

## CS 5220 – Project 2

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We did the domain decomposition for speed up and it works!

## Our work

There are three major optimization base on the original code.

1. Profiling
2. Parallelization
3. Tuning

## Profiling

We did some hotspot analysis of the serial code, and this is what we find out:

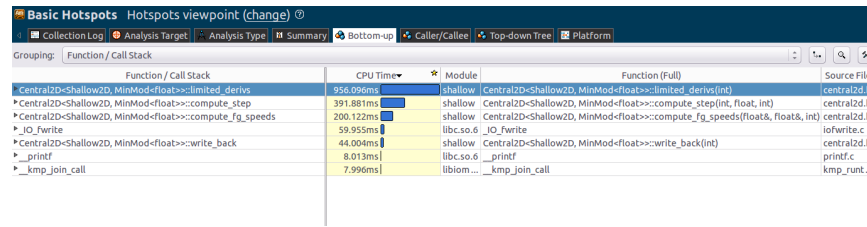


Figure 1: Hotspot analysis by ampxe-cl

We find that more 58% of time is spent on the function call "limited\_derivs". However, to our surprise, the function has been vectorized on assembly language level. We think since the data is not continuous in memory, there is a delay for acquiring data from memory to register. Probably that's why this function call is so slow. To further optimize this function call, we have to work on data dependence.

0x4036f5	197	vmovssl 0x8(%r8,%rsi,1), %xmm2
0x4036fc	197	vmovssl 0x14(%r8,%rsi,1), %xmm1
0x403772	197	vminss %xmm5, %xmm14, %xmm14
0x403782	197	vmovssl (%r11,%rsi,1), %xmm2
0x403788	197	vmovssl (%rbx,%rsi,1), %xmm1
0x4037fa	197	vminss %xmm4, %xmm14, %xmm14
0x403809	197	vmovssl 0x4(%r11,%rsi,1), %xmm2
0x403810	197	vmovssl 0x4(%rbx,%rsi,1), %xmm1
0x403882	197	vminss %xmm3, %xmm10, %xmm10
0x403892	197	vmovssl 0x8(%r11,%rsi,1), %xmm1
0x403899	197	vmovssl 0x8(%rbx,%rsi,1), %xmm0
0x40390d	197	vminss %xmm2, %xmm6, %xmm6
0x40391c	197	vmovssl (%r9,%rsi,1), %xmm10
0x403922	197	vmovssl (%r14,%rsi,1), %xmm9
0x403997	197	vminss %xmm1, %xmm3, %xmm3
0x4039a4	197	vmovssl 0x4(%r9,%rsi,1), %xmm2
0x4039ab	197	vmovssl 0x4(%r14,%rsi,1), %xmm1
0x403a23	197	vminss %xmm0, %xmm8, %xmm8
0x403a32	197	vmovssl 0x8(%r9,%rsi,1), %xmm1
0x403a39	197	vmovssl 0x8(%r14,%rsi,1), %xmm0
0x403aa9	197	vminss %xmm0, %xmm2, %xmm2

Figure 2: Assmby for hotspot

## Parallelization

We parallelized our code based on domain decomposition idea. The `-d` option specify how many time we split the domain along one direction.

In our code, each pocessor has its own domain of simulation with `nghost` layer of ghost cells. We maintain an extra data structure `sg_` for the whole simulation board. And there are three stages in each iteration of our parallel code:

- **Apply periodic:** In this stage, each thread copy the periodic boundary to their own ghost cells. After this stage, each one of those subdomains is ready for simulation.
- **Run step:** In this stage, each thread do simulation on their own subdomain, and the water height and water speed of each cell is updated.
- **Write back:** In this stage, every thread copy their updated subdomain back to the `sg_` data structure. There is a barrier after this stage to make sure before next iteration, the data needed by ghost cells is up-to-date.

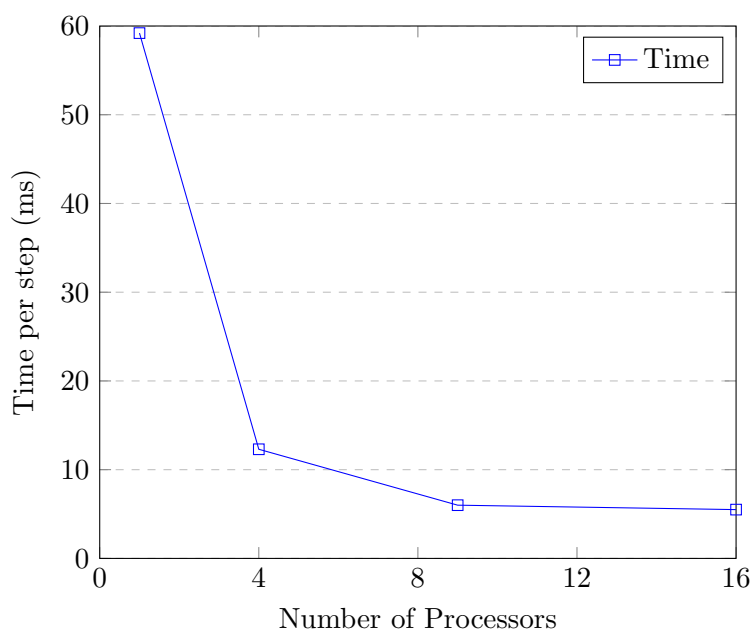


Figure 3: Strong Scaling

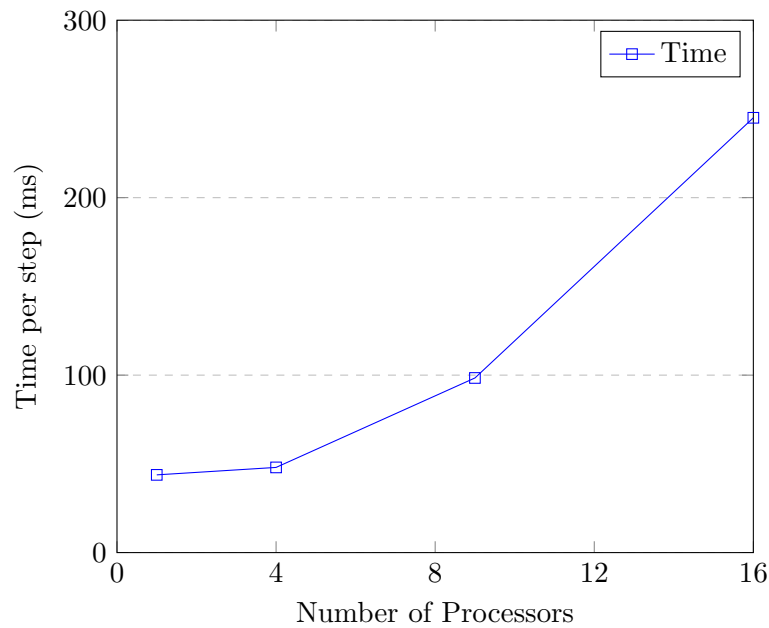


Figure 4: Weak Scaling

## Summary

Through this project:

- We analysed the hotspot of our code, yet we haven't done any tuning.
- Strong scaling for the parallel code is okay for the first three points, but the fourth point is bad.
- Weak scaling for the parallel code is okay for the first three points, but the fourth point is terrible.

In the future:

- We will try to cut down the time for calling `limited_diffvs` by tuning.
- We will use more layers of ghost cells and batch of steps simulations.
- We will further optimize the code by hotspot analysis.