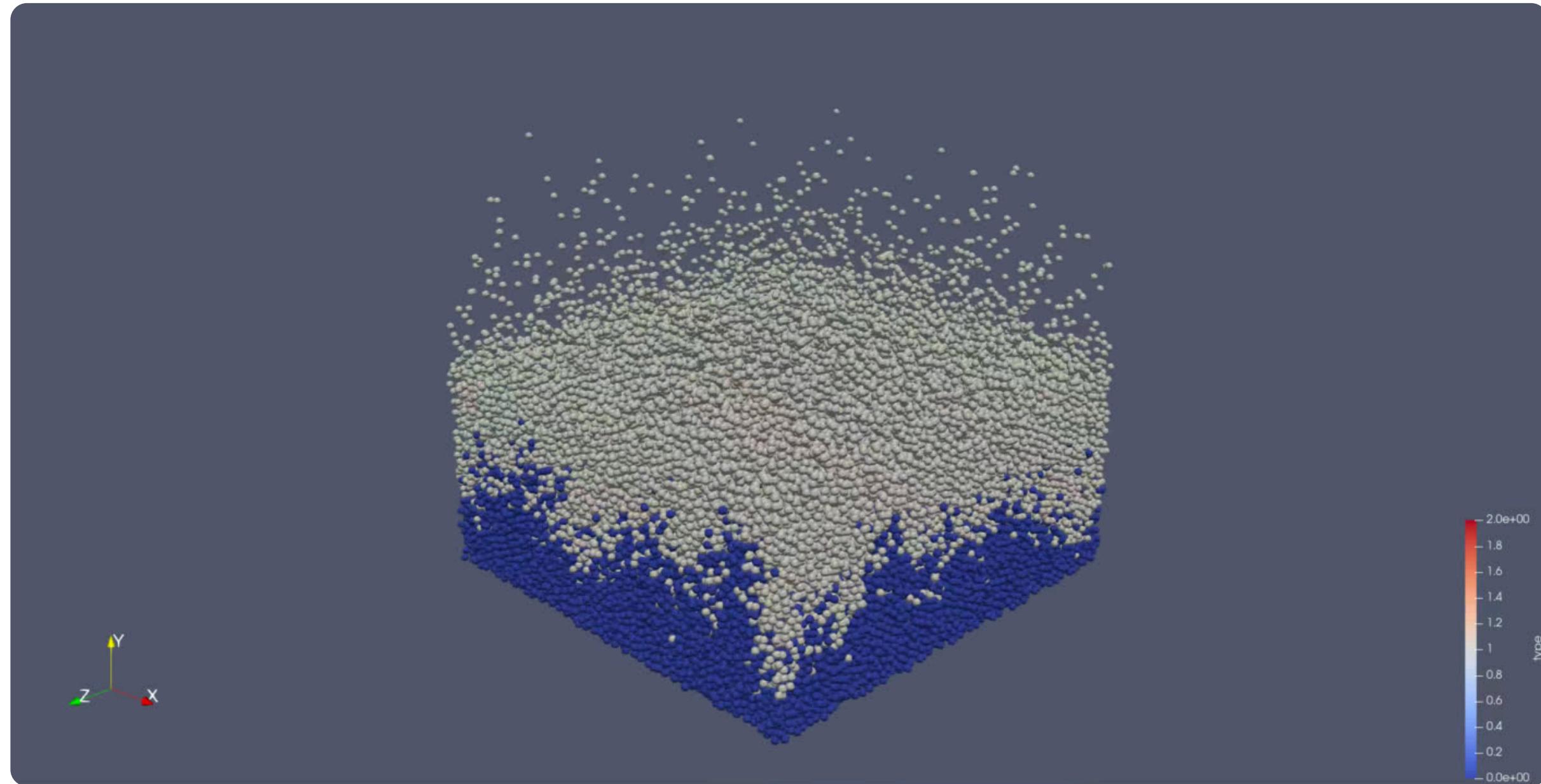
Input and force calculation was already adapted for 3D simulations. Ran simulations according to worksheet from different angles and perspectives.

Local hardware: Ryzen 9 7950X3D 16K/32T, 64GB DDR5 RAM      **08**

# SIMULATION 1

Colored by type (the denser fluid is white)

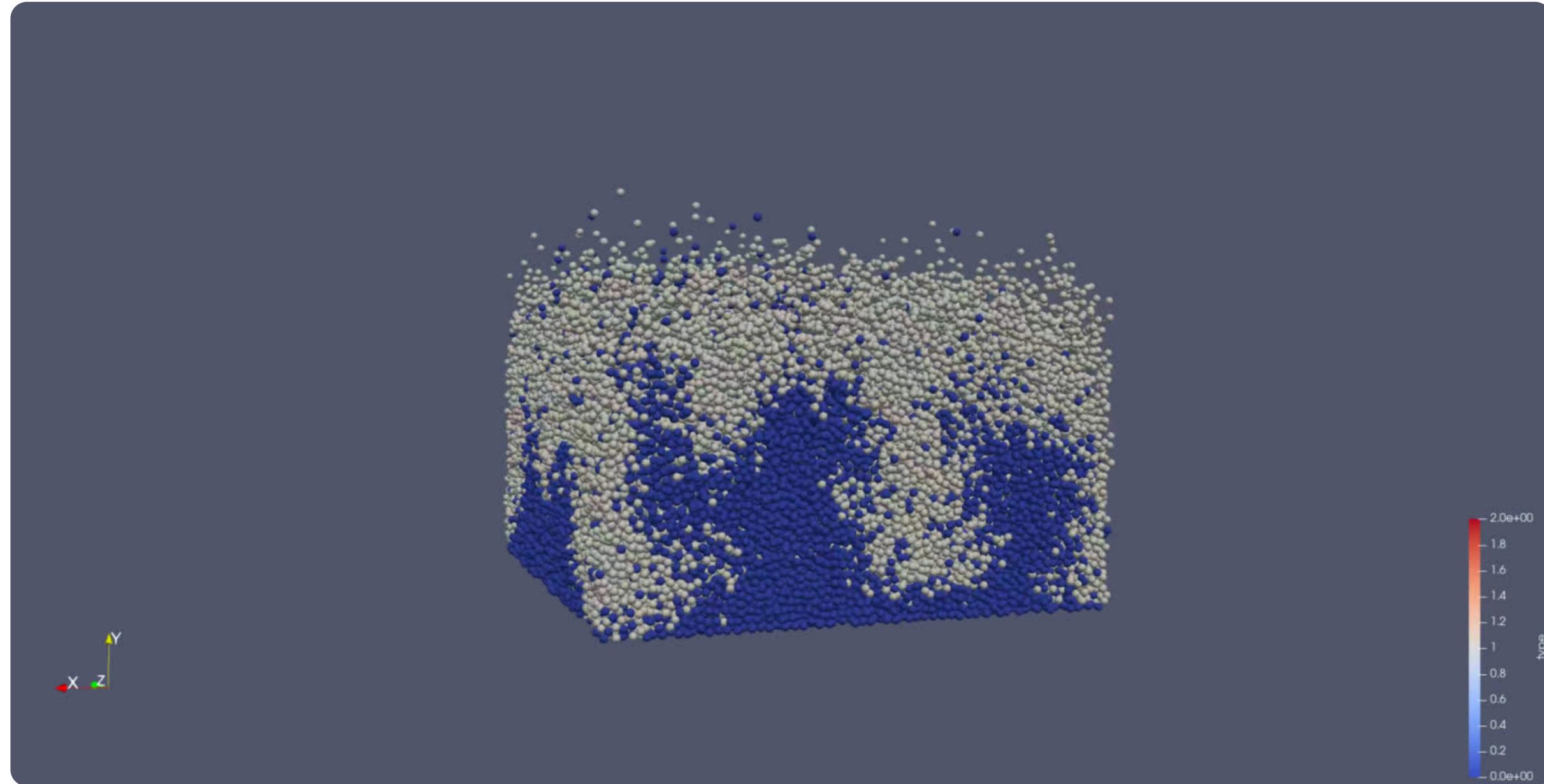
09



# SIMULATION 1

Colored by type (the denser fluid is white)

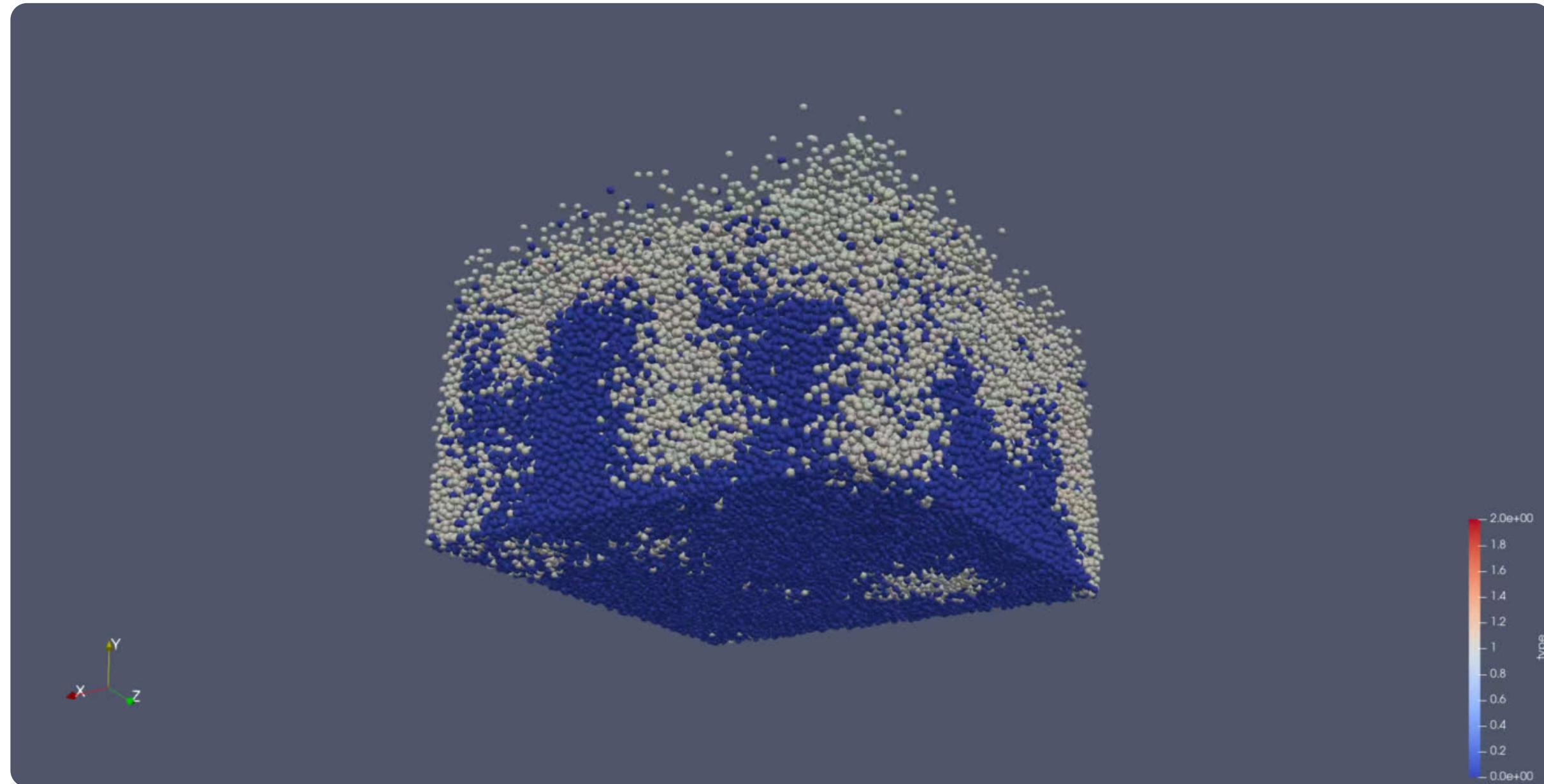
10



# SIMULATION 1

Colored by type (the denser fluid is white)

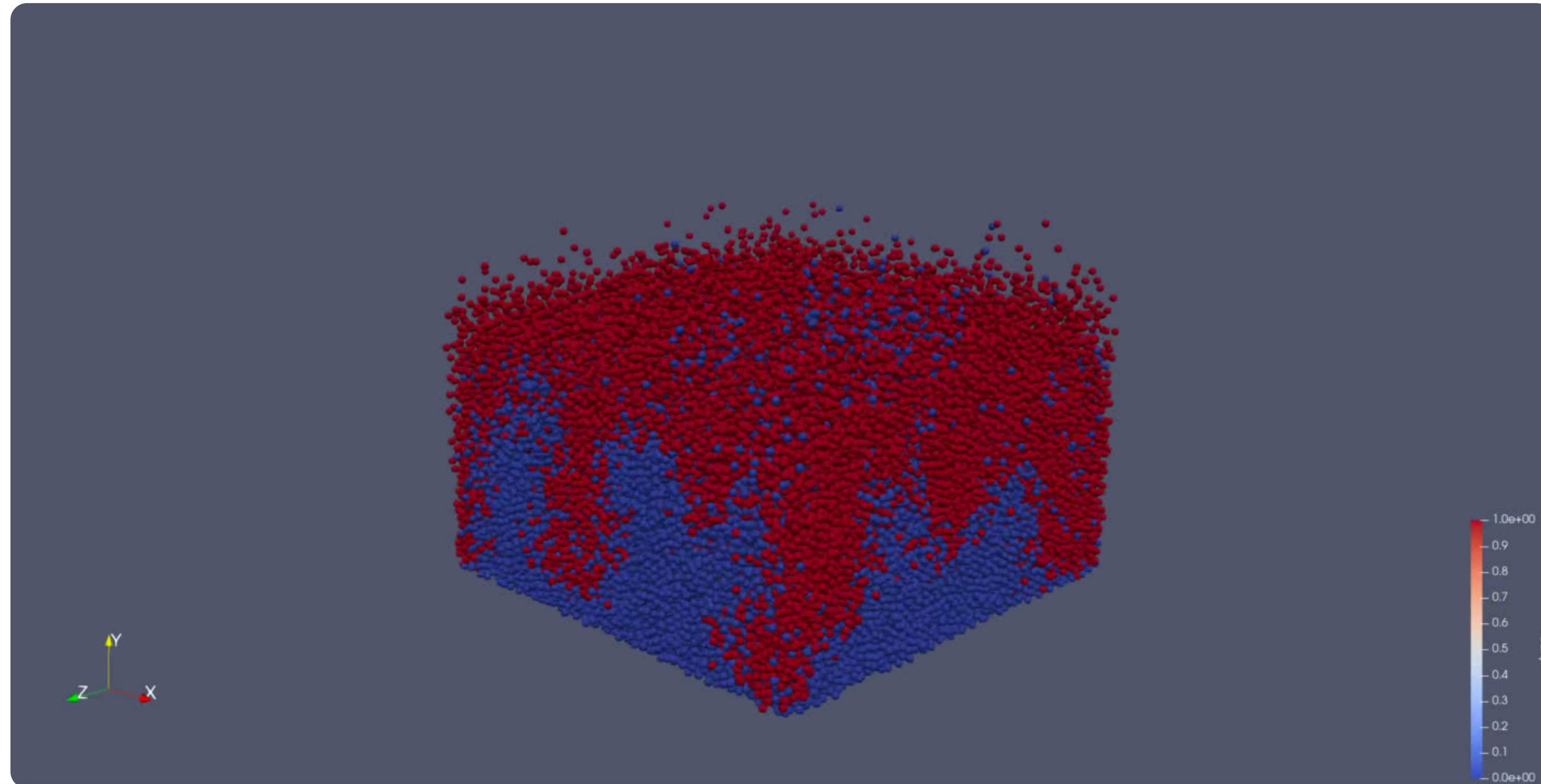
11



# SIMULATION 1

Colored by type (the denser fluid is red)

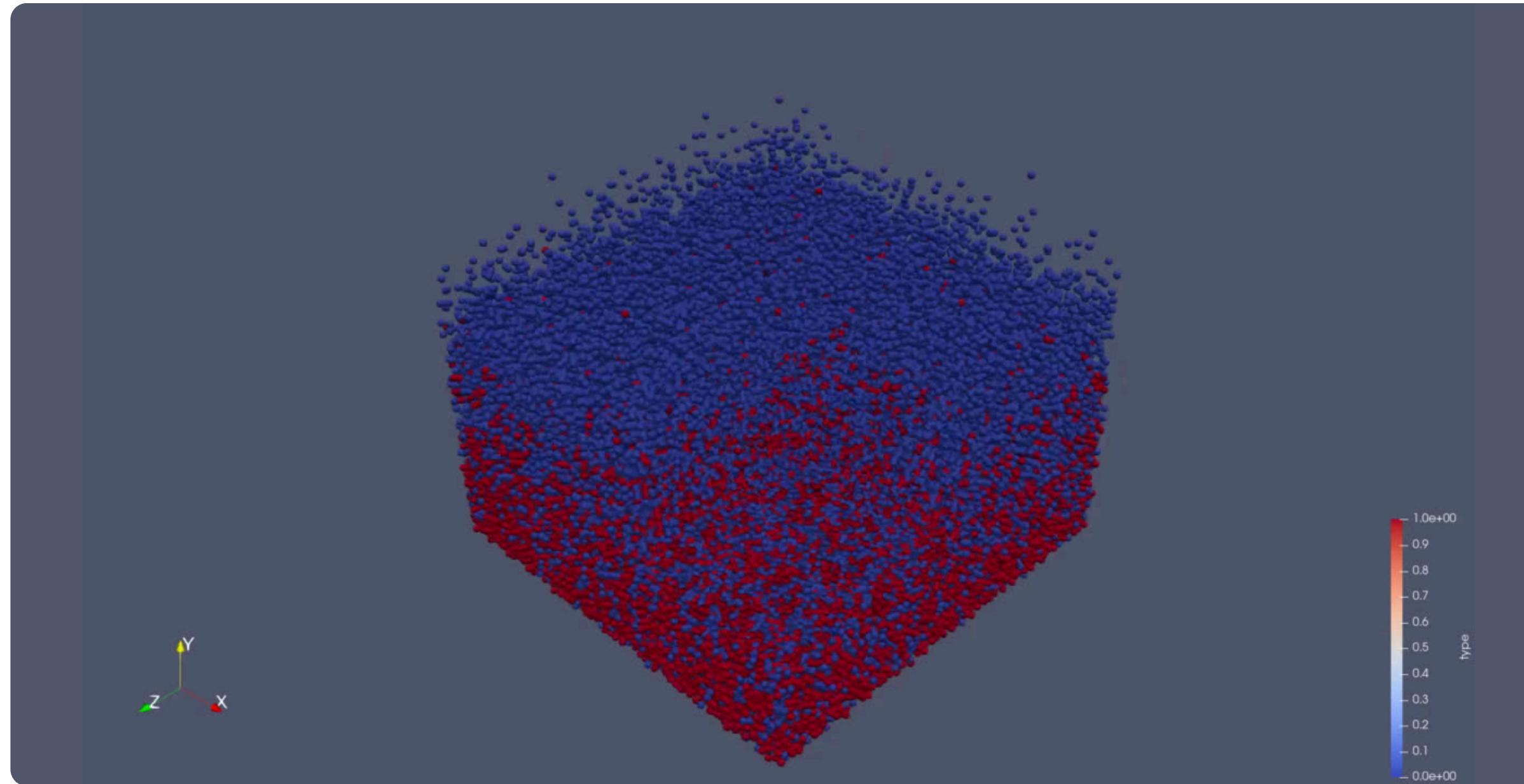
12



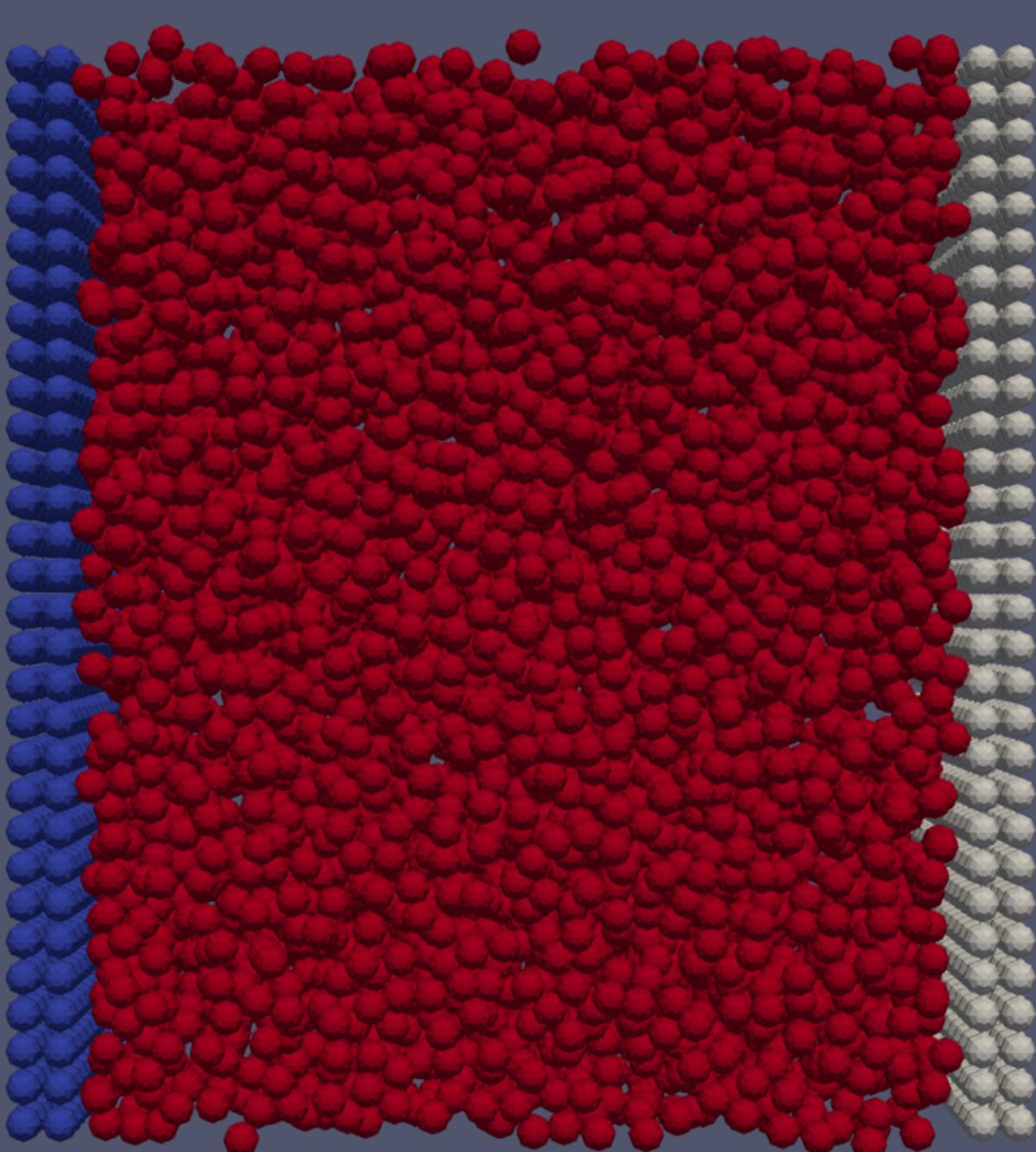
# SIMULATION 1

Colored by type (the denser fluid is red)

13



# Nanotube



## TASK 4 - OPTION A

Nano-scale flow simulation

### Fixed walls

No position updates or velocity calculations for fixed molecules, implemented through an extra optional attribute in XML that is saved in config, specifying if a molecule is fixed

Different input parameters for sigma, epsilon etc. enable different types of materials, e.g. for the walls

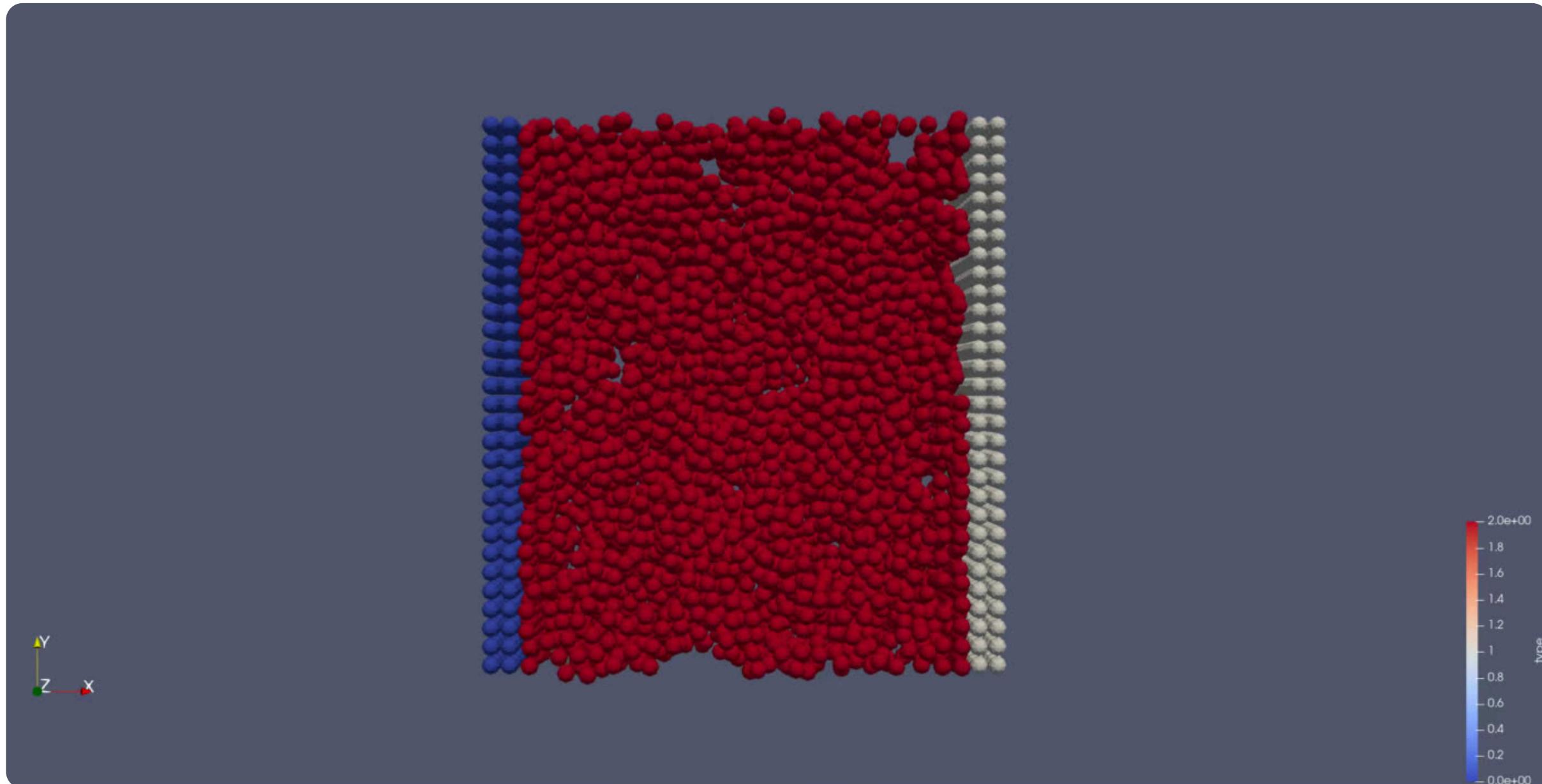
### Performance

Local runtime for specifications on worksheet with output writing enabled but with  $t_{end} = 50$  is **08:42:776 (mm:ss::ms)**

# SIMULATION 2

Colored by type (walls are blue and white, fluid is red)

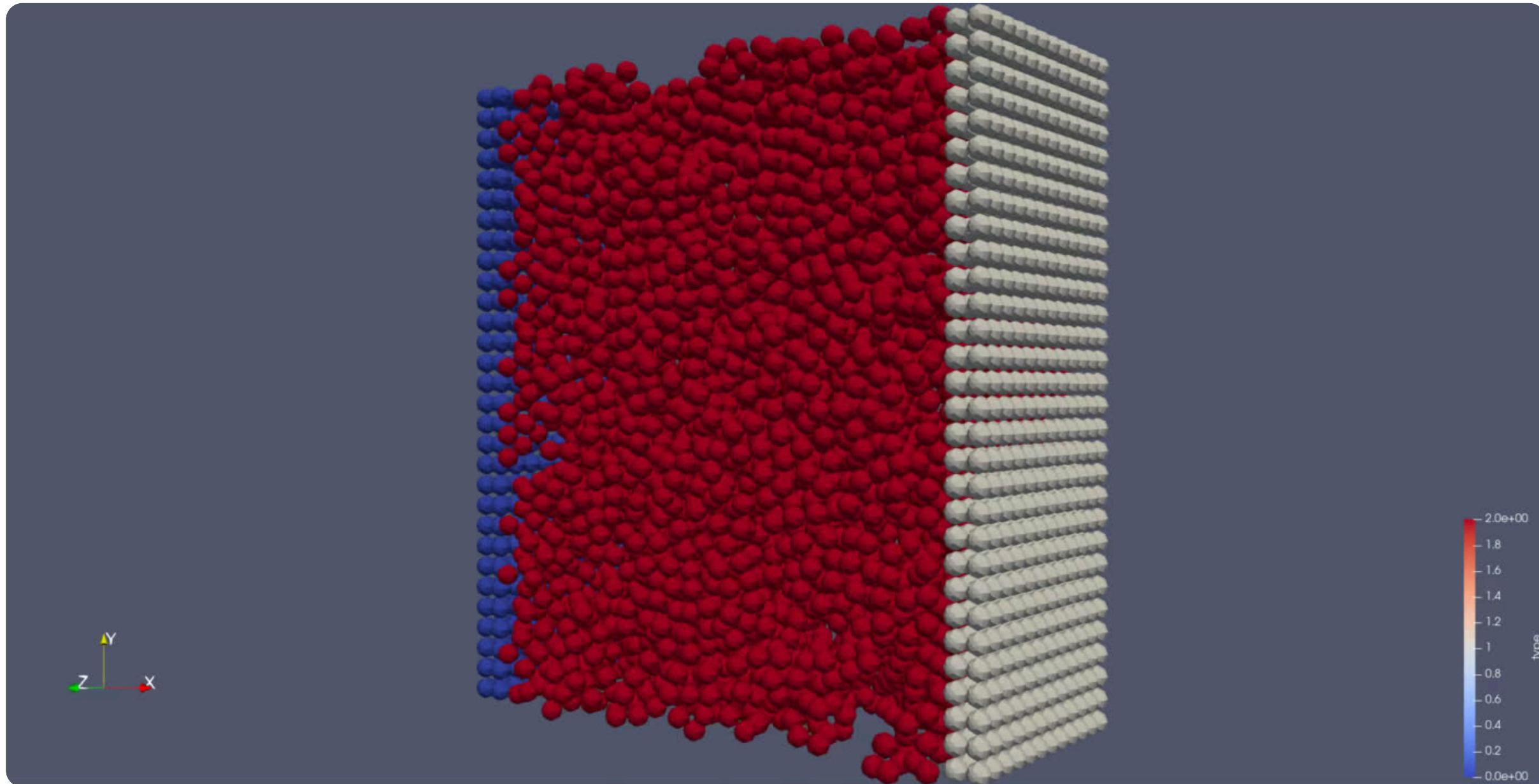
15



# SIMULATION 2

Colored by type (walls are blue and white, fluid is red)

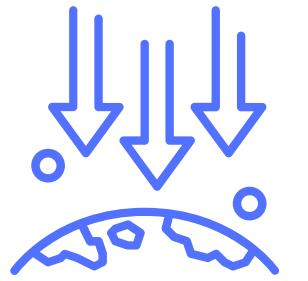
16



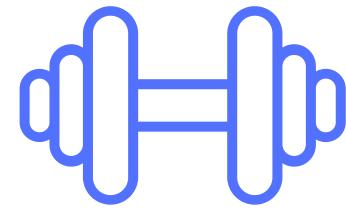
# OTHER NANOTUBE SCENARIOS

Task 4 - Option A

High gravity



Large mass walls



Large epsilon fluid

$\epsilon$

Large sigma fluid

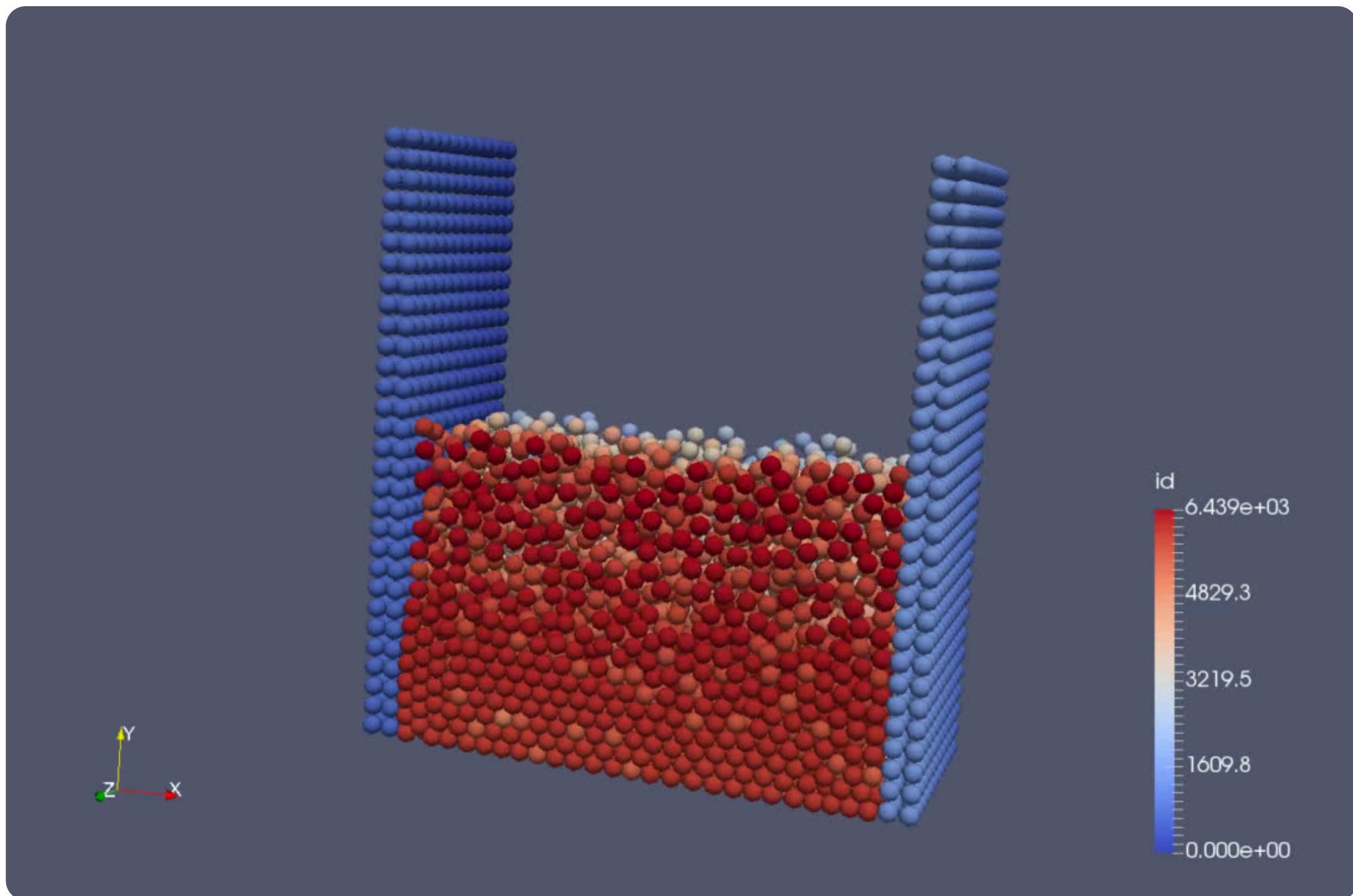
$\sigma$

# SIMULATION 3

Colored by id (walls are blue, fluid is red)

High gravity

18

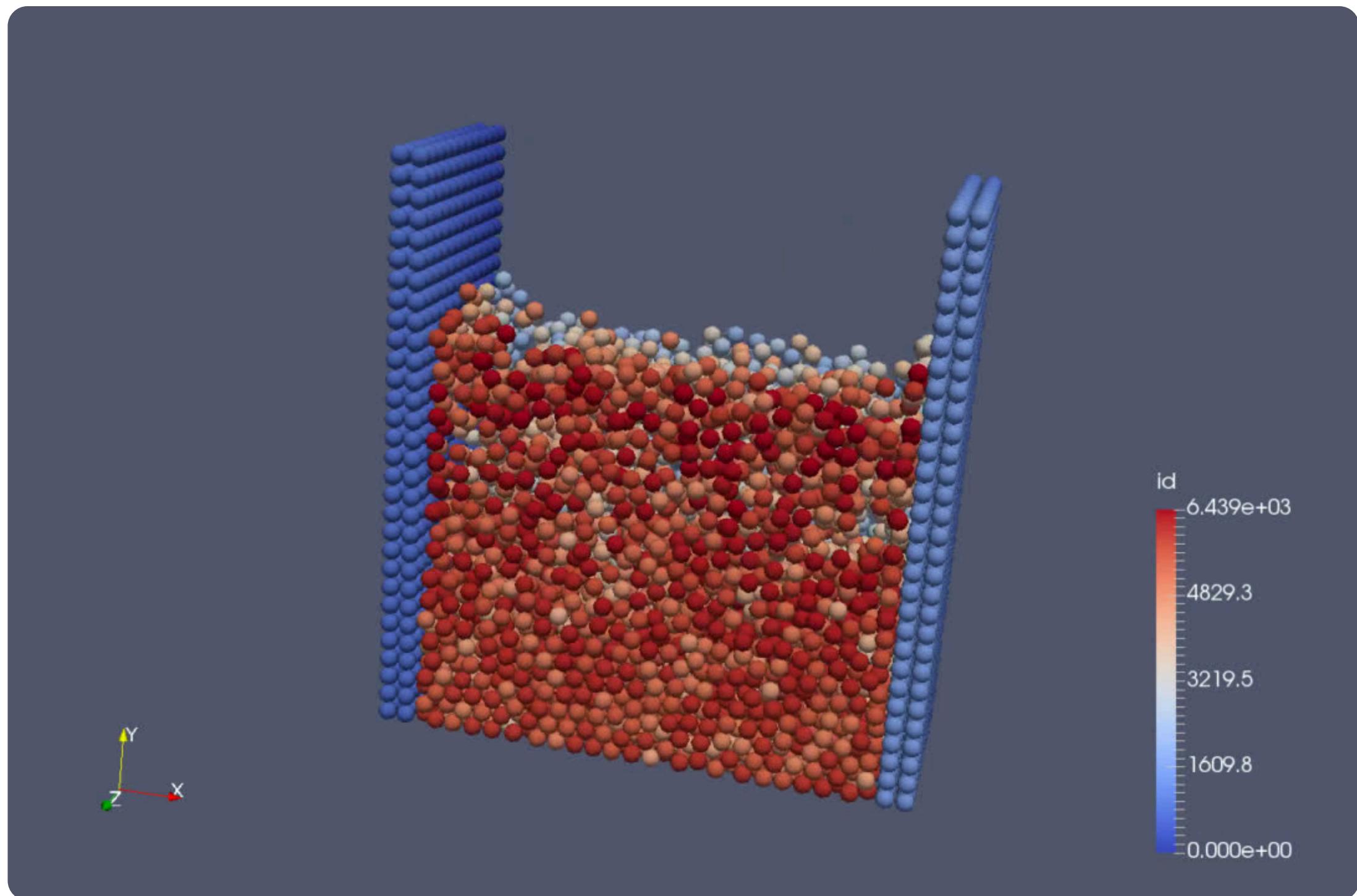


# SIMULATION 4

Colored by id (walls are blue, fluid is red)

Large mass walls

19

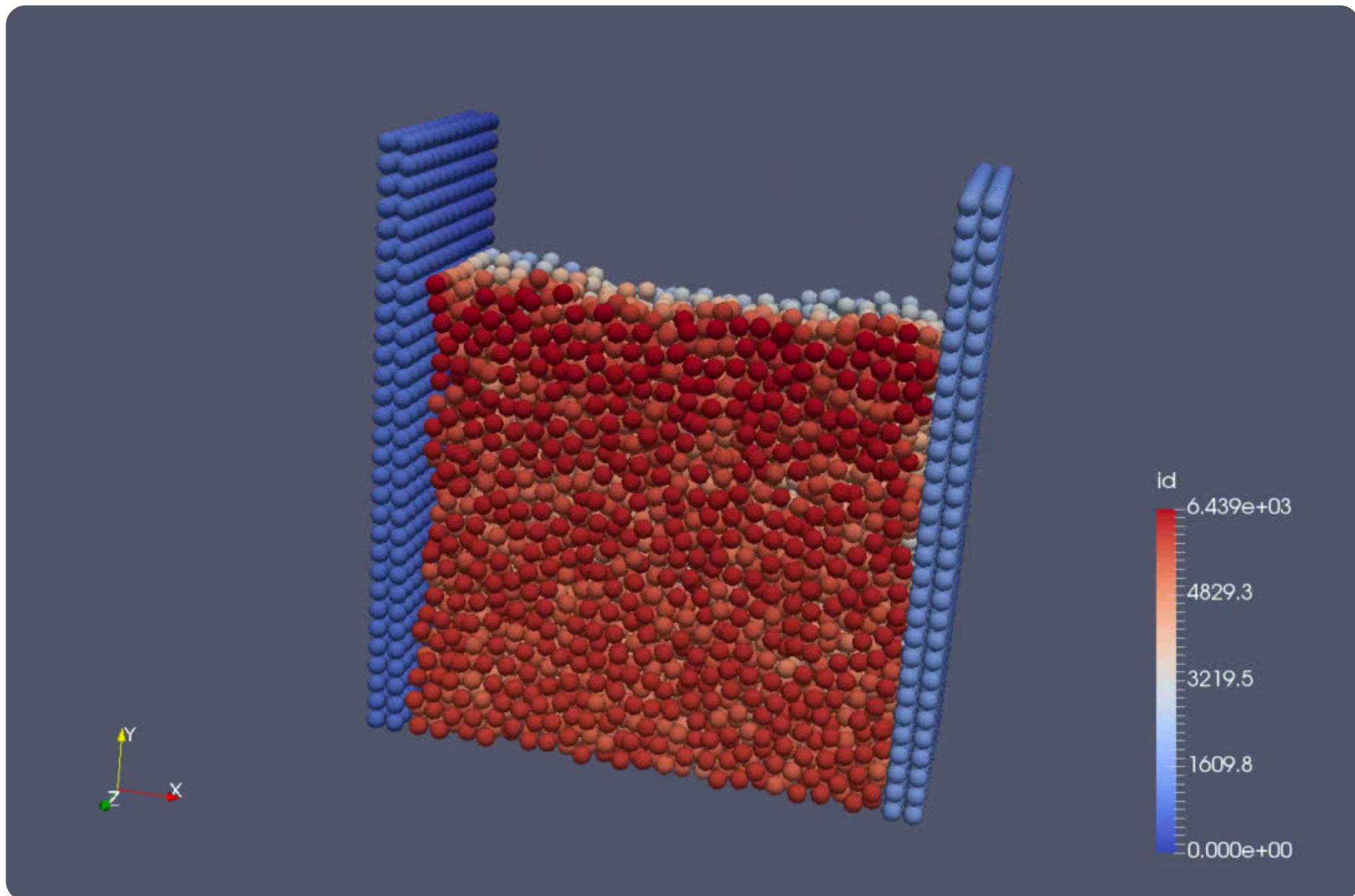


# SIMULATION 5

Large epsilon fluid

Colored by id (walls are blue, fluid is red)

20

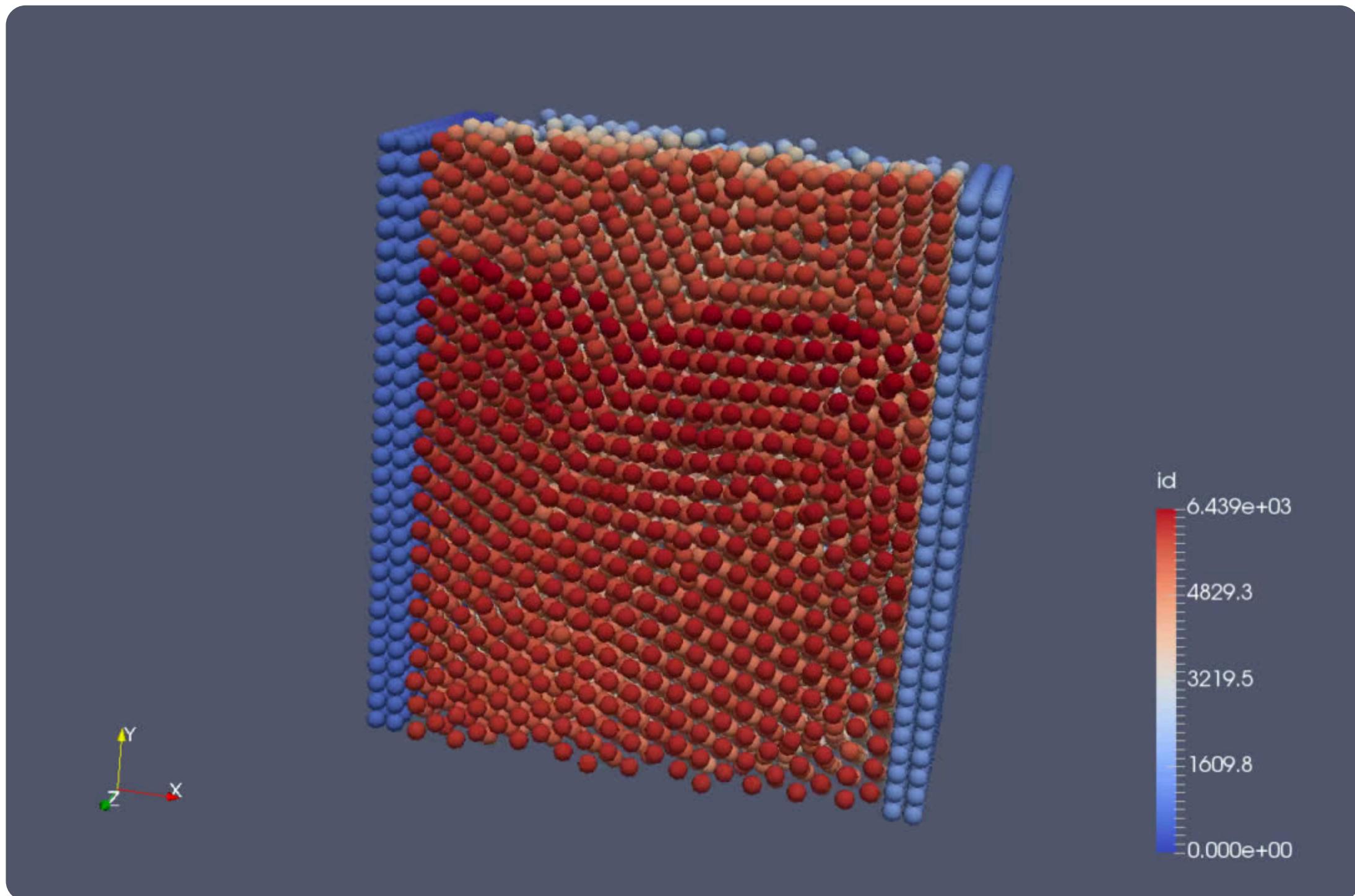


# SIMULATION 6

Large sigma fluid

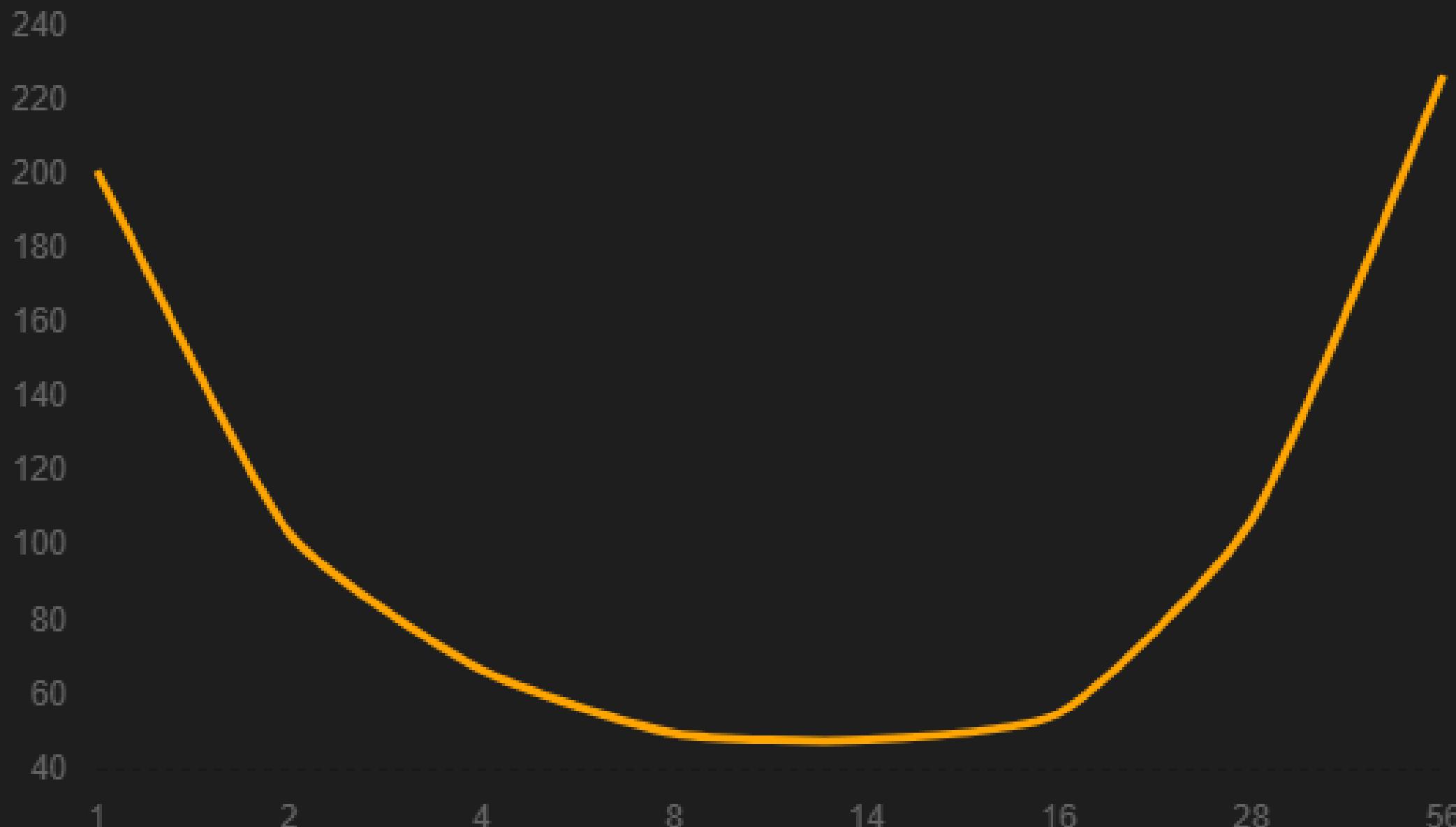
Colored by id (walls are blue, fluid is red)

21



**3D Rayleigh-Taylor: I5 12600k**

Y Execution Time (seconds) von X Number of Threads



# STRONG-SCALING PLOT

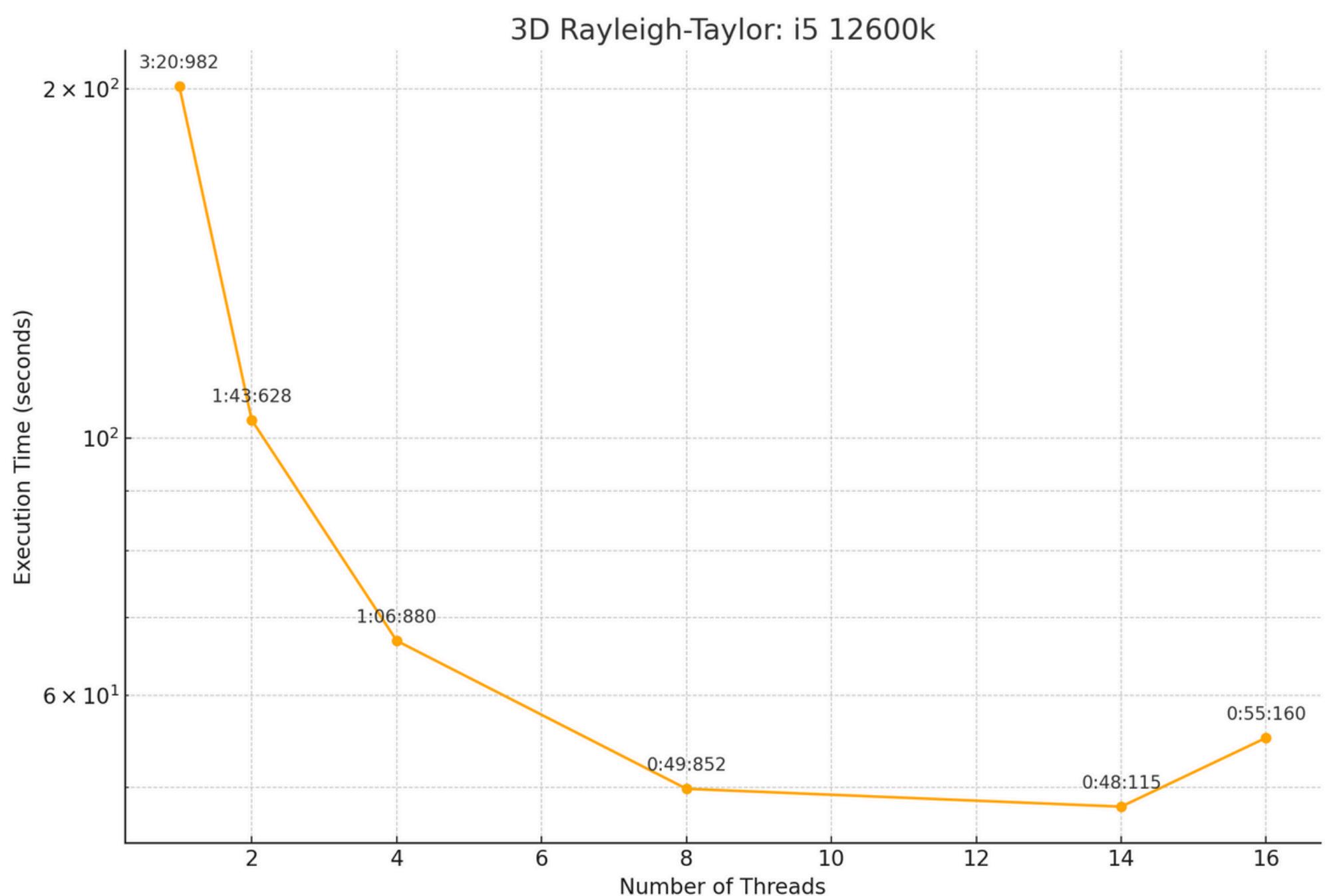
Calculated on local hardware for 1-16 “real” threads according to the used hardware

28 and 56 threads are not real threads but simulated as if they were

we did local computing due to problems on the cluster

Intel i5 12600K 10K/16T  
32GB DDR4 RAM

# STRONG-SCALING PLOT



Plot for the 1-16 actual threads

Intel i5 12600K 10K/16T  
32GB DDR4 RAM

**4-5 hours**

Rayleigh-Taylor Instability 3D

00:**43:933**

Rayleigh-Taylor Instability 3D Contest

08:**42:780**

Nanoscale flow WS5

**LOCAL MACHINE**

**did not run**

02:**03:821**

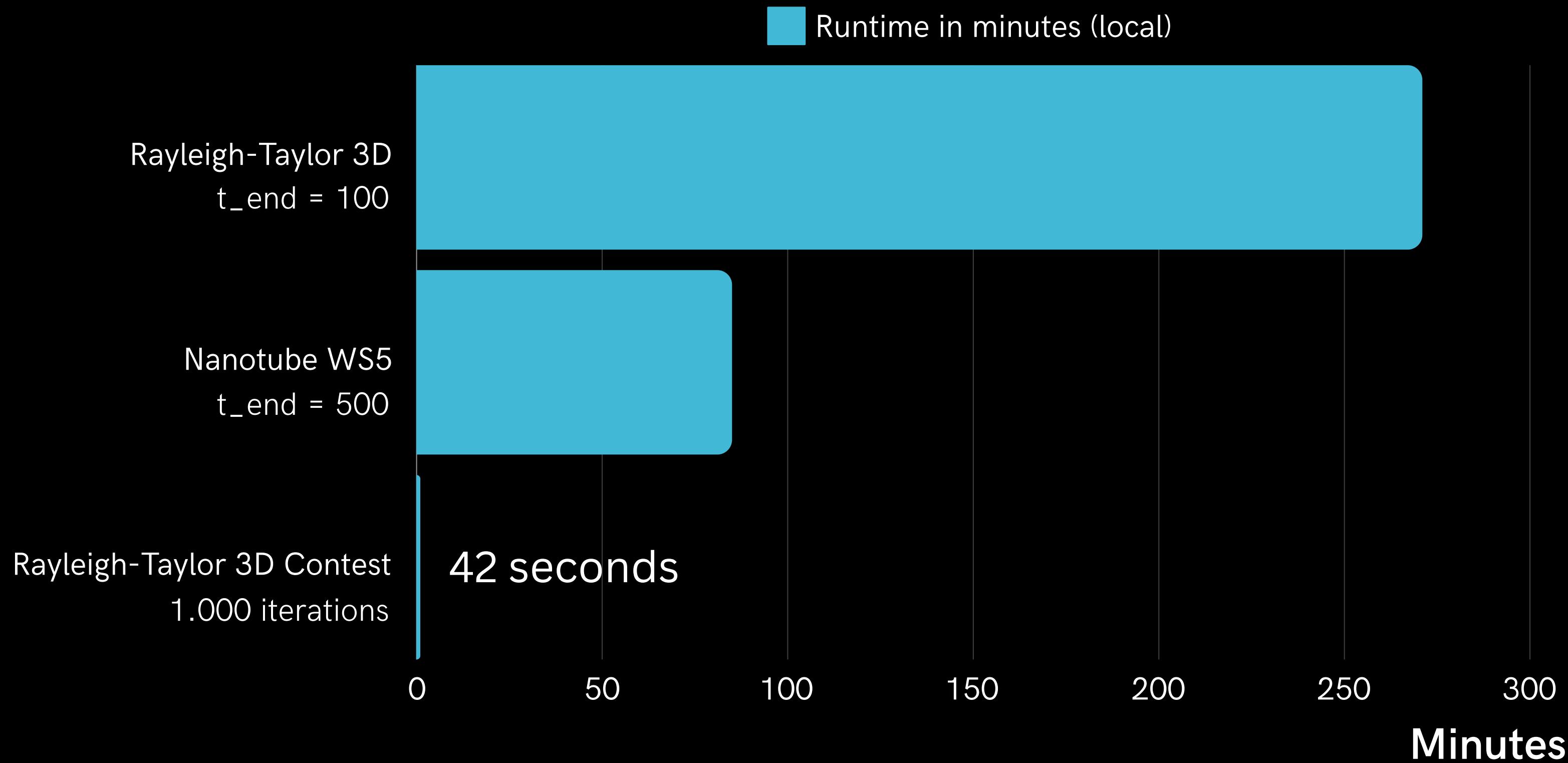
**did not run**

**CLUSTER**

Format  
mm:ss:ms

## BENCHMARKS

Local Hardware: AMD Ryzen 9 7950X3D  
16K/32T, 64GB RAM



Local Hardware: AMD Ryzen 9 7950X3D 16K/32T, 64GB RAM

## **RAYLEIGH TAYLOR CONTEST 2**

1000 iterations of the Taylor-Rayleigh instability simulation in 3D

02:03:821 (mm:ss:ms)

On linux cluster

00:43:76 (mm:ss:ms)

On local machine (Ryzen 9)

## **RAYLEIGH TAYLOR 3D WITH T-END 100**

1000 iterations of the Taylor-Rayleigh instability simulation in 3D

approx. 4-5 hours

(unfortunately got interrupted)

On local machine (Ryzen 9)

08:41:23 (hh:mm:ss)

On local machine (Laptop 12K/24T)

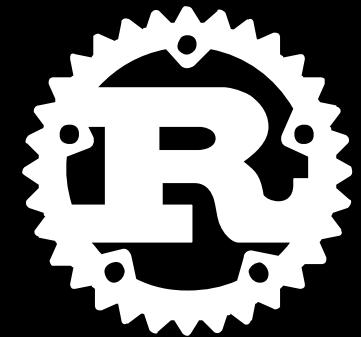
## **CONTEST 2**

## **RT 3D LONG**

# FUTURE IDEAS

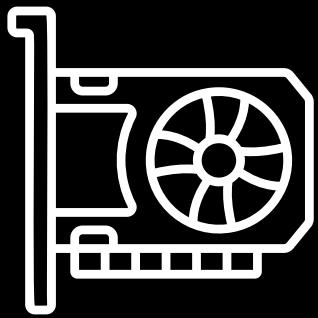
## LANGUAGE

Implementing the same / similar molecular simulator in Rust instead of C++ which has comparable or even faster performance and its own unique components



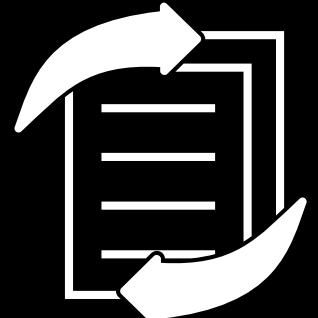
## GPU PERFORMANCE

Optimizing the code for GPU computation to try out performance changes



## REFACTORING

Rewriting the whole code based on the now known features it should support would yield better performance and streamlining





28

Scientific Computing (PSE) Molecular Dynamics  
Group D

Johannes H. | Julius K. | Tim S.

**THANK YOU FOR  
LISTENING**