

# Asynchronous Smoothed Particle Hydrodynamics

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## 1 Abstract

A new method for parallelizing asynchronous Smooth-Particle Hydrodynamics is presented, extending the single-threaded and multi-core asynchronous SPH algorithms of Reinhardt et al[17] to the GPU. The method will then be evaluated for performance against single and multi-core implementations, and further directions for improvement will be discussed.

## 2 Introduction

Computational Fluid Dynamics is predominantly concerned with the numerical integration of the Navier-Stokes equations. These are second order partial differential equations with respect to time (Equation 1) which describe fluid in terms of the *continuum hypothesis*, which models fluids as smooth vector fields[4]. Despite the true nature of fluids being an emergent property of their molecular interactions, the macroscopic behaviours of a fluid can be predicted by this model with sufficient accuracy for most applications[4].

$$d\mathbf{u}/dt = -(\mathbf{u} \cdot \nabla)\mathbf{u} + \nu\Delta\mathbf{u} + \mathbf{g} \quad (1)$$

Velocity of the fluid is represented with vector field  $\mathbf{u}$ , and density is represented with scalar field  $\rho$  (Greek rho, not to be confused with pressure  $p$ ).

Fluid simulation can broadly be grouped into two categories: *Eulerian* and *Lagrangian*. Eulerian simulations observe physical quantities which move past fixed locations in space. Often these fixed points form a regular grid, however some methods also use meshes. Lagrangian simulations instead track the movement of physical quantities through the flow of the fluid. There also exist hybrid approaches which attempt to make use of both paradigms, such as Particle in a Cell (PIC).[13]

Eulerian methods, especially those using grid structures, lend themselves particularly well to parallelization, since individual steps in a simulation can be designed as filters. Eulerian simulations of free surface flows, however, tend suffer from loss of fluid mass over time. In these simulations the distinct boundary between different fluids is typically modelled by a signed distance function, which to flow along with the fluid is advected with the other physical properties, and then since this distorts the distance values it must be corrected back to a signed distance function again.[13] Over time numerical diffusion of this boundary leads to observable mass loss.

Lagrangian methods track fluid using discrete masses, making them immune to this problem altogether. However, these methods suffer another problem. The forces involved when particles interact with each other or with a boundary can be very large for an infinitesimally small amount of time. This is especially true for incompressible fluids, which resist any and all potential compression with tight spring-like forces. However, if we integrate using a larger timestep, these large forces will cause particles to move much farther than they should. This instability cascades into the simulation diverging, or “exploding”. To ensure stability the timestep must always be sufficiently small enough that this will never occur.

Particle-based fluid simulations are also a difficult problem to parallelize efficiently. Similar to the notorious N-Body problem, particles in CFD simulations may potentially interact with any other particle in the system, at any point in time. The random access which this requires is particularly unfriendly to GPUs, which are typically optimized for streamed memory. To make matters worse, to obtain any meaningful results requires a very large number of particles (typically on the order of millions) and the calculations must be done at extremely small time-scales in order to ensure the stability of the system.

## 3 Literature Review

### 3.1 Smoothed-Particle Hydrodynamics

Smoothed-particle Hydrodynamics (SPH) is a Lagrangian simulation method first proposed independently by L.B. Lucy and R.A. Gingold and J.J. Monaghan for simulations in astrophysics.[14, 6] J.J. Monaghan later extended the method to free surface flows.[15]

Smoothed-particle Hydrodynamics is a Lagrangian method which treats discrete masses of fluid as particles which can move freely throughout the simulation domain. These particles are subject to forces due to pressure and viscosity of the fluid. Forces are calculated for each particle based on its current position and velocity and then new position and velocity are computed by integrating these forces over a given time step.

The SPH update step is presented in Algorithm 1. The computation of acceleration has been split as suggested by Ihmsen et al. so that advection forces are considered when computing pressure forces.[11]

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**Algorithm 1** Full SPH step using split forces calculation.

---

```

1: for  $i \in \{1, 2, \dots, n\}$  do
2:   Compute  $\mathbf{a}_i^{visc}$ 
3:   Compute  $\mathbf{a}_i^{ext}$ 
4:    $\mathbf{u}_i \leftarrow \mathbf{u}_i + (\mathbf{a}_i^{visc} + \mathbf{a}_i^{ext})dt$ 
5: end for
6: for  $i \in \{1, 2, \dots, n\}$  do
7:   Compute  $\rho_i$  by Equation 3
8:   Compute  $p_i$  by Equation 2
9: end for
10: for  $i \in \{1, 2, \dots, n\}$  do
11:   Compute  $\mathbf{a}_i^{pres}$ 
12: end for
13: for  $i \in \{1, 2, \dots, n\}$  do
14:    $\mathbf{u}_i \leftarrow \mathbf{u}_i + \mathbf{a}_i^{pres}dt$ 
15:    $\mathbf{x}_i \leftarrow \mathbf{x}_i + \frac{1}{2}(\mathbf{a}_i^{visc} + \mathbf{a}_i^{ext} + \mathbf{a}_i^{pres})dt^2 + \mathbf{u}_i dt$ 
16: end for

```

---

There are two main forces at play in SPH. The first is pressure force  $\mathbf{a}_i^{pres}$

which is calculated from density using an Equation of State (EOS). There are many different choices for this, the one in Equation 2 is commonly used for water. The adiabatic index  $\gamma$  is typically set to 7 for water,  $\rho_0$  is the reference density of the fluid, and  $c_s$  is the speed of sound in the fluid.

$$p_i = \frac{\rho_0 c_s}{\gamma} \left( \left( \frac{\rho_i}{\rho_0} \right)^\gamma - 1 \right) \quad (2)$$

This requires that we determine the density of the fluid at each particle's center point  $\mathbf{x}_i$ . Using the Parzen-Rosenblatt kernel estimation we can define the density field in terms of the particles as point samples as shown in Equation 3. The volume scaling factor  $V_i$  is approximated as  $m_i/\rho_i$  using the physical quantities of mass  $m_i$  and density  $\rho_i$ , which allows us to cancel density. We can then use this to estimate the density at each particle,  $\rho_i$ , which was previously unknown.

$$\rho(\mathbf{x}) = \sum_i^n \rho_i W(\mathbf{x} - \mathbf{x}_i) V_i = \sum_i^n m_i W(\mathbf{x} - \mathbf{x}_i) \quad (3)$$

There are multiple different choices of discretization for the gradient operator, the one presented here is due to Adams and Wicke.[1] One important property which they note the gradient of pressure should possess is that the resulting pressure force is symmetric between pairs of particles (respecting Newton's third law). Because of this, the derivation of the gradient operator is somewhat roundabout, requiring some algebraic manipulation to arrive at the form below.

$$\nabla p_i / \rho_i = \sum (p_j / \rho_j^2 + p_i / \rho_i^2) (x_j - x_i) \nabla W(|x_j - x_i|) m_j \quad (4)$$

The second force we must calculate is the viscosity force  $\mathbf{a}_i^{visc}$ . Viscosity is modelled as the Laplacian of the velocity field. As with the gradient, we also insist in a discretization of the Laplacian which results in symmetric viscosity forces.[1]

$$\nabla \mathbf{u}_i = \sum (\mathbf{u}_j - \mathbf{u}_i) \nabla W(\mathbf{x}_j - \mathbf{x}_i) m_j / \rho_j \quad (5)$$

Until now we have avoided discussing the kernel  $W$ . The kernel should obey three properties: first is that the area under its curve must add to one; second, the kernel should vanish after  $h$  units (so that  $W(h) = 0$ ), where  $h$  is called the support distance; and third, the kernel must be differentiable everywhere on its domain.[13] Many different kernels have been suggested, and we must be careful to use the kernel for the appropriate dimension. Using a kernel designed for  $\mathbb{R}^2$  will not have an integral of 1 in  $\mathbb{R}^3$  (or vice versa). The kernels presented in Equations 6 and 7 are due to M. Müller.[16]

$$W(\mathbf{r}) = \begin{cases} 315 \frac{(1-|\mathbf{r}|/h)^3}{64\pi h^3} & \text{if } |\mathbf{r}| < h \\ 0 & \text{otherwise} \end{cases} \quad (6)$$

$$\nabla W(\mathbf{r}) = \begin{cases} -45\mathbf{r} \frac{(1-|\mathbf{r}|/h)^2}{\pi h^4} & \text{if } |\mathbf{r}| < h \\ \mathbf{0} & \text{otherwise} \end{cases} \quad (7)$$

### 3.2 Neighborhood Search

Every particle in the simulation enacts a force on and likewise has a force enacted on it by every other particle. This means there are  $O(n^2)$  interactions between particles in the system, which is obviously intractable when dealing with millions of particles. Thankfully, the magnitude of these forces vanishes as distance increases, and can be safely ignored after a certain threshold. If we assume an upper limit on the number of possible neighbors a given particle will interact with, we can reduce the simulation from quadratic to linear.

Tree structures are ill suited to the streaming memory models of the GPU and were avoided when optimizing the neighborhood search. A simple alternative which is better suited to the GPU is the cell list. In a cell list, the simulation domain is split into a regular grid of cubes known as cells. Each cell contains an index to an arbitrary particle contained by it. Each particle has associated with it an index to the next particle in the cell. A

unique index value is reserved for denoting that a cell is empty, and that the final particle in a cell list has been reached.[13]

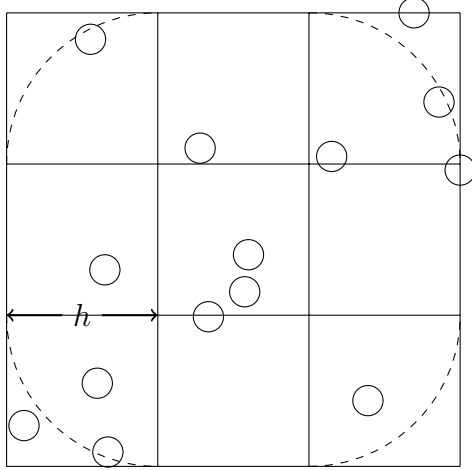


Figure 1: Neighborhood of a cell. Particles inside dashed region are possibly within  $h$  of a point in the center cell.

Now when we wish to list neighbors of a given particle we can look up the particle’s grid index, and then iterate the particles that share that cell, and those neighboring cells within the given distance threshold. Commonly the cell size is chosen to be the same as the distance threshold, which is also usually the support size of our kernel  $W$ , so that we only need to search a  $3 \times 3 \times 3$  cube of cells. This is the approach also taken in this implementation.

### 3.3 GPU Acceleration

The work of Ihmsen et al. provided much of the groundwork with an early parallel implementation.[11] For this search they used a Z-index sort, a space filling curve which provides a cache-friendly ordering for the particles. Amada et al. present a partial GPU implementation which relies on the CPU for the neighborhood search, providing the information to the GPU as a texture.[2] Harada et al. present an early fully GPU implementation[8]. Later H  rault make use of the programmable pipeline to create a CUDA implementation[10], which they later released as open source[9]. Finally, Rustico et al. extend this to multiple GPUs.[18]

### 3.4 Time-step

M. Desbrun and M. Gascuel applied the Courant-Friedrichs-Lewy criterion to SPH, providing an upper bound on the time step based on the kernel support size and the maximum particle velocity[5]. They suggest a using an adaptive time-step based on this condition.

Another approach is to avoid a global time-step altogether. P. Goswami and C. Batty propose segmenting the time-step by spatial chunks.[7]. Asynchronous SPH allows every particle to have its own time frame[17][3][5]. This is more efficient when there are only a few fast particles, as is often the case.

The method of asynchronous SPH suggested by Reinhardt et al. integrate each particle forward in time individually.[17] When describing the method it will be useful to speak in terms of a particle's age, which is the total time which it has been integrated during the simulation. All particles in the simulation start at the same age, and ages of particles are tracked individually. A queue was used to sort the particles by their age, so that the youngest particle was always the next one to be stepped forward in time. This ensures that the neighborhood around the particle would all be older, and therefore only requiring that we backtrack particles - keeping the simulation stable.

To make the algorithm multithreaded, Reinhardt et al. suggest dividing the particles into multiple queues, a method due to Kale and Lew.[17, 12] However, now the particle being integrated might not be the youngest particle in its neighborhood, since other threads may be responsible for particles in the particles' neighborhood. In order to solve this, Reinhardt et al. suggest testing for this before integrating. If a younger particle is found, we place the particle in a second list of pending particles. This list is routinely added back to the queue, in the hope that on dealing with a particle the second time another thread will have taken care of the younger particles.

## 4 Problem Statement

One of the greatest difficulties with SPH over other methods is the incredibly small timesteps required in order for the simulation to remain stable. Since the CFL condition is bounded by the maximum velocity of all the particles in the simulation, only a single fast particle can slow down the entire simulation. By making the simulation asynchronous, we have the opportunity to

localize the effect of individual fast particles on the time-step. The problem which will be examined in this report is how to extend the work of Reinhardt et al.[17] and implement asynchronous smoothed-particle hydrodynamics efficiently using the GPU.

## 5 Method

Originally a per-particle approach was employed in the GPU implementation, much as in the original method. However, instead of a queue, all of the particles were given their own individual thread. The particles who are the youngest of their neighbors are then integrated. However, this was found to be prohibitively expensive to implement on the GPU. While in theory this should be an improvement over the single threaded approach, in tests the simulation ran much slower. The overhead of backtracking the neighborhood of each individual particle costs more than any time saved.

Instead of assigning a single thread to every particle, it was found that instead the problem should be divided by the cells used in the neighborhood search (see Section 3.2). All the particles in a cell are then integrated at once. This allows us to re-use the backtracked neighborhood surrounding the cell for each particle in the cell, greatly reducing the overhead of backtracking while still allowing different areas of the simulation to be integrated at different rates. This is similar to the regional time stepping of P. Goswami and C. Batty, while retaining individual timeframes per particle.

A cell is only updated if it contains the youngest particle of all its neighboring cells. This ensures that no region of the simulation is stepped too far ahead of other regions. This also ensures that it is never necessary to extrapolate the location of particles, only interpolation of physical values is required to bring a neighborhood of particles to the same frame of reference in time, increasing the stability of the simulation.

All the particles in a cell are integrated simultaneously by the same time step. The time-step  $dt$  is determined by Equation 8, due to Reinhardt et al., which ensures by the Courant-Friedrichs-Lewy condition that the simulation remains stable. The addition of constant term  $\epsilon = 0.001$  in the denominators assures the time-step remains defined even for particles which are not moving or accelerating. The scaling factors were determined experimentally.



$$dt = \min_{x_i \in \mathcal{N}} \left\{ 0.02 \sqrt{\frac{h}{\epsilon + |a_i|}}, 0.05 \frac{h}{\epsilon + |u_i|} \right\} \quad (8)$$

## 5.1 Implementation

The implementation was written in CUDA and tested on a RTX 3060. The details of this implementation will now be examined. The code for advancing a frame in the simulation is provided below.

First we update the cells for the neighborhood search using `updateCells` (line 1), and copy the result to the GPU (lines 2-9). This is done only once per global update, and so is currently done on the CPU.

```

1 | updateCells();
2 | cudaMemcpyAsync( particles_dev , particles.data() ,
3 |   numParticles*sizeof( Particle ) , cudaMemcpyHostToDevice );
4 | cudaMemcpyAsync( nextParticles_dev , particles_dev ,
5 |   numParticles*sizeof( Particle ) , cudaMemcpyDeviceToDevice );
6 | cudaMemcpyAsync( cells_dev , cells.data() ,
7 |   200*200*200*sizeof( Cell ) , cudaMemcpyHostToDevice );
8 | cudaMemcpyAsync( nextCells_dev , cells_dev ,
9 |   200*200*200*sizeof( Cell ) , cudaMemcpyDeviceToDevice );

```

Next we increase the global `time` of the simulation (line 10). Every cell must be at least this old before we finish this cycle in order for us to produce the next frame of animation.

With the updated particle and cell lists on the device, we can begin simulation. This is done by `stepCell`, configured with 8000 workgroups aligned in a  $20 \times 20 \times 20$  grid with 160 threads each (line 15).

Memory cannot be written in place, since this would lead to race-time conditions between workgroups. Therefore, two device buffers are used for both the cells and particle lists. One is read from while the other is written to. In order to save on copies, a so called ‘ping-pong buffer’ strategy is employed, where the roles of the buffers are reversed by swapping the pointers (lines 18-19).

```

10 | time += 1.0f/60.0f;
11 |
12 | device_ptr<Cell> p = device_pointer_cast(cells_dev);
13 | do {
14 |     for (int k = 0; k < 50; k++) {
15 |         stepCell<<<<dim3(20,20,20),160>>>>(cells_dev ,
16 |             nextCells_dev , particles_dev , nextParticles_dev ,
17 |             time);
18 |         std::swap(cells_dev , nextCells_dev);
19 |         std::swap(particles_dev , nextParticles_dev);
20 |     }
21 |     cudaDeviceSynchronize();
22 | } while ( find_if(thrust::device , p, p+200*200*200,
23 |                 isUpdated()) == p+200*200*200);

```

The kernel is called in batches of 50, after which we synchronize the host and device and check if every cell is older than `time`. To do this we use the Thrust library provided by NVIDIA to do perform a fast search through reduction on the cells.

Once all the cells have been sufficiently updated, we then copy the cells and particles back to the host. Before we do this we might optionally sort the particles by cell index, in order to make access to the particle buffer more contiguous and therefore cache friendly. This step was left out of the final implementation as it was measured to be slightly slower. However, for larger numbers of particles or on different GPUs where cache may be more of a bottleneck, this step may serve to help.

```

24 | // Optional sort
25 | device_ptr<Particle> p = device_pointer_cast(particles_dev);
26 | sort(device , p, p+numParticles);
27 |
28 | cudaMemcpyAsync(particles.data() , particles_dev ,
29 |     numParticles*sizeof(Particle) , cudaMemcpyDeviceToHost);
30 | cudaMemcpyAsync(cells.data() , cells_dev ,
31 |     200*200*200*sizeof(Cell) , cudaMemcpyDeviceToHost);
32 | cudaDeviceSynchronize();

```

Now let's examine the kernel `stepCell` itself. Each invocation of `stepCell` is responsible for one main cell, and populates its neighbor list from the neighboring cells. Each workgroup is divided into 160 threads, each sharing the backtracked neighbor list.

Due to the linear nature of the cell list, thread 0 is tasked with populating this list first from the cells. It iterates through all the particles in each cell backtracking them to the same time and then adding the backtracked version to the shared list (lines 58-90).

```

33  __global__
34  void stepCell(Cell* cells, Cell* nextCells, Particle* particles,
35              Particle* nextParticles, float timeLimit) {
36      constexpr static glm::ivec3 deltas[] = {
37          glm::ivec3(-1,-1,-1),
38          // ...
39          glm::ivec3(+1,+1,+1),
40      };
41
42      ivec3 center(90+blockIdx.x, 90+blockIdx.y, 90+blockIdx.z);
43      int centerIndex = center.x+center.y*200+center.z*200*200;
44
45      nextCells[centerIndex] = cells[centerIndex];
46      nextCells[centerIndex].updateCounter = 0;
47
48      Cell& centerCell = cells[centerIndex];
49      if (centerCell.firstParticle == -1
50          || centerCell.time > timeLimit)
51          return;
52
53      __shared__ Particle neighbors[160];
54      __shared__ int numNeighbors;
55      __shared__ float neighborTime;
56      __shared__ bool doStep;
57
58      if (threadIdx.x == 0) {
59          doStep = true;
60          numNeighbors = 0;
61
62          for (int i = centerCell.firstParticle;
63              i != -1;
64              i = particles[i].nextParticle) {
65              neighbors[numNeighbors++] = particles[i];
66              assert(numNeighbors != 160);
67          }
68
69          neighborTime = std::numeric_limits<float>::max();
70          for (ivec3 delta : deltas) {
71              const ivec3 index = center+delta;

```

```

72         const Cell& cell =
73             cells[index.z*200*200+index.y*200+index.x];
74         if (cell.firstParticle == -1)
75             continue;
76         else if (cell.time < centerCell.time)
77             doStep = false;
78
79         neighborTime = min(neighborTime, cell.time);
80         for (int i = cell.firstParticle;
81             i != -1;
82             i = particles[i].nextParticle) {
83             float t = (centerCell.time - cell.time + cell.
84                 deltaTime) / cell.deltaTime;
85             neighbors[numNeighbors++] = backtrace(particles[
86                 i], t);
87             assert(numNeighbors != 160);
88         }
89     }
90     --syncthreads();

```

Once this is finished, the  $n$ -th thread searches for the  $n$ -th particle in the main cell. It then calculates the forces and integrates using `stepParticle` (line 96), storing the result to the next particle buffer (rather than the one being read). As a final step, the first thread updates the main cell's time.

```

90     for (int k = 0, i = centerCell.firstParticle;
91         i != -1;
92         i = particles[i].nextParticle, k++) {
93         if (k == threadIdx.x) {
94             if (doStep)
95                 nextParticles[i] = stepParticle(neighbors[k],
96                     neighbors, numNeighbors, centerCell.
97                         deltaTime);
98             else
99                 nextParticles[i] = particles[i];
100             break;
101         }
102     }
103     if (threadIdx.x == 0 && doStep) {
104         nextCells[centerIndex].updateCounter = 1;
105         nextCells[centerIndex].time =

```

```

106 |         centerCell.time+centerCell.deltaTime;
107 |     }
108 | }

```

Note that it is important that we copy the particles even if we do not step the cell forward in time (line 101). Otherwise information is not propagated when using the ping-pong buffer strategy.

## 6 Results

Now the results of the implementation described will be examined. In Figure 6 we see the classic ball drop test. The frames have been displayed by ray-marching a signed distance function produced by spheres centered at each particle and globally smoothed.

The test only involves a small number of particles (approximately a thousand) but shows that the method works and exhibits fluid-like behavior (more obvious in the animations). The OpenMP implementation is roughly four times slower on average, with a peak of 4.5s to calculate the second frame. The CUDA implementation remains nearly constant at 0.25s per frame for the duration of the test.

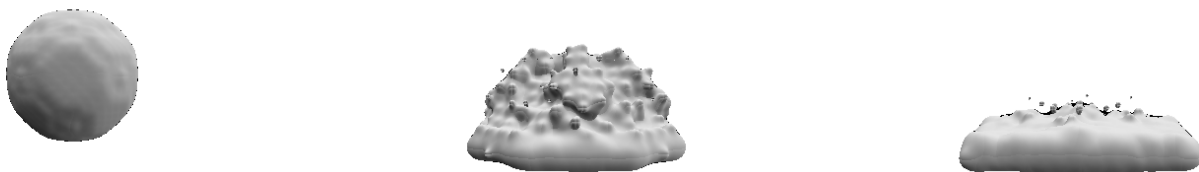


Figure 2: Frames 1,14 and 20 from ball drop test.

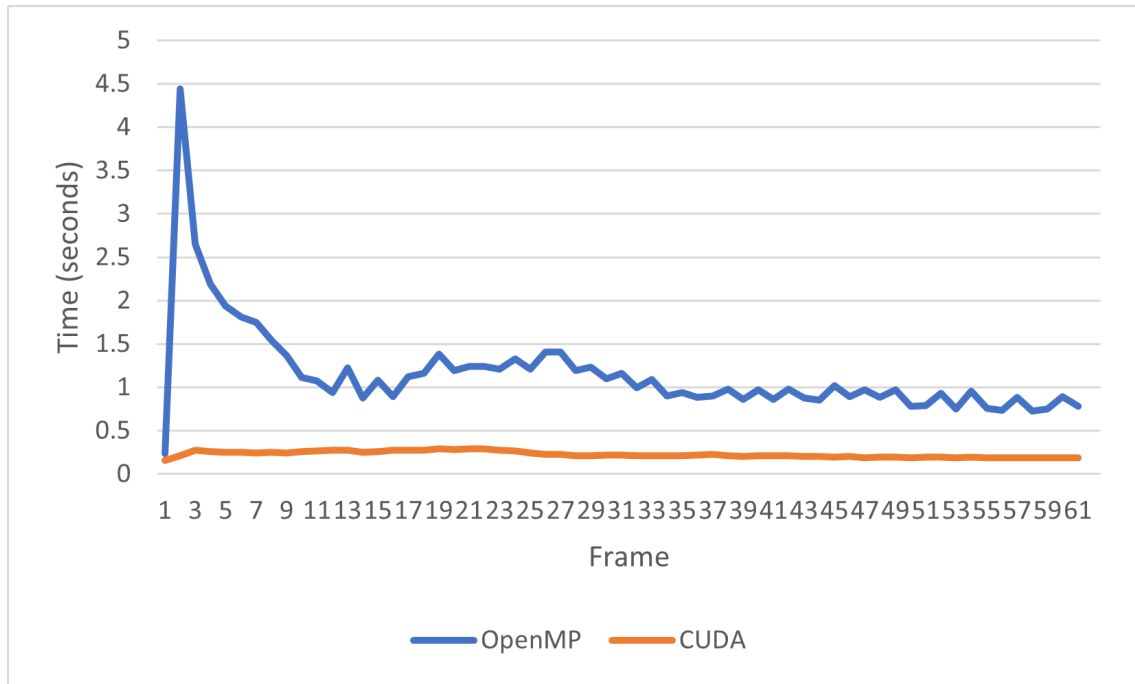


Figure 3: Performance of Ball Drop test.

Figure 6 shows the results of a larger test. Note the steep spike in computation time for frames 16 to 45. This was likely due to the nature of the test: as the particles become more compressed this increases the number of interactions between particles. Then as the pressure force restores the density of the fluid to the target value the performance begins to increase.

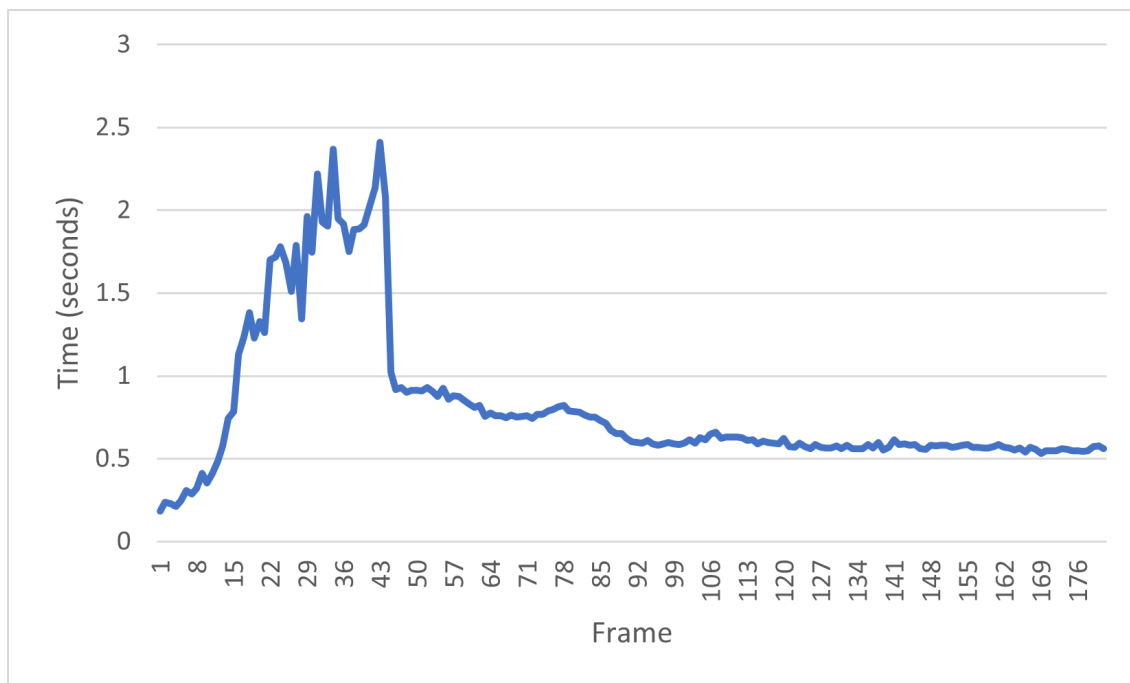


Figure 4: Performance of Dam Break test.

The types of physics simulations which use SPH often require up to tens of millions of particles. Therefore, as a final test, another ball-drop test with approximately one million particles was performed. Unfortunately, the implementation was not able to handle the large amount of neighbors once the ball began to compress after impact with the floor plane. In the next section some potential directions for solving this problem will be discussed.

## 7 Future Work

A current restriction is the fixed size of the simulation domain. This was done to allow the cells to be parallelized as individual workgroups. A direction of future research would be how to effectively extend the boundary beyond the current limitations in size placed by the number of GPU workgroups. A multigrid approach would be one potential solution: the domain could be broken into chunks of cells which would be small enough to be parallelized. In such an approach one would need to be careful to account for the boundary

between chunks to that information properly propagates between them. One possibility here would be to use regions which overlap by a few cells.

Another restriction with the current implementation is the fixed number of neighbors, limited by the size of shared memory between threads in a workgroup. Each potential neighbor also adds a thread, which increases the overall computational cost. In a perfectly incompressible fluid, the upper limit on the possible number of neighbors a particle can have is fixed, since the number of spheres that can be packed in a given volume is fixed. SPH, however, there is a delay between fluids moving too close together and the pressure forces moving them apart again. This delay allows even incompressible fluids such as water to compress momentarily.

This could be addressed by using PCISPH, which enforces the incompressibility of the fluid by relaxing the particles based on the density field. This would ensure that the number of neighbors never rises beyond a constant number.

## 8 Conclusion

A new method for parallelizing asynchronous SPH for GPGPUs using CUDA has been demonstrated, extending the work of [1]. Though similar to the work of P. Goswami and C. Batty, the particles each carry their own time frame, making the simulation fully asynchronous. The method provides visually correct results and shows promising performance on smaller tests. However, future work is needed to scale up to the number of particles required in typical scientific simulations. In particular, the fluid must not be compressible in order to maintain the assumption that the number of neighbors in a cell is fixed.

## 9 Appendix

The implementation of `stepParticle` is provided below, following the algorithm presented in Section 3.1.

```

109 | __device__
110 | Particle stepParticle(Particle& particle_, Particle* neighbors,
111 |                     int numNeighbors, float dt) {
112 |     Particle particle = particle_;

```



```

113 particle.lastPosition = particle.position;
114 particle.lastVelocity = particle.velocity;
115 particle.lastDensity = particle.density;
116 particle.lastPressure = particle.pressure;
117
118 // Compute density
119 particle.density = 0.0f;
120 for (int i = 0; i < numNeighbors; i++) {
121     auto xij = particle.position - neighbors[i].position;
122     particle.density += Particle::mass*W(xij);
123 }
124
125 // Compute F* (Fvisc + Fext)
126 constexpr float nu = 0.015f;
127 particle.accel = glm::vec3(0.0f, -9.81f, 0.0f);
128 for (int i = 0; i < numNeighbors; i++) {
129     auto vij = particle.velocity - neighbors[i].velocity;
130     auto xij = particle.position - neighbors[i].position;
131     if (length2(xij) > 0.0) {
132         particle.accel += 2.0f * nu
133             * Particle::mass / neighbors[i].density
134             * vij * dot(xij, dW(xij))
135             / (dot(xij, xij) + 0.01f * sq(Particle::radius))
136         ;
137     }
138 }
139 // Compute velocity using forces
140 particle.velocity += dt*particle.accel;
141
142 // Compute new density
143 particle.density = 0.0f;
144 for (int i = 0; i < numNeighbors; i++) {
145     auto xij = particle.position - neighbors[i].position;
146     auto vij = particle.velocity - neighbors[i].velocity;
147     particle.density += Particle::mass*W(xij);
148     particle.density += dt*dot(dW(xij), vij);
149 }
150
151 // Compute pressure and pressure forces
152 constexpr float kappa = 0.5f;
153 auto accelP = glm::vec3{0.0f,0.0f,0.0f};
154 particle.pressure = kappa
155     * std::max(particle.density - Particle::rho0, 0.0f);
156 for (int i = 0; i < numNeighbors; i++) {

```

```

157         auto xij = particle.position - neighbors[i].position;
158         if (length2(xij) > 0.0) {
159             accelP -= dW(xij) * Particle::mass
160                 * (particle.pressure / sq(particle.density)
161                 + neighbors[i].pressure / sq(neighbors[i].
162                     density));
163         }
164         particle.accel += accelP;
165
166         // Integrate particle over time using dt
167         particle.velocity += dt*accelP;
168         particle.position += dt*particle.velocity
169             + sq(dt)/2.0f*particle.accel;
170
171         glm::vec3 r = glm::vec3(0.4f,0.3f,0.4f);
172         glm::vec3 a = 5.0f-r, b = 5.0f+r;
173         for (int d = 0; d < 3; d++) {
174             if (particle.position[d] < a[d]) {
175                 particle.position[d] = a[d]
176                     +std::min(b[d]-a[d], a[d]-particle.position[d]);
177                 particle.velocity[d] *= -0.2f;
178             }
179             if (particle.position[d] > b[d]) {
180                 particle.position[d] = b[d]
181                     -std::min(b[d]-a[d], particle.position[d]-b[d]);
182                 particle.velocity[d] *= -0.2f;
183             }
184         }
185         particle.time += dt;
186         return particle;
187     }

```

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