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Abstract. Phase unwrapping is an important problem in the areas of optical metrology, synthetic aperture radar (SAR) image analysis, and magnetic resonance imaging (MRI) analysis. These images are becoming larger in size and, particularly, the availability and need for processing of SAR and MRI data have increased significantly with the acquisition of remote sensing data and the popularization of magnetic resonators in clinical diagnosis. Therefore, it is important to develop faster and accurate phase unwrapping algorithms. We propose a parallel multigrid algorithm of a phase unwrapping method named accumulation of residual maps, which builds on a serial algorithm that consists of the minimization of a cost function; minimization achieved by means of a serial Gauss–Seidel kind algorithm. Our algorithm also optimizes the original cost function, but unlike the original work, our algorithm is a parallel Jacobi class with alternated minimizations. This strategy is known as the chessboard type, where red pixels can be updated in parallel at same iteration since they are independent. Similarly, black pixels can be updated in parallel at same iteration. We present parallel implementations of our algorithm for different parallel multicore architecture such as CPU-multicore, Xeon Phi coprocessor, and Nvidia graphics processing unit. In all the cases, we obtain a superior performance of our parallel algorithm when compared with the original serial version. In addition, we present a detailed comparative performance of the developed parallel versions. © *2018 Society of Photo-Optical Instrumentation Engineers (SPIE)* [DOI: 10.1117/1.OE.57.4.043113]

Keywords: phase unwrapping; synthetic aperture radar interferograms; parallel computing; multicore CPU; Xeon Phi; graphics processing unit.

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1 Introduction

Phase unwrapping is an important problem in the areas of optical metrology, synthetic aperture radar (SAR) image analysis, and magnetic resonance imaging analysis. In this paper, the case study is developed for unwrapping large dimensions SAR interferograms^{1–3} using parallel computing.

An SAR signal contains amplitude and phase information. Amplitude is the strength of the radar response and phase is the fraction of one complete sine wave cycle.⁴ Therefore, the SAR interferogram is generated by two complex SAR images that observe the same area from slightly different look angles. This can be done with two radars mounted on the same platform or with a radar at different times by exploiting repeated orbits of the same satellite. Then, the interferogram is obtained by cross multiplying, pixel-by-pixel, the first SAR image $u_1 = |u_1|e^{i\phi_1}$ with the complex conjugate of the second $u_2^* = |u_2|e^{-i\phi_2}$.⁵⁻⁷ Thus, the interferogram amplitude is the amplitude of the first image multiplied by that of the second one $|u_1||u_2|$, whereas its interferometric phase is the phase difference between the images $\phi = \phi_1 - \phi_2$.

The relationship between the actual phase ϕ and the computed wrapped phase g can be mathematically expressed as

$$g_r = W\{\phi_r + \eta_r\},\tag{1}$$

where *r* indicates a pixel position in a regular lattice \mathcal{L} ; η_r represents noise and in the case of SAR phase it is correlated with the signal magnitude; finally,

$$W\{z\} \stackrel{\text{def}}{=} z + 2n\pi,\tag{2}$$

is the nonlinear wrapping operator, with *n* an integer such that $W\{z\} \in (-\pi, \pi]$. If the Nyquist criterion is violated

$$\phi_r + \eta_r - \phi_s - \eta_s \neq W(g_r - g_s), \tag{3}$$

with $s \in \mathcal{N}_r$ and $\mathcal{N}_r = \{s \in \mathcal{L} : ||r - s||_2 = 1\}$ the set of first neighbor pixels to the pixel *r*, then the unwrapping process needs to be implemented as the solution to an illposed inverse problem (i.e., a problem with many possible solutions) to estimate the unwrapped phase *f*. To this end, the efficient accumulation of residual maps (ARM) method was recently reported in Ref. 8, which unwraps phase maps with the additional advantage that can be implemented in parallel.

Parallel computing consists of the simultaneous execution of calculations, instructions, processes, or tasks using more than one processor. There are different architectures to

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achieve parallel computing; for example multicore computers, graphics processing units (GPUs), Xeon Phi coprocessors (XPCs), computer clusters, and field programmable gate arrays (FPGAs). According to each one of these architectures, there are different programming paradigms and languages, which exploit their capacities. Some of them are OpenMPI for clusters; OpenMP for multicore CPUs and XPCs; Cg, compute unified device architecture (CUDA), OpenCL, and OpenACC for GPUs and Xilinx tools for FPGAs. Furthermore, the architectures can be merged to improve the performance of some problems. It is possible to have a multicore computer with one or more GPUs, or to have a cluster with one or more GPUs and FPGAs in each node,⁹ or a grid of GPUs, etc.

This work was planned to present a comparison as fair as possible between parallel implementations of the ARM method using a multicore CPU, XPC and GPU, to help implementers to take a decision in future parallel implementations of phase unwrapping algorithms that are being developed. In the state of the art, it is common to find this type of comparisons using implementations of simple algorithms of linear algebra methods (vector and matrix multiplications, factorizations), transformations (Fourier transform), or interpolations (bilinear, bicubic, and splines). These implementations are useful as benchmark tests but are far from reflecting the actual working conditions of current image analysis methods. Our study is complementary to such tests and throws another type of conclusions, more focused on the implementation of image analysis algorithms. For this reason, we are using a complex algorithm of the state of the art (multigrid phase unwrapping) implemented in different frameworks and we evaluate them in production conditions. The evaluated frameworks are more similar to the actual working conditions. These frameworks are summarized in Table 1. Framework 1 consists of using MATLAB with OpenMP directives into a mex-C file to execute the program in the multicore CPU or XPC, and using CUDA kernel integration in MATLAB (without mex-C) to process in the GPU. Framework 2 consists of using C/C++ with OpenMP directives to execute the program in the multicore CPU or in the XPC, and with CUDA kernel functions to execute the program in a GPU. The hardware architectures we are not considering are computer clusters and FPGAs. Our reasons are that computer clusters seem not to be proper for cellular processes with high messaging interchange (the kind of processing often used in image processing), and FPGA requires of specific and dedicated hardware.

The remaining sections of this paper are organized as follows: Sec. 2 gives a brief review of the serial ARM method. Section 3 describes the parallel implementation of the method and gives the implementation details of the most

Table 1 Parallel implementation frameworks.

Architecture	Framework 1 (MATLAB)	Framework 2 (C/C++)
Multicore CPU or XPC	OpenMP with mex-C file	OpenMP
GPU	CUDA kernel integration in MATLAB	CUDA kernels

demanding process. Sections 4 and 5 present a comparison of the processing time and speedup between the multicore CPU, XPC, and GPU architectures, using simulated and real data, respectively. Section 5 also discusses the wrapped phase generation from two SAR images. Finally, Sec. 6 gives our remarks and conclusions.

2 Serial Implementation of Accumulation of Residual Maps Method

The ARM method consists of an incremental scheme for unwrapping the wrapped phase g. Let

$$\rho_{rs} \stackrel{\text{def}}{=} W(g_r - g_s),\tag{4}$$

be the wrapped first differences of the wrapped phase and $f^{(k)}$ be the current estimate of the unwrapped phase. Then, we can compute the current residual wrapped differences as

$$\rho_{rs}^{(k)\text{def}} W\{\rho_{rs} - f_r^{(k)} + f_s^{(k)}\}.$$
(5)

If the residual differences field $\rho_{rs}^{(k)} \neq 0$, then one could try to estimate an update field $\delta^{(k)}$ such that

$$\sum_{(r,s)\in\mathcal{L}} W\{\rho_{rs}^{(k)} - \delta_r^{(k)} + \delta_s^{(k)}\}^2 \le \sum_{(r,s)\in\mathcal{L}} [\rho_{rs}^{(k)}]^2.$$
(6)

In any case, the field δ will compensate only the conservative component of the residual differences field ρ_{rs} ; i.e., monopoles in the sense of wrapped phase could not be solved.¹⁰⁻¹³

In the ARM algorithm, the current phase is updated (accumulated) with

$$f_r^{(k+1)} = f_r^{(k)} + \delta_r^{*(k)},\tag{7}$$

the residual wrapped differences that are computed with Eq. (5), and the update field $\delta^{*(k)}$ is computed by minimizing the half-quadratic cost function

$$\begin{split} \delta^{*(k)}, \omega^{*(k)} &= \arg\min_{\delta^{(k)}, \omega^{(k)}} U[\delta^{(k)}, \omega^{(k)}; \rho^{(k)}] \\ &= \frac{1}{2} \sum_{r \in \mathcal{L}} \sum_{s \in \mathcal{N}_r} ([\omega_{rs}^{(k)}]^2 \{ [\rho_{rs}^{(k)} - \delta_r^{(k)} + \delta_s^{(k)}]^2 \\ &+ \lambda [\delta_s^{(k)} - \delta_r^{(k)}]^2 \} + \mu [1 - \omega_{rs}^{(k)}]^2), \end{split}$$
(8)

where λ and μ are the positive parameters of the algorithm (see Ref. 8 for a deeper discussion on their selection). The data term in Eq. (8) is a weighted version of the regularized least square potential.^{13–15} The second term {membrane potential: $[\delta_s^{(k)} - \delta_r^{(k)}]^2$ } penalizes large local variations on the unwrapped phase,^{16,17} which reduces noise.

The solution to Eq. (8) can be computed by alternating minimization with respect to $\delta^{(k)}$ and $\omega^{(k)}$. Thus, if $\omega^{(k)}$ is fixed, then the solution of the positive-definite diagonal-dominant linear system given by $\partial U/\partial \delta_r^{(k)} = 0$ can be computed with a Gauss–Seidel (GS) iterative scheme

$$\delta_r^{*(k)} = \frac{\sum_{s \in \mathcal{N}_r} [\omega_{rs}^{(k)}]^2 [\rho_{rs}^{(k)} + \delta_s^{(k)} (1+\lambda)]}{\sum_{s \in \mathcal{N}_r} [\omega_{rs}^{(k)}]^2 [1+\lambda]},\tag{9}$$

where $\delta_s^{(k)}$ (for $s \in \mathcal{N}_r$) are the first neighbor values to the $\delta_r^{(k)}$. In the case of a Jacobi update scheme, $\delta_s^{(k)} \equiv \delta_s^{(k-1)}$. In the GS scheme, the updated values are used; i.e., in a pixel scanning top-down/left-right, the left and upper pixels are at iteration k and the right and down pixels are at iteration k - 1. Our parallel implementation uses the red-black update rule, explained in Sec. 3. Similarly, from $\partial U/\partial \omega_{rs}^{(k)} = 0$, we obtain the closed equation:

$$\omega_{rs}^{*(k)} = \frac{\mu}{\mu + [\rho_{rs}^{(k)} - \delta_r^{(k)} + \delta_s^{(k)}]^2 + \lambda [\delta_s^{(k)} - \delta_r^{(k)}]^2}.$$
 (10)

It is well known that GS is prone to having a slow reduction of low-frequency residuals. Thus, the convergence is accelerated using a multigrid strategy;¹⁸ this strategy is summarized in Algorithm 1. It shows a simple multigrid scheme where the solution at the level N is used as initial guess for level N - 1. This proposal implements a full-nested multigrid strategy detailed in Algorithm 2. In the implementation of Algorithm 2, the computation of residual wrapped phase at the end of each iteration is required. The residual wrapped phase between two given phases q and f is denoted as

residual
$$(g, f) \stackrel{\text{def}}{=} W(g - f).$$
 (11)

Algorithm 1 Multigrid unwrapping. Multigrid strategy with N scale levels.

1: **function** MULTIGRIDUNWRAP $(g, f, \lambda, \mu, N, T)$,

2: if N > 0 then,

- 3: $\hat{g} = \text{DOWNSAMPLE}(g); \triangleright \text{Down sampling}$
- $\hat{f} = \text{DOWNSAMPLE}(f);$ 4:
- \hat{f} =MULTIGRIDUNWRAP ($\hat{g}, \hat{f}, \lambda, \mu, N 1, T$); \triangleright Change level, 5:
- $f = \text{UPSAMPLE}(\hat{f}); \triangleright \text{Up sampling},$ 6:
- 7: end if
- 8: //Unwrap current level,
- 9: Set $\delta = 0, \forall r$,

Set $\omega_{rs} = 1$, $\forall (r, s)$; $\triangleright A$ different initial value for ω can be used, 10:

Compute $\rho_{rs}^{(k)}$, $\forall (r, s)$, with Eq. (5) 11:

- for t = 1, 2, ..., T do, 12:
- 13: Update δ , keeping fixed ω , with Eq. (9),
- 14: Update ω , keeping fixed δ , with Eq. (10),
- end for 15:
- 16: $f = f + \delta$,
- 17: end function
- 18: return f.

Algorithm 2 Nested Multigrid Unwrapping. Nested multigrid strategy for Algorithm 1 with N iterations of N scale levels.

1: function NESTMULTIGRIDUNWRAP (g, λ, μ, N, T)

- 2: $f = 0; N_0 = N$
- 3: while N > 0 do
- 4: $f' = MULTIGRIDUNWRAP (g, \lambda, \mu, N_0, T); \triangleright Simple multigrid$
- 5: f = f + f':
- 6: $g = \text{residual}(g, f); \triangleright \text{Residual wrapped phase},$
- 7: N = N - 1:
- 8: end while
- 9: return f + q;
- 10: end function

To implement this function, one can use the equation

$$\operatorname{residual}(g, f) = \operatorname{atan2}[\sin(g - f), \cos(g - f)]. \tag{12}$$

The serial ARM method was implemented in MATLAB using mex-files and it is available in Ref. 19.

3 Parallel Implementation

In our parallel implementation, we develop two frameworks (see Table 1). Framework 1 uses MATLAB and improves the processing time of Eqs. (9) and (10). We include OpenMP directives into the mex-C file to execute the heavy work in the multicore CPU or in the XPC. In contrast, we use the CUDA kernel integration in MATLAB (without mex-C) to process in the GPU. Framework 2 uses C/C++ and improves the processing time of the whole method. Then, the Algorithms 1 and 2 are translated from MATLAB scripts to C/C++ files, this framework uses only OpenCV for reading and writing input images. With the whole code in C/C++, we can include OpenMP directives to execute the program in the multicore CPU or in the XPC, and we can create CUDA kernel functions to execute the program in a GPU. An implementation of the proposed frameworks can be found in the following code ocean capsules: https://codeocean.com/2018/04/ 09/comparison-of-multi-hardware-parallel-implementationsfor-a-phase-unwrapping-algorithm/ and https://codeocean. com/2018/04/23/comparison-of-multi-hardware-parallelimplementations-for-a-phase-unwrapping-algorithm/.

From the serial ARM method, we note that the solver for Eqs. (9) and (10) is the computationally heaviest part, then we parallelize these equations as a first approach. Note that, for updating ω in Eq. (10) as well as the sentences in the Algorithms 1 and 2, we need only pixel-wise operations, applying an independent parallelism model,²⁰ whereas for updating δ in Eq. (9), we need a strategy to parallelize the GS method because the communication between a pixel and its neighborhoods. The following subsections describe the GS method and the implementation details, focusing on the parallelization of Eq. (9).

3.1 Gauss-Seidel Method

The GS method is one of the basic iterative methods for solving linear systems.²¹ In our particular case, we want

Algorithm 3 GS with natural ordering. Computes δ ; T is the maximum iteration number; cols and rows are the height and width of the image, respectively,

1: function GS $(\delta, \omega, \tilde{\rho}, \lambda, T)$

2: for t = 1, 2, ..., T do

- 3: **for** r = (1,1) to (cols, rows) **do**
- 4: Update δ_r with Eq. (9);
- 5: end for
- 6: end for
- 7: return δ ;
- 8: end function

to compute Eq. (9) with a GS iterative scheme. An advantage of GS is that it is only necessary one array to allocate and update the values of δ_r . In contrast, depending on the order in which we loop the grid pixels, we will get different implementations of the GS method.

We implement our algorithm taking into account two ordering ways such as the natural ordering (see Algorithm 3) and the red-black ordering (see Algorithm 4). Note that in both algorithms the for loops run over all image pixels rper row. The difference is that in the red-black ordering the pixels are considered red and black following a chessboard pattern. We consider a pixel r = (i, j) red if i + jis even and black if i + j is odd. Then, when the red pixels are updated in the first for loop, they only need the black pixel values and vice versa in the second loop. With the red-black ordering, we can implement the algorithm in parallel using a multicore CPU, XPC, or GPU.

3.2 Implementation Details

In both frameworks, we use the OpenMP and CUDA languages to parallelize the code. In the following, we describe the directives and functions created to parallelize Eq. (9).

OpenMP is an API for shared-memory parallel programming.²² The "M" in OpenMP stands for "multiprocessing," a term that is synonymous with shared-memory parallel computing. Thus, OpenMP is designed for systems in which each thread or process can potentially have access to all available memory. OpenMP provides a set of pragma directives that are used to specify parallel regions, manage threads inside parallel regions, and distribute for loops in parallel, among other things.

Note that, the internal for loops of Algorithm 4 can be parallelized with OpenMP directives as follows:

```
omp_set_num_threads(NUM_THREADS);
#pragma omp parallel
{
    #pragma omp for private ()
    for (:,:,:){
    }
}
```

```
Algorithm 4 GS with red–black ordering. Computes \delta; T is the maximum iteration number; cols and rows are the height and width of the image, respectively;
```

1: function GS RED–BLACK ($\delta, \omega, \tilde{\rho}, \lambda, T$)

- 2: for t = 1, 2, ..., T do
- 3: for r = (1,1) to (cols, rows) that are red do
- 4: Update δ_r with Eq. (9);
- 5: end for
- 6: for r = (1,1) to (cols, rows) that are black do
- 7: Update δ_r with Eq. (9);
- 8: end for
- 9: end for
- 10: **return** δ;
- 11: end function

where NUM_THREADS is the thread number to launch. #pragma omp parallel opens a parallel region and #pragma omp *for* parallelizes the for loop. In shared-memory programs, the individual threads can have private and shared memories. Communication is usually done through shared variables. Inside the #pragma omp *for*, all variables are shared by default for all threads, then, to declare shared and private variables we use only the private directive, to declare the variables that are private for each thread launched. In this way, we can process a mex-C or a C/C++ code in a multicore CPU.

Intel XPC, also known as Intel many integrated core architecture (or Intel MIC), is a coprocessor computer architecture that offers additional power-efficient scaling, vector support, and local memory bandwidth while maintaining the programmability and support associated with Intel Xeon processors.²³ The XPC runs Linux and has its own IP address. Then, we can log onto the XPC in a terminal window. There are two programming models for computing in an XPC, the native and the offload model. In the native model, the application runs in the coprocessor, whereas in the offload model, the application runs a main host program in the CPU and offloads work to the coprocessor. We use the offload model, because our application can run the main program from MATLAB or OpenCV and offloads the heavy work in the coprocessor through mex-C file or the C/C++ code. Moreover, the XPC provides a wide interoperability with OpenMP, bringing a set of specialized directives; thus, we add the following at Algorithm 4

```
#pragma offload target (mic:0) inout() in ()
{
    omp_set_num_threads (NUM_THREADS);
    for (t=0;t<T;t++){</pre>
```

```
#pragma omp parallel
{
    #pragma omp for private ()
```

```
for (:,:,:){// for red pixels
}
#pragma omp for private ()
for(:,:,:){//for black pixels
}
}
```

}

where #pragma offload target(mic:0) opens a code region, which will run in the XPC. Note that we use the attribute inout for including variables that will be transferred from CPU to XPC memory and vice versa. Also, we use the attribute *in* for including variables that will only be transferred from CPU to XPC. The rest are pragmas of OpenMP to parallelize the internal for loops. These pragma directives can be ignored, and the program should simply work in a nonparallel mode (sequential). When a compiler recognizes OpenMP directives (requires the OpenMP switch on the Intel compilers), then the directives are interpreted to give direction on how to create parallel tasks to speed the execution of a program through parallelism.

In contrast, CUDA is a language that contains a set of instructions for processing in a GPU. The advantage of using a GPU is that it contains multiple transistors for the arithmetic logic unit, based on the single instruction and multiple threads programming model, which is exploited when multiple data are managed from one simple instruction in parallel, similar to single instruction multiple data (SIMD) model.^{24,25} In framework 1, we use the CUDA kernel integration in MATLAB.²⁶ Then, we create an executable kernel from CU or PTX (parallel thread execution) files, and run that kernel on a GPU from MATLAB. The kernel is represented in MATLAB by a CUDAKernel object, which can operate on MATLAB array or gpuArray variables. We implement the kernel of Algorithm 4 as follows:

%Copy memory from CPU to GPU

delta_dev=gpuArray (double(delta));

omega_dev=gpuArray (double(omega));

%Create the kernel functions

GS_kernel=parallel.gpu.CUDAKernel('solverGS.ptx',' solverGS.cu');

Omega_kernel=parallel.gpu.CUDAKernel('Omega. ptx',' Omega.cu');

%Size of Grid and Thread Blocks

blocksize_x=32;%Fix this parameter according

blocksize_y=32; %to GPU capabilities

grid_x=ceil (rows/blocksize_x);

grid_y=ceil (cols/blocksize_y);

solverGS_kernel. ThreadBlockSize=[blocksize_x, blocksize_y,1];

solverGS_kernel. GridSize=[grid_x,grid_y];

Omega_kernel. ThreadBlockSize=[blocksize_x, blocksize_y,1];

Omega_kernel. GridSize=[grid_x,grid_y];

%kernel executions

```
for t=1:niter
```

- [delta_dev]=**feval** (GS_kernel,delta_dev,omega_dev, ...,0); %*Red-* ->0
- [delta_dev]=**feval** (GS_kernel,delta_dev,omega_dev, ...,1); %*Black-->1*
- [omega_dev]=**feval** (Omega_kernel,omega_dev, delta_dev,...);

end

%Copy memory from GPU to CPU

delta=gather(delta_dev);

Note that in framework 1, using any device (an XPC or a GPU), the number of times that the program needs to transfer memory between CPU and the device is $N \times N$, where N is the number of levels for both the multigrid Algorithm 1 and the nested Algorithm 2. To reduce this number of transferences, we develop framework 2, where we translate the MATLAB script to C/C++ code. In this framework, we can transfer the memory from Algorithm 2 just once.

With the implementations above described, we can run our program in a multicore CPU, XPC, or GPU. In the following section, we present some experiments and results comparing the execution time in such architectures.

4 Synthetic Experimental Evaluation

The experiments were executed on two servers, the first one (server K20) is a server with Intel(R) Xeon(R) CPU E5-2620 v2 2.10 GHz, Ubuntu 14.04 (64 bits), 24 hyperthreading cores, a XPC 3120A 1.1-GHz with 57 cores and 6-GB RAM, and a video card Tesla K20 with 4-GB RAM. The second one (server K40) is a server with Intel(R) Xeon(R) CPU E5-2690 v2 3.00 GHz, Ubuntu 14.04 (64-bits), 20 physical cores, a XPC 3120A 1.1 GHz with 57 cores and 6-GB RAM, and a video card Tesla K40 with 12-GB RAM.

Figure 1 shows a synthetic wrapped phase map with a size of 512×512 pixels, which we use in our experiments. This synthetic wrapped phase map was used in Ref. 8 to compare the ARM method with different state-of-the-art methods. The second and third rows of the Fig. 1 show the rewrapped phase from the unwrapped phase of the ARM method, in serial and parallel versions, respectively, using different numbers of iterations T, N = 5 levels of nested and multigrid algorithms, $\lambda = 0.1$ and $\mu = \pi/10$. With T = 2000, we note that the rewrapped phase in the serial and parallel versions converges to the same result; thus, in the rest of the synthetic experiments we fix T = 2000 iterations.

Tables 2 and 3 show a comparative performance of our two respective frameworks of parallel ARM method for server K20, on the other hand, the Table 4 shows a comparative performance of the parallel ARM method framework 2 for server K40. These tables show processing times using the double-precision floating-point format in different architectures (multicore CPU, XPC, and GPU) and different image sizes (512×512 , 1024×1024 , 2048×2048 , and 4096×4096). Processing time is measured in seconds from the start of Algorithm 2 until this is finalized, which means that we are considering both the processor work and the memory transfer time.

With a size of image $\leq 512^2$, the best processing time is obtained using the multicore CPU of server K20 (see

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Fig. 1 Rewrapped phases from serial (second row) and parallel (third row) versions at different numbers of iterations *T*. (a) Wrapped synthetic phase, (b) T = 250, (c) T = 500, (d) T = 1000, and (e) T = 2000.

Table 2 Processing time in seconds of our parallel ARM method framework 1 in the server K20. The smallest time per row is shown in bold.

Size of image	Serial 1 <i>C</i>	Multicore CPU						XPC			
		2 <i>C</i>	4 <i>C</i>	8 <i>C</i>	12 <i>C</i>	16 <i>C</i>	20 <i>C</i>	112 <i>C</i>	224 <i>C</i>	GPU	
512 ²	36.5	19.3	11.2	6.8	5.7	10.1	7.1	17.9	24.6	10.8	
1024 ²	144.6	80.3	43.3	30.8	26.7	26.9	28.6	50.5	88.8	21.9	
2048 ²	814.3	449.6	249.1	154.6	155.8	158.4	150.1	316.2	360.7	66.9	
4096 ²	3529.9	1937.8	1067.9	616.3	604.6	779.3	700.3	1206.9	1291.1	248.4	

Table 3	Processing time in s	econds of our parallel A	ARM method framework 2 in th	ne server K20.	The smallest time per r	ow is shown in bold.
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Size of image	Serial 1 <i>C</i>		Multicore CPU						XPC	
		2 <i>C</i>	4 <i>C</i>	8 <i>C</i>	12 <i>C</i>	16 <i>C</i>	20 <i>C</i>	112 <i>C</i>	224 <i>C</i>	GPU
512 ²	33.9	20.4	10.6	5.7	5.5	7.2	5.3	30.1	32.0	6.04
1024 ²	136.0	81.8	49.3	27.4	22.9	23.5	19.9	60.3	60.6	11.2
2048 ²	660.8	360.6	282.0	97.8	122.5	121.4	127.1	449.8	410.6	34.2
4096 ²	2815.1	1452.6	775.0	411.6	603.7	514.6	499.3	3210.5	3192.5	123.3

Tables 2 and 3), whereas with a size of $>= 1024^2$ the best time is obtained using the GPU. Server K20 has 24 hyperthreading cores, but physically it has two sockets with 6 cores per socket, what means the server has 12 physical cores and 24 logical cores. We can see that the best performance is obtained with 12 and 8 cores. In contrast, the server K40 has 20 cores (20C) without hyperthreading. Note that the best times for the multicore CPU were obtained when we fixed the NUM_THREADS to 20C (see Table 4). When the hyperthreading is enabled, each physical core is

Size of image	Serial 1 <i>C</i>	Multicore CPU						XPC			
		2 <i>C</i>	4 <i>C</i>	8 <i>C</i>	20 <i>C</i>	24 <i>C</i>	28 <i>C</i>	112 <i>C</i>	224 <i>C</i>	GPU	
512 ²	26.3	15.3	7.9	4.1	2.2	3.4	3.5	9.5	10.6	2.5	
1024 ²	107.2	62.4	31.5	16.2	7.1	12.7	11.5	24.8	23.0	7.2	
2048 ²	515.7	281.9	146.7	67.4	50.5	72.4	69.4	130.7	100.6	25.7	
4096 ²	2193.8	1047.1	536.6	273.2	208.8	285.4	273.0	716.9	636.2	99.6	

Table 4 Processing time in seconds of our parallel ARM method framework 2 in the server K40. The smallest time per row is shown in bold.

divided in two logical cores, sharing resources such as the instruction pointers, integer and floating point registers, scheduling queues, caches, and execution units. The performance of a parallel program using hyperthreading declines if a logical core monopolizes some critical resource as the floating point registers or the caches. The purpose of parallel programming is increased performance, which is fundamentally a program optimization problem. As the memory hierarchies are radically different between platforms and the connection of cores on a single processor varies widely, this optimization problem is complicated.²⁷ In our implementation, we can see that it is better to adjust NUM_THREADS with the number of physical cores (N_{pc}) of CPU than with the number of logical cores, to obtain good performance.

We observe that the results of the XPC in server K20 are better than the serial program and comparable with the 2Cmulticore option, but these are overcome by the other architectures. One might think that the bad performance of the XPC in Table 2 is due to the memory transfers as the case of the GPU, but, in Table 3, we can see that even with the reduction of memory transfers, the XPC has the worst times. However, in Table 4, we can see an improvement in the processing time of the XPC; this is due to the directive #pragma omp simd that we have included in the OpenMP code. This directive transforms a loop into a loop that will be executed concurrently using SIMD instructions to help with the vectorization inside the XPC.²³ Vectorization refers to applying a single instruction to a group of data items of the same data type (or vector) that can be processed at once. On the XPC, the vector processing unit supports 512-bit vector width, our implementation uses double-precision floatingpoint format numbers; thus, eight values can be processed simultaneously.

With respect to the GPU, we can see that the memory transfers impact its performance (see Tables 2 and 3). Note that between the GPUs K20 and K40 (see Tables 3 and 4), there is a little difference in the processing time, which is due to the differences in the clock speed of both servers and the CUDA cores, 2688 of K20 and 2880 of K40.

The speedup of a parallel program can be defined as

$$S = \frac{T_s}{T_p},\tag{13}$$

where T_s is the processing time of a serial program and T_p is the processing time of a parallel program.²² Figures 2 and 3 show the speedups of the two frameworks of parallel the ARM method in server K20, and Fig. 4 shows the speedups of framework 2 in server K40. In the case of multicore CPU, Figs. 2 and 3 show that *S* increases when we use from 2*C* to 8*C* or 12*C*, while this decreases when we use 16*C* and 20*C*. In Fig. 4, we can see an increase in *S* until N_{pc} , then with NUM_THREADS > N_{pc} , *S* decreases and it is comparable



Fig. 2 Evaluation of speedups of parallel ARM method framework 1 in server K20, using different architectures (multicore CPU from 2*C* to 20*C*, XPC with 112*C* and 224*C*, and GPU K20) and different image sizes.



Fig. 3 Evaluation of speedups of parallel ARM method framework 2 in server K20, using different architectures (multicore CPU from 2*C* to 20*C*, XPC with 112*C* and 224*C*, and GPU K20) and different image sizes.



Fig. 4 Evaluation of speedups of parallel ARM method framework 2 in server K40, using different architectures (multicore CPU from 2*C* to 28*C*, XPC with 112*C* and 224*C*, and GPU K40) and different image sizes.

with the speedup of 8*C*. Next, in the case of the XPC, we can see that *S* is almost the same that 2*C* multicore in Figs. 2 and 3, whereas, in Fig. 4, the speedup of XPC is over the speedup of 4*C* multicore, this due to the included simd directive. Note that in the case of the GPU, *S* has notable differences between the different sizes of processed images. Furthermore, *S* is higher in framework 2 than framework 1, reaching a speedup of approx. $23 \times$ for an image of 4096 × 4096 pixels.

5 Phase Unwrapping for Large Synthetic Aperture Radar Interferograms

Before introducing the conducted experiments with large SAR inteferograms, we show the generation of interferometric synthetic aperture radar (InSAR) for RADARSAT-2, which is a Canadian satellite system equipped with a powerful SAR instrument. It offers powerful technical advancements that enhance marine surveillance, environmental monitoring, resource management, and mapping around the world,. From this process, we obtain the wrapped phase that will be later unwrapped with the ARM method.

5.1 Interferometric Synthetic Aperture Radar Formation

InSAR exploits the phase difference between two single look complex (SLC) images acquired over the same area from slightly different sensor positions.⁷ Figure 5 shows the processing chain for InSAR formation. The block diagram can be applied for SLC images using the opensource software called sentinel application platform (SNAP)²⁸ as follows

• Data coregistration: To create the interferogram, the two SLC images must be coregistered into a stack. One image is selected as the master or reference and the other image is the slave. The pixels in the slave image will be moved to align with the reference image to subpixel accuracy. This ensures that each ground target contributes to the same pixel in both the master and slave images.



Fig. 5 InSAR formation as applied to RADARSAT-2 SLC products.

- Interferogram formation: Once the images have been coregistered, the inteferogram is formed by cross multiplying the master image $u_m = |u_m|e^{i\phi_m}$ with the complex conjugate of the slave image $u_s^* = |u_s|e^{-i\phi_s}$. The amplitude of both images is computed by $|u_m||u_s|$, whereas the phase represents the phase difference between the two images $\phi = \phi_m \phi_s$.
- Topographic removal: The phase difference can have contributions from the Earth phase, which is the phase contribution due to the Earth curvature. In the interferometric processing, the flat Earth phase is removed to eliminate sources of error, to be left with only the contributor of interest, which is typically the elevation or the displacement. Therefore, the interferogram is flattened after removing the topographic phase.
- Phase filtering: Inteferometric phase is normally corrupted with noise; therefore, to properly unwrap the phase, the signal-to-noise ratio needs to be increased by filtering it.

5.2 Unwrapping Synthetic Aperture Radar Interferograms

For unwrapping SAR interferograms, we apply our ARM method framework 2 in server K40, with $\lambda = 10$ and $\mu = 100$. In Fig. 6, panel (a) shows a wrapped SAR phase taken from a tutorial available in Ref. 29, which has a size of 561×1591 pixels. We apply our method with N = 3 and T = 4000. Panels (b) and (c) show the unwrapping and rewrapped phase, and panel (d) shows a



Fig. 6 Unwrapped phase from SAR interferogram taken from Ref. 29. (a) Wrapped phase, (b) unwrapped phase, (c) rewrapped phase, and (d) mesh of unwrapped phase.

mesh of the unwrapped phase, where we can see the shape of a volcano. The processing time is ~105.2 s using the serial code, 17.7 s using the XPC (a speedup of ~6×), 4.5 s using OpenMP with 20C (a speedup of ~23×), and 7.5 s using the GPU (a speedup of ~14×).

In contrast, the panel (a) of Fig. 7 shows a huge wrapped SAR phase with size of 8000×10500 pixels, which we create from two SLC RADARSAT-2 images using SNAP and following the steps above described for InSAR formation. The images were acquired on May 4, 2008 and May 28, 2008, over the Phoenix area. Here, we are interested in full-resolution SAR inteferograms, but a subset around a particular area in which you are interested can be created to reduce the amount of processing needed. For unwrapping the wrapped SAR phase, we fix N = 7 and T = 1000. Panels (b) and (c) show the unwrapped and rewrapped phase, and panels (d) and (e) show the unwrapped phase and mesh of the region bounded by the red rectangle of size of 801×2001 pixels. The processing time is ~6216.8 s using the serial code, 739.7 s using OpenMP with 20C (a speedup of $\sim 8\times$), and 397.4 s using the GPU (a speedup of $\sim 16 \times$). In this experiment, the memory of the XPC was insufficient.

6 Remarks and Conclusions

In this work, different parallel architectures were used, comparing its processing time for solving a phase unwrapping problem. We translate the whole code from MATLAB to



Fig. 7 Unwrapped phase from our SAR interferogram taken over Phoenix area. (a) SAR wrapped phase, (b) unwrapped phase, (c) rewrapped phase, (d) a region (red rectangle) of the unwrapped phase, and (e) mesh of unwrapped phase region (red rectangle).

C/C++ to parallelize the method and improve the speedup, avoiding several copies of memory between CPU-RAM and the coprocessors XPC or GPU RAM. For our parallel implementation using a multicore CPU, it is important to know the number of physical cores, because with this number, we obtain the highest speedup between the possible configurations for launching threads in the CPU. In the case of XPC, it is notable that its speedup can be improved if we consider the vectorization; however, it is very poor compared with the rest. Although it is easier to translate a serial program to parallel using OpenMP than CUDA, we can see that the GPU remains a powerful tool, with which it is possible to obtain high speedup values above $15\times$, considering image sizes greater than 1024² pixels such as SAR interferograms, and thus it is worthwhile to invest time in to translate the code using CUDA.

As future work, we think that another interesting comparison could be using OpenCL, to create a kernel function, which can be offloaded in the different architectures here used. We want to improve the performance of our GPU implementation using texture and unified memory. Moreover, we want to compare the ARM method with another one (such as Snaphu³⁰) that is used to process SAR interferograms.

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