

# CS5220 HW2 Report: Shallow Water Simulation

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## 1. Introduction

In this report, we explore parallelization strategies for solving the shallow water equation. We start by profiling the naive implementation of the code in order to identify bottlenecks as well as build our intuition for developing a relatively accurate performance model. This information is used to guide the parallelization of the code for both the compute nodes (i.e., Intel Xeon E5-2620) and the accelerator boards (i.e., Intel Xeon Phi 5110P) on the Totient cluster. We further implement and evaluate several optimizations for tuning the parallel code.

## 2. Profiling the Shallow Water Simulation

Understanding which areas of the code most time is spent, and where opportunities for performance gains are available is key to achieving good speedups in general, and in particular for our parallel implementation.

We will focus on run-time profiling of the code, as well as compile-time reports generated by the Intel compiler to guide our efforts.

### 2.1. Identifying Bottlenecks

As a first pass, we profiled the provided default code using the *amplxe* tool, with truncated results given below.

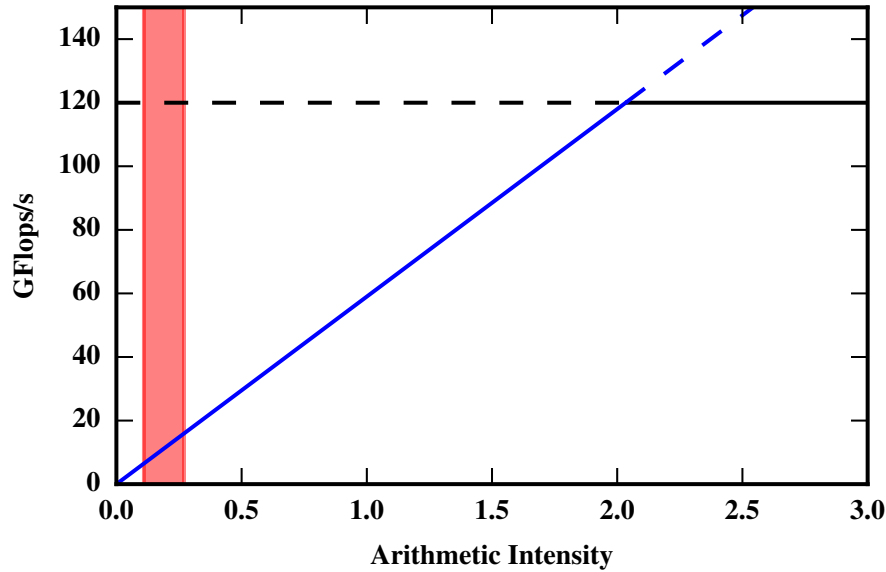
Function	CPU Time
Central2D<Shallow2D, MinMod<float>>::limited_derivs	1.350 s
Central2D<Shallow2D, MinMod<float>>::compute_step	0.652 s
Central2D<Shallow2D, MinMod<float>>::compute_fg_speeds	0.236 s

Expectedly, the vast majority of the time is spent inside the functions for the limiter, computing the step, and computing the wave speeds. What is surprising is the amount of time spent inside the limiter. Given a cursory glance at the code, one would assume that the `compute_step` function would be much more expensive than `limited_derivs`, yet we are seeing just over 2 times as much time spent inside `limited_derivs`. Identifying the cause and improving this performance bottleneck should be a main priority.

One clue may be that the arithmetic intensity of `limited_derivs` is much lower than `compute_step` and `compute_fg_speeds`. We analyzed the binary with *MAQAO*, and arithmetic intensity of the loops were computed. The results can be seen in the table below.

Function	Arithmetic Intensity (AI)
<code>limited_derivs</code>	0.11
<code>compute_step</code>	0.23-0.27
<code>compute_fg_speeds</code>	0.21

Figure 1 displays the roofline model for a single node, with the red shaded area representing our range of arithmetic intensity<sup>1</sup>. As can be seen in the figure, we are well into the memory-bound range, and thus regularizing memory access is likely to lead to good speedups.



**Figure 1:** Roofline Model of our computation. Red shaded area represents our estimated arithmetic intensity.

We can also look at more detailed profiling of individual functions. As an example, below is an excerpt from a profiling report for the `compute_step` function.

```
// Predictor (flux values of f and g at half step)
for (int iy = 1; iy < ny_all-1; ++iy)
    for (int ix = 1; ix < nx_all-1; ++ix) {
        vec uh = u(ix, iy);
        for (int m = 0; m < uh.size(); ++m) {
            uh[m] -= dtcdx2 * fx(ix, iy)[m];
            uh[m] -= dtcdy2 * gy(ix, iy)[m];
        }
        Physics::flux(f(ix, iy), g(ix, iy), uh);
    }

// Corrector (finish the step)
for (int iy = nghost-io; iy < ny+ngghost-io; ++iy)
    for (int ix = nghost-io; ix < nx+ngghost-io; ++ix) {
        for (int m = 0; m < v(ix, iy).size(); ++m) {
            v(ix, iy)[m] =
                0.2500 * ( u(ix, iy)[m] + u(ix+1, iy)[m] +
                          u(ix, iy+1)[m] + u(ix+1, iy+1)[m] ) -
                0.0625 * ( ux(ix+1, iy)[m] - ux(ix, iy)[m] +
                          ux(ix+1, iy+1)[m] - ux(ix, iy+1)[m] +
```

<sup>1</sup>Maximum memory bandwidth found at

[http://ark.intel.com/products/83352/Intel-Xeon-Processor-E5-2620-v3-15M-Cache-2\\_40-Ghz](http://ark.intel.com/products/83352/Intel-Xeon-Processor-E5-2620-v3-15M-Cache-2_40-Ghz)

and maximum compute speed found at

[http://download.intel.com/support/processors/xeon/sb/xeon\\_E5-2600.pdf](http://download.intel.com/support/processors/xeon/sb/xeon_E5-2600.pdf)

```

                uy(ix ,   iy+1)[m] - uy(ix ,   iy)[m] +
                uy(ix+1,iy+1)[m] - uy(ix+1,iy)[m] ) -      0.009s
dtcdx2 * ( f(ix+1,iy  )[m] - f(ix ,iy  )[m] +
           f(ix+1,iy+1)[m] - f(ix ,iy+1)[m] ) -      0.049s
dtcdy2 * ( g(ix ,   iy+1)[m] - g(ix ,   iy)[m] +      0.004s
           g(ix+1,iy+1)[m] - g(ix+1,iy)[m] );      0.044s
    }
}

// Copy from v storage back to main grid
for (int j = nghost; j < ny+nghost; ++j){
    for (int i = nghost; i < nx+nghost; ++i){
        u(i , j) = v(i-io , j-io);
    }
}

```

Although not very enlightening, it verifies our intuition that the corrector portion should be most heavily targeted for optimization, although a non-negligible amount of time is spent in other areas.

### 2.1.1. Vectorization

Significant performance gains can be obtained by vectorizing our functions. Vectorization is “the unrolling of a loop combined with [...] SIMD instructions”<sup>2</sup>.

We decided that rather than attempt to write SSE/AVX instructions by hand, we would rely on the autovectorization capabilities of the Intel compiler. Guiding our efforts is the optimization/vectorization report generated by the intel compiler.

Looking at the vectorization report generated by compiling the default code, we see that the compiler did not vectorize any loop, with many message like this one, which corresponds to the `compute_fg_speeds` function.

```

LOOP BEGIN at central2d.h(268,9)
    remark #15344: loop was not vectorized: vector dependence prevents
                    vectorization
    remark #15346: vector dependence: assumed FLOW dependence between
                    _M_elems line 74 and _M_elems line 76
    remark #15346: vector dependence: assumed ANTI dependence between
                    _M_elems line 76 and _M_elems line 74
LOOP END

```

Full definitions for the different types of vector dependence can be found in the Intel vectorization document linked above, but the basic idea is that the compiler has to assume that arrays may refer to overlapping memory locations. We can invite the compiler to ignore this potential dependency (if we as the programmers know it to be false), with the `#pragma ivdep` directive.

Doing so for the `compute_fg_speeds` results in the compiler vectorizing the loop, and realizing a potential speedup of 3.84.

```

LOOP BEGIN at central2d.h(269,9)
    remark #15300: LOOP WAS VECTORIZED

```

<sup>2</sup><https://software.intel.com/sites/default/files/m/4/8/8/2/a/31848-CompilerAutovectorizationGuide.pdf>

```

remark #15460: masked strided loads: 6
remark #15462: unmasked indexed (or gather) loads: 6
remark #15475: — begin vector loop cost summary —
remark #15476: scalar loop cost: 308
remark #15477: vector loop cost: 76.870
remark #15478: estimated potential speedup: 3.840
remark #15479: lightweight vector operations: 109
remark #15481: heavy-overhead vector operations: 1
remark #15487: type converts: 8
remark #15488: — end vector loop cost summary —
LOOP END

```

Full vectorization of the remaining code is not so simple. The way the data is currently laid out, as an array of structs representing solution values, rather than as a struct of arrays, makes it hard for the compiler for autovectorize. Such a hierarchy is likely why Professor Bindel elected to re-write everything in C – a more in depth discussion of how to remain in C++ will be covered in section 4.1.

To vectorize the *limited\_derivs*, we ended up borrowing some Professor Bindel’s C version of *minmod.h*, which combined with a *#pragma ivdep* allowed the compiler to vectorize.

## 3. Parallelizing the Shallow Water Simulation

### 3.1. Parallelizing for the Compute Node

A relatively simple way to parallelize work in any structured grid computation, including the shallow water equation solver in this assignment, is to divide the grid into blocks and assign each thread a block for which to compute the solution. We will refer to the cells in this block as *live cells*. Depending on the stencil radius of the computation kernel, we also need a ring of padding cells called *ghost cells* that are consumed in order to calculate the solution for the boundary cells of the block. Here we redefine a *block* as the live cells and ghost cells required by a thread to calculate the solutions for all of the live cells. We say ‘consume’ since after advancing a timestep, an additional outermost *r* layers of the ghost cell ring become obsolete, where *r* is the stencil radius. The radius of the ring of ghost cells we need depends on the number of timesteps we want each thread to execute before synchronizing (i.e., updating the ghost cells with new solutions from other threads). Executing more than one timestep before synchronization is referred to as *batching* and will be discussed in Section 4. For this section, we assume each thread only executes a single timestep before synchronization.

In order to apply the blocking parallelization to the shallow water equation solver, we first need to identify which functions can or cannot be parallelized in the computation kernel. `Central2D::run()` calls the following functions:

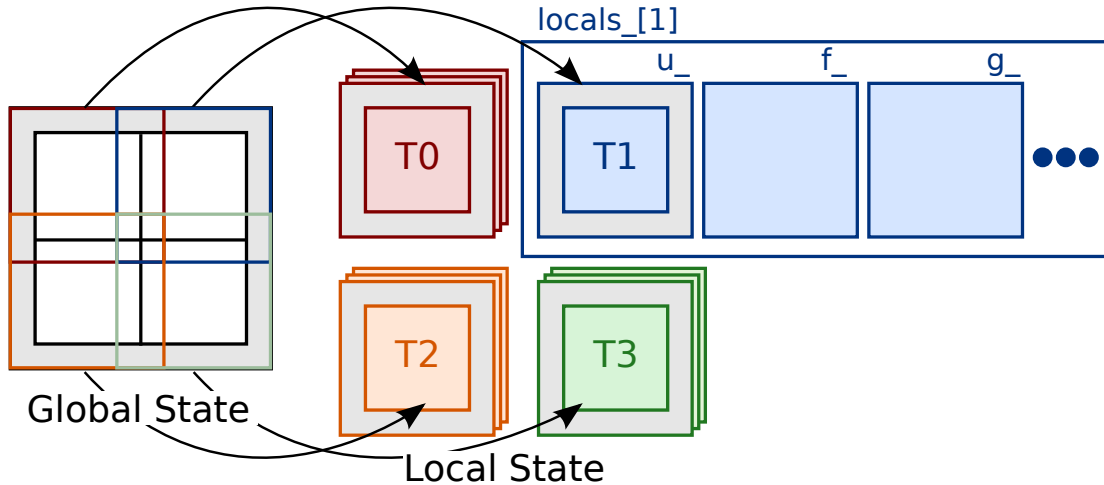
- `apply_periodic()`
- `compute_fg_speeds()`
- `limited_derivs()`
- `compute_step()`

The `apply_periodic()` function updates the ghost cells of the grid by copying the outermost *r* layers of the live cells in a periodic fashion, where *r* is the ghost cell radius. This function forces a synchronization

point since the ghost cells in a given block depend on the live cells of other blocks, meaning each thread has to communicate the solutions of the live cells in its block to all of the other threads. As such, this function only needs to be called at the beginning of every super-step (i.e., main computation sub-step + staggered computation sub-step).

The `compute_fg_speeds()` function is actually comprised of two separate logical functions: (1) calculating the maximum wave speeds in the x/y directions, and (2) updating the flux vectors. (1) is only used to calculate the `dt` which does not change within a super-step and uses expensive square root operations, so we separate this into a `compute_wave_speeds()` function that is called once before every super-step. This function is similar to `apply_periodic()` in that it requires synchronization across threads, in this case to calculate the maximum wave speeds across all blocks in the grid, thus making it difficult to parallelize naively. On the other hand, (2) still needs to be called at every sub-step, but can independently calculate the `f` and `g` vectors for a block without synchronization.

The `limited_derivs()` function calculates the `ux`, `uy`, `fx`, and `gy` vectors from the `u`, `f`, and `g` vectors to implement the MinMod limiter. The `compute_step()` function then uses these output vectors to calculate the solution vectors, `u`, for the current timestep. These functions can also independently calculate the output vectors without synchronization.



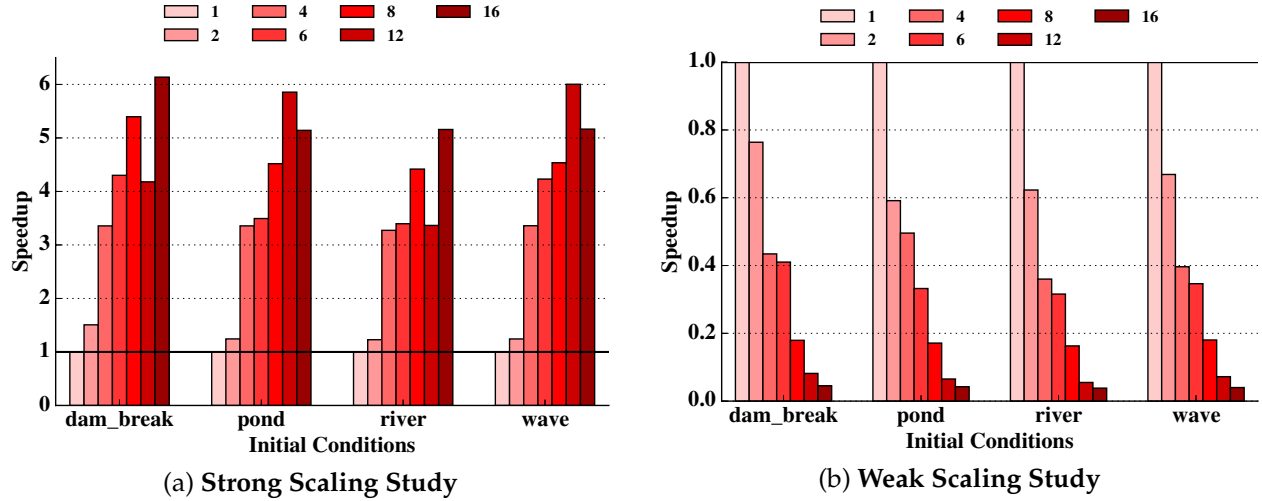
**Figure 2: Parallelizing Computation Using Per-Thread Local State** – In this example, the global grid is divided into four local blocks, each of which is copied to per-thread memory allocated separately from the global grid. The `LocalState` class encapsulates all vectors in a local block including `u_`, `f_`, `g_`, `ux_`, `uy_`, `fx_`, `gy_`, and `v_`. Each thread is only responsible for computing the solution of the live cells (i.e., non-gray area) in its local block. In order to do so, the vectors in the ghost cells (i.e., gray area) must be used. Note that the ghost cells of one local block overlap with the live cells of other local blocks grid, causing redundant computation.

For now, we apply the blocking parallelization to the `compute_flux()`, `limited_derivs()`, and `compute_step()` functions and leave parallelization of the other functions for future optimizations. Because the ghost cells of each block become contaminated with incorrect vectors and ghost cells of one block overlap with live cells of other blocks, it is important that we allocate per-thread memory for storing various vectors of the block separate from the *global* state. As such, we encapsulate this per-thread information in a class called `LocalState` that holds the vectors for the *local* state: `u_`, `f_`, `g_`, `ux_`, `uy_`, `fx_`, `gy_`, and `v_`. In fact, all of these vectors with the exception of `u_` are only used in the local state computation; we only need to keep a separate global state for `u_`. Figure 2 shows a visual representation of how the global grid is divided into local blocks. A vector of pointers to per-thread `LocalState` objects is kept as a member variable of `Central2D` called `locals_` that can be indexed by the thread ID.

To facilitate copying of vectors between the global and local states, we implement the `copy_to_local()` and `copy_from_local()` functions. Note that these functions can easily be parallelized since copying to the local state only has conflicting accesses on reads, and copying from the local state only writes the live cells in the global grid which do not overlap between local blocks. The ghost cells of the global grid are updated by the `apply_periodic()` function.

The pseudocode for our current parallelization is as follows:

```
while (t < tfinal) {
    apply_periodic();
    real dt = calculate_dt(compute_wave_speed());
    #pragma omp parallel num_threads(nthreads)
    {
        int tid = omp_get_thread_num();
        copy_to_local(tid);
        for (int io : sub_steps) {
            compute_flux(tid);
            limit_derivs(tid);
            compute_step(tid);
        }
        copy_from_local(tid);
    }
    t = update_time(dt);
}
```



**Figure 3: Performance Results of Parallel Implementation of Shallow Water Equation Solver Running on Compute Nodes** – Performance of the parallel implementation of the shallow water equation solver running on the Totient compute nodes compared against the provided serial implementation for all initial conditions. All speedups are the execution time normalized to the serial implementation. For the strong scaling experiment, we use the default 200x200 grid. For the weak scaling study, we increase the problem size at the same factor as the increasing number of threads (e.g., 200x200 grid for 1 thread, 400x400 grid for 4 threads, etc.).

The preliminary results comparing the performance of the serial and parallel implementations of the shallow water equation solver running the compute nodes is shown in Figure 3.

For the strong scaling study, we sweep the number of threads used for the parallel implementation using the same 200x200 grid size and normalize the execution time results against the serial implementation. In general, we see a pseudo-linear speedup with the number of threads until 6 threads, then the performance tapers off with any additional increase in parallelization. This is to be expected, especially with a smaller dataset, since the synchronization overhead begins to dominate the amount of work per thread with more threads. There is an interesting phenomenon for the performance of the 6 and 12 threads configurations in which the performance stops monotonically increasing and is inconsistent across different initial conditions. We believe the reason for this is two-fold. First, there is suboptimal load balancing for block dimensions that do not evenly divide the grid dimensions. Specifically, for a 200x200 grid, splitting the grid into 3 blocks in any dimension (e.g., 3x2 for 6 threads, 4x3 for 12 threads) will assign less work to the boundary blocks. Second, there might be an increase in cache conflicts for certain block dimensions. This effect would be most noticeable during the copying of data to and from the per-thread local state, which occurs after every timestep. The interplay between this effect and the fact that smaller blocks are more likely to fit in each thread's (i.e., core's) private L1 cache, could be a possible cause of the inconsistency we see for the performance of certain block dimensions.

For the weak scaling study, we increase the grid size with the same factor as the increase in the number of threads. For example, the 4 thread configuration uses a 400x400 grid, whereas the serial configuration uses a 200x200 grid. An ideal weak scaling parallel implementation would have a roughly constant speedup for increasing numbers of threads. These studies help programmers to identify the impact of synchronization overheads in their parallel implementations. In our results, we see an approximate exponential decay in performance as we scale the problem size with the number of threads, implying that our synchronization overhead does not scale linearly with the number of threads. However, it is important to keep in mind that this trend is likely partly due to the fact that increasing the grid dimensions increases the computational complexity by an order of  $N^3$ , rather than  $N^2$ . This is because we are not only increasing the total number of elements on the grid, but we are also increasing the total number of timesteps required to compute a single frame (i.e., requires more timesteps to reach convergence for a larger grid). Since we are increasing the number of timesteps, it is natural that the number of synchronization boundaries between timesteps increase as well.

### 3.2. Parallelizing for the Accelerator Board

There are several ways we can leverage the Intel Xeon Phi accelerator boards to further increase the speedups we see with the parallelized implementation for the compute nodes. One approach is to execute the code natively on the accelerator itself with no fine-grain offloading. Although this approach makes it easier to tailor the code for the accelerator and reduces overheads of copying data between the host and the device memories, we are forced to run all the computation from start to end on the accelerator. Another approach is to only offload certain sections of the computation to the accelerator. With this approach, there is a greater burden on the programmer to identify and parallelize the most compute-intensive section of the code, but allows him or her more flexibility in choosing which sections to accelerate.

As a starting point, we chose the second approach of offloading specific sections of the computation to the accelerator and build off of the parallel implementation for the compute nodes discussed in Section 3.1. In this naive first-pass, we offload the parallel section of the code and pass in the dimensions of the global grid and the blocks, as well as a pointer to the global grid itself. Inside the offloaded kernel, we still spawn off the specified number of threads normally, except that instead of using pre-allocated member `LocalState` objects, each thread creates its own `LocalState` object on the stack. The `copy_to_local()` and `copy_from_local()` functions were modified to copy the flattened elements of the global grid passed in by the host to the local grid vectors. Once the copy is complete, the other functions in the offloaded kernel can take a pointer to the per-thread `LocalState` object as an argument and access the vectors in

this object similar to before. Because this naive implementation requires us to offload computation before every timestep, there is an unnecessarily high overhead of transferring data between the host and device memories.

We continued to optimize the parallel implementation for the accelerator by offloading the entire `run()` function of the simulator. In this approach, one main thread on the accelerator is responsible for running the `apply_periodic()` and `compute_wave_speeds()` functions before parallelizing computation across multiple threads on the accelerator for the other functions mentioned above. The benefit of this approach is that we do not have to incur the overhead of copying data between the host and the accelerator memories for offloading the computation kernel for *every* timestep. Instead, we only pay this overhead once per frame in order to keep data local to the accelerator for as long as possible. Note that here we only need to copy the global grid vectors (i.e., `u_offload`) and the simulator parameters (e.g., `nx`, `nxblocks`, etc.) during the offload. Functions meant to be called from the accelerator are annotated with the `__declspec(target(mic))` attribute and must essentially be pure functions. In order to make it easier to pass in simulator parameters to these functions that normally rely on the simulator's member variables, we encapsulate the parameters copied to the accelerator in a separate `Parameters` class.

## 4. Tuning the Shallow Water Simulation

### 4.1. Vectorizing with AVX Extensions

Vectorizing with AVX Extensions can fall under two general categories:

1. Having the compiler auto-vectorize your code for you, and
2. Writing your own vectorized kernels for use as subroutines.

Both approaches have benefits and detriments. For example, when using the compiler to vectorize your code you save yourself the pain of having to reason about which register has what data, what SSE/AVX function callbacks wrap the appropriate instruction level code, etc. This provides great convenience, but does come at a cost. The compiler is only as smart as you let it be, and if you arrange your code improperly, you may not only slow down your program but you may very well yield the wrong result by accident. Writing your own vectorized code, on the other hand, can be quite difficult to get right. Often times it may be best to take a mixed approach, where you examine the instructions the compiler generated for a given code segment and determine whether or not you can refine this approach further.

#### 4.1.1. Auto-vectorization Using ICC

As discussed previously, you can use the `amplxe` tool to help with profiling, as well as use the intel compiler to generate an `optrpt` file describing what was / was not vectorized, how effective it was if vectorization occurred, and why vectorization did not occur if that were the case. On the note of when vectorization does not occur, we would like to mention a couple of things:

1. Not all loops are created equal

That is not every loop can be vectorized, either because its length cannot be known at compile-time, or because the data elements being accessed cannot be made to execute in parallel.

2. Understanding the codes in the `.optrpt` file can help you rearrange your code so that it is vectorized (recalling from 1 that not all of this is possible).



We found the following presentation extremely useful in rearranging loop statements:

<https://engineering.purdue.edu/milind/ece573/2011spring/lecture-14.pdf>

3. Introducing compile-time constants / `constexpr` members of a class can go a long way in assisting the compiler understand what can / cannot be vectorized.
4. Taking great care to enforce memory alignment as well as declare said alignment to the compiler will also enable it to vectorize even more.

We leave items 1 and 2 as an exercise for the reader ;) For item 3, we must first acknowledge that the purpose of the data type `std::array` is largely just for compilation hints, and we can wield this to our advantage. In the original implementation of `Shallow2d.h`, we had that

```
// Type parameters for solver
typedef float real;
typedef std::array<real,3> vec;
```

were the primary solver types used throughout the program. The issue, though, is that regardless of us declaring with `<real, 3>`, with high probability (given the architectures we are compiling on) this will get padded to 16 bytes regardless. The issue with this padding though, is that it is not guaranteed to be a `float` and treating it as such can potentially give problems. Recognizing this, we can modify this definition to be:

```
// Type parameters for solver
#define VEC_DIM 4 // change this and we all die...
typedef float real;
#ifdef __INTEL_COMPILER
    typedef __declspec(align(16)) std::array<real, VEC_DIM> vec;
#else // GCC
    typedef __attribute__((aligned(16))) std::array<real, VEC_DIM> vec;
#endif

// allow loop unrolling over 'vec'
static constexpr int vec_size = VEC_DIM;
static constexpr int VEC_ALIGN = 16;
```

Woah. Ugly. But necessary. What this does is ensure that we will have 4 floats per type `vec`, and depending on the compiler you are using also declares the alignment of this type. This stage may not be necessary. We also define two `static constexpr` members to allow us to use say `Physics::vec_size` as a compile time constant to allow us to unroll loops / `__assume_aligned` on memory. For example, the original computation of the corrector step in `Central2d.h` was:

```
// Corrector (finish the step)
for (int iy = nghost-io; iy < ny+ngghost-io; ++iy) {
    for (int ix = nghost-io; ix < nx+ngghost-io; ++ix) {
        for (int m = 0; m < v(ix, iy).size(); ++m) {
            v(ix, iy)[m] =
                0.2500 * ( u(ix, iy)[m] + u(ix+1, iy)[m] +
                          u(ix, iy+1)[m] + u(ix+1, iy+1)[m] ) -
                0.0625 * ( ux(ix+1, iy)[m] - ux(ix, iy)[m] +
                          ux(ix+1, iy+1)[m] - ux(ix, iy+1)[m] +
                          uy(ix, iy+1)[m] - uy(ix, iy)[m] +
                          uy(ix+1, iy+1)[m] - uy(ix+1, iy)[m] ) -
                dtcdx2 * ( f(ix+1, iy)[m] - f(ix, iy)[m] +
                          f(ix+1, iy+1)[m] - f(ix, iy+1)[m] ) -
                dtcdy2 * ( g(ix, iy+1)[m] - g(ix, iy)[m] +
                          g(ix+1, iy+1)[m] - g(ix+1, iy)[m] );
        }
    }
}
```

Though extremely verbose, we can now use these new additions to write a new loop:

```
// Corrector (finish the step)
for (int iy = nghost-io; iy < ny+ngghost-io; ++iy) {
    for (int ix = nghost-io; ix < nx+ngghost-io; ++ix) {
        /* Nomenclature:
        *   u_x0_y0 <- u(ix, iy)
        *   u_x1_y0 <- u(ix+1, iy)
        *   u_x0_y1 <- u(ix, iy+1)
        *   u_x1_y1 <- u(ix+1, iy+1)
        */
        // The final result
        real *v_ix_ey = v(ix, iy).data(); __assume_aligned(v_ix_ey, Physics::VEC_ALIGN);

        // grab u
```

The reasoning is that these traits now enable the intel compiler to do what it does best: vectorize like there ain't no tomorrow:

```

      LOOP END
      LOOP END
      LOOP END

```

The last stage in this phase, which we are uncertain if it is working correctly, is to create a custom allocator for `std::vector`. Unfortunately there is no byte aligned allocator in the standard yet, so we snagged somebody else's that seems to be popular on the interweb:

<https://gist.github.com/donny-dont/1471329>

We believe by virtue of the fact that it gives the right simulation that we are at least have a valid allocator, but honestly at this point we are very frustrated with how difficult it has been to use `std::vector`. After declaring

```
#define BYTE_ALIGN 64
#ifdef __INTEL_COMPILER
    typedef __declspec(align(BYTE_ALIGN)) std::vector<vec, aligned_allocator<vec, BYTE_ALIGN>> aligned_vector;
#else // GCC
    typedef __attribute__((aligned(BYTE_ALIGN))) std::vector<vec, aligned_allocator<vec, BYTE_ALIGN>> aligned_vector;
#endif
aligned_vector u_;           // Solution values
aligned_vector f_;           // Fluxes in x
aligned_vector g_;           // Fluxes in y
aligned_vector ux_;          // x differences of u
aligned_vector uy_;          // y differences of u
aligned_vector fx_;          // x differences of f
aligned_vector gy_;          // y differences of g
aligned_vector v_;           // Solution values at next step
```

it turns out that memory aligned allocators (in general) greatly conflict with other locations in the code that were previously vectorized. So although we were able to get some interesting vectorization results from the above, we may switch over to C-style arrays in the near future. The chaos of the above code, in particular the newly vectorized loop, do not seem worth the trade-off of how ugly / difficult to follow it is.

We may also explore other allocators e.g. using Eigen's aligned allocator, but generally do not see the benefit as we have probably spent more time trying to utilize these vectors through arcane trickery than we would have to just rewrite it based off of say `float *u` and use a memory aligned malloc. Time will tell whether we decide to keep these or not.

#### 4.1.2. Manual Vectorization

When it comes to writing custom kernels, we actually are very excited to do this. After much deliberation we have finally been able to get `#pragma offload target(mic)` to cooperate with things like `std::vector` and `std::array`.

Getting them working on the Phi's was as far as we got at this point, as discussed in section 3.2, so we hope to be able to play with this more and compare with the intel compiler's vectorization and *maybe* even beat it! If anything, we can just pull the same trickery we did in the corrector step described in the previous section.

## 4.2. Batching Multiple Timesteps

Another tuning optimization we implemented for the parallel implementations was batching multiple timesteps for each thread before having to synchronize. A single timestep of the computation (i.e., both even and odd sub-steps) requires 3 additional ghost cells in every dimension. This is because although each sub-step only requires 1 additional ghost cell, the staggered sub-step always computes the vector for the cells with a (-1,-1) offset from the normal sub-step. Coupled with the fact that we lose the fidelity of the data on the boundaries one additional cell per sub-step, we actually need an extra ghost cell on top of the 2 ghost cells to calculate the ghost cell required by the staggered sub-step. However, for every *additional* timestep we want to batch, we only need an extra 2 ghost cells on top of the initial 3 ghost cells for the first timestep due to the fact that the last layer of ghost cells is still accurate after the first timestep completes. As such, we modified the parallel implementation to allow a configurable number of batchable timesteps and adjust the number of ghost cells required accordingly.

The tradeoff here is that although we reduce the amount of synchronization required, we increase the amount of data each thread needs to hold in its local state. To a lesser extent, the latter also means that the amount of data we need to copy in `apply_periodic()` also increases. Another challenge with batching is synchronizing the `dt` across all threads. Because this value is calculated from the maximum velocities (i.e., `cx`, `cy`) across the *entire* grid and because this needs to be computed again for every timestep, we have a few options:

- Have each thread compute its local maxima and synchronize across threads to find the global maxima (defeats the purpose of batching);
- Have each thread compute its local maxima and run with different `dt` values (affects correctness of results);
- Calculate the `dt` once before each batch-step and use it for all timesteps in the batch for all threads;

Although it is conservative, we choose the third option, which is the only reasonable option not requiring a highly optimized local maxima sharing synchronization. One consequence of using this option with batching is that if the remaining time to be simulated is less than the time that will be simulated per batch-step, we could end up unnecessarily running multiple timesteps with a very small `dt` to finish the simulation. In order to address this issue, we dynamically adjust the number of timesteps in a batch (instead of the `dt` when we encounter this situation so that we always run the minimum number of timesteps required to simulate the specified amount of time.