

Figure 1: 4 process decomposition of a 22x16 stencil. Process labels start in the top left and proceed clockwise. Local variables of state are also illustrated for indexing a 2D array, $0.1, B_x^1$, etc.

1 Basic Terminology And Motivation

The solution of partial differential equations (PDEs) is a well known problem in computing addressed by many numerical methods. To solve a PDE numerically, a method must choose a discrete form of its equation. A key characteristic of the discrete form chosen is its *stencil*. A stencil defines a dependency structure that constrains the execution of the PDE solution method. The dynamics of this dependency structure is the topic of this section. These dynamics further define a central performance bottleneck in solving PDEs on parallel processors which is the motivation for the algorithm presented in the sections that follow.

Figure 1 illustrates an example of a stencil for a 2D PDE at grid cell (B_x^1, B_y^1) of process 4. At cell (B_x^1, B_y^1) there is a single blue circle surrounded by 4 circles. This structure is known as a 5-point stencil for the 5 blue circles in the image, and it characterizes a well known discretization for the solution of 2D PDEs. The grid cell containing the center circle is the spatially dependent value of the PDE at a given point in time. This value can undergo an update as a function of the values of the grid cells in the adjacent circles for that same point in time. We therefore refer to those adjacent grid cells as dependencies. The unit of time taken for a cell to update the value of its PDE is referred to as the timestep of that cell. Given a well defined stencil, the relationship between the timestep of a cell and its adjacent cells can be studied independently of the values of the underlying PDE.

Figure 2 illustrates the timestep at each cell of process 4 from figure 1, for some arbitrary point during the computation of a PDE's solution. It demonstrates how the spatial dependencies of each cell can change with time. The cell (B_x^1, B_y^1) has a time step of 3, and so do all of its adjacent cells. When a cell has neighbors above, below, left, and right of it which contain values computed from the same timestep, we'll say the cell's dependencies are valid at a given point in time. When any one of its neighbors does not contain a value from the same timestep, then its dependencies are *invalid*. An example of a cell with invalid dependencies is given in cell $(B_x^1, 1)$ in figure 2, where the value of the PDE is only completed up to the 2nd timestep, and it's neighboring cells contain values from different timesteps. Given that timesteps increment for every update, a cell with valid dependencies in one timestep may acquire invalid dependencies immediately following an update, and any numerical solver is constrained by this.

The constraints resulting from the stencil's dependency structure further effects the performance of the parallelization strategy for computing a 2D PDE. A typical strategy starts with the definition of a global 2D area over which a PDE is being solved, followed by a subdivision of that area into separate 2D grids for each processing unit to manage. Figure 1 illustrates the general principle for 4 processing units, each with separate

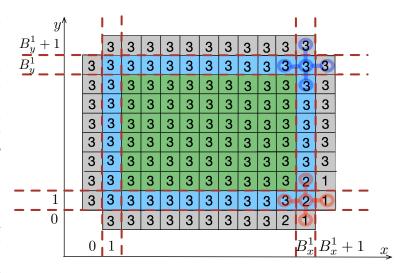
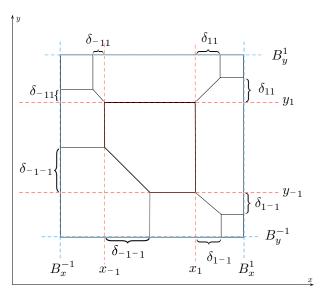
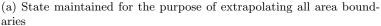
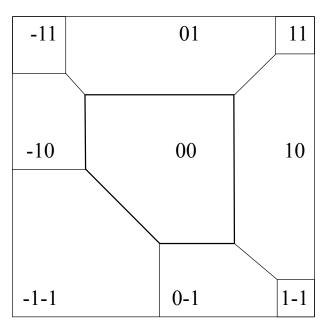


Figure 2: An illustration of grid cell timesteps, and their relationship to stencil dependencies. In cell (B_x^1, B_y^1) dependencies are valid and an update of the cell can be performed. In cell $(B_x^1, 1)$ the opposite is true. Transparent blue and red circles are overlaid respectively for valid and invalid dependencies.

11x8 grid portions of the total 22x16 global grid size. For a parallel numerical method that cannot exploit shared memory, the 11x8 grids are insufficient to capture interprocess dependencies, requiring 13x10 grid cells for each process subdivision illustrated. The







(b) Region labels, each given by 2 digits in the form $a_x a_y$ throughout the rest of the document. $a_x a_y$ additionally provides a shorthand for the vector $[a_x, a_y] = \mathbf{a}$, and we therefore refer to these region labels as region vectors, or area vectors throughout the document.

Figure 3: (3a) An illustration of state labels that allow each of the 9 regions to relax their boundaries. (3b) Shorthand labels for each region, also known as area vectors.

additional gray (ghost) cells which account for the increased dimensions are redundant relative to the global PDE solution, and they hold information which originates from the blue (border) cells held by the adjacent processes. To give a precise example in figure 1, cells $(1,1)-(B_x^1,1)$ of process 1 need to be reflected in cells $(1,B_y^1+1)-(B_x^1,B_y^1+1)$ on process 4 in order for process 4 to compute cells $(1,B_y^1)-(B_x^1,B_y^1)$. The analogous constraint is true for the other border/ghost cell pairs for each of the processes. This data sharing between processes defines an additional process level dependency that is a direct result of the cell level dependency structure introduced by a 5 point stencil.

This dependency structure between processes is known as a nearest neighbor dependency structure, and divides each process' local grid into 9 distinct regions depending on their degree of interprocess dependence. Continuing to use process 4 in figure 1 as an example, the regions are the green shaded interior region, and the 8 blue shaded border regions. The interior region depends only on border regions, and is therefore relatively independent of neighbor processes. The border regions have 2 types of dependence. The first type depends on only one neighbor process. This is given by cell ranges on the 4 sides of the 2D grid: $(2,1) - (B_x^1 - 1,1)$, $(1,2) - (1,B_y^1 - 1)$, $(2,B_y^1) - (B_x^1 - 1,B_y^1)$, and $(B_x^1,2) - (B_x^1,B_y^1 - 1)$. In these ranges only the border cells received from a single adjacent process are needed to successfully compute grid cells at each point. The other type of region depends on 2 processes and exists exclusively in the corners of the 2D grid at cells: (1,1), $(1,B_y^1)$, (B_x^1,B_y^1) , and $(B_x^1,1)$.

Given the 9 static regions described, if a process wishes to execute a complete timestep for each cell in its grid, it must synchronize the pace of its timesteps with each neighbor process. If synchronizations are expensive this creates a performance bottleneck, and it becomes beneficial to develop strategies to relax these synchronization requirements. In order to maintain valid stencil dependencies while relaxing synchronizations requires relaxing the boundaries of the 9 classically static regions of a 2D grid on each process. In the next section we'll describe how this is done, before developing a concise approach for performing region updates that guarantee valid cell level dependencies.

2 State Variables For Region Decomposition

Figure 3b illustrates a concise labeling of each of the 9 regions described above, whose boundaries have been relaxed to arbitrary positions. The regions correspond to our prior explanations as follows:

- 00 is the interior region with no interprocess dependencies.
- $\{-10, 10, 01, 0^{-1}\}$ is the set of border regions with single process dependencies.

Region labels are also referred to as region vectors throughout this paper, and the general form used above, $a_x a_y$, is shorthand for the vector $[a_x, a_y] = \mathbf{a}$.

Figure 3a illustrates the minimal set of variables needed to relax the boundaries of the regions just described. With these dynamic variables of state, each process has enough information to ensure the grid cells of each region have valid dependencies¹. This is because each state variable captures information about the timestep, and therefore dependency information of each region. To illustrate this point, consider the interior region, 00, which has no explicit dependencies. Given that the number of grid cells in 00 dominates the number of grid cells in the border region, we're typically safe to continue shrinking this region and updating it's grid cells, regardless of the last timestep received in the ghost region. Therefore, we define the maximal timestep possible as being the uniform timestep reached by the interior most region, 00,

$$t_{max} \equiv \text{The uniform timestep of each cell in region 00.}$$
 (1)

From this definition, every region's timestep can be further defined relative to t_{max} and the state variables drawn in figure 3a. For example, variable x_{-1} is the difference in the number of timesteps completed by the interior region and the number of timesteps completed at the leftmost edge of region $^{-}10$. x_{-1} is also equal to the number of timesteps since the last time a ghost cell was received along that border. Further, δ_{-1-1} can be thought of as the number of times region $^{-}10$ or region $0^{-}1$ have been updated without a corresponding update to region $^{-}1^{-}1$, making δ_{-1-1} negative as illustrated. Figure 3a also illustrates a positive δ_{1-1} term. This can occur when 00 has updated several times without any updates from regions $0^{-}1$ and 10. These intuitions are summarized analytically for all the dynamic variables illustrated in figure 3a as

$$t(a_x a_y, i_x, i_y) = t_{max} + a_x \cdot x_{a_x} + a_y \cdot y_{a_y} - a_x \cdot i_x - a_y \cdot i_y + |a_x| \cdot |a_y| \cdot \delta_{a_x a_y}, \tag{2}$$

where $t(a_x a_y, i_x, i_y)$ provides the timestep for region $a_x a_y$ in grid cell (i_x, i_y) .

The only variables equation (2) doesn't capture from figure 3a are the static boundary terms, denoted by the B variables. Each B^{-1} term denotes a lower bound, and B^{1} term denotes an upper bound for the grid cells along dimensions x and y which are correspondingly subscripted B_x and B_y .

ALGORITHM 1: Procedure to update state values for each region $a_x a_y$

```
procedure update\_state(a_x a_y) is
          when a_x a_y = 00
                x_{-1} = x_{-1} + 1; \quad y_{-1} = y_{-1} + 1;
 3
                x_1 = x_1 - 1; y_1 = y_1 - 1;
 4
               \delta_{-1-1} = \delta_{-1-1} + 1; \ \delta_{1-1} = \delta_{1-1} + 1; \ \delta_{11} = \delta_{11} + 1; \ \delta_{-11} = \delta_{-11} + 1;
 5
          when a_x a_y = 11^- 1
 6
            \delta_{-1-1} = \delta_{-1-1} + 1;
          when a_x a_y = 1^{-1}
 8
            \delta_{1-1} = \delta_{1-1} + 1;
 9
          when a_x a_y = 11
10
11
            \delta_{11} = \delta_{11} + 1;
          when a_x a_y = 11
12
            \delta_{-11} = \delta_{-11} + 1;
13
          when a_x a_y = 10^{-1}
14
               \delta_{-1-1} = \delta_{-1-1} - 1; \ \delta_{-11} = \delta_{-11} - 1;
15
               x_{-1} = x_{-1} - 1;
16
          when a_x a_y = 10
17
                \delta_{11} = \delta_{11} - 1; \ \delta_{1-1} = \delta_{1-1} - 1;
18
               x_1 = x_1 + 1;
19
          when a_{x}a_{y} = 0^{-}1
20
                \delta_{-1-1} = \delta_{-1-1} - 1; \ \delta_{1-1} = \delta_{1-1} - 1;
21
               y_{-1} = y_{-1} - 1;
22
          when a_x a_y = 01
23
                \delta_{-11} = \delta_{-11} - 1; \ \delta_{11} = \delta_{11} - 1;
24
25
                y_1 = y_1 + 1;
```

While we've defined the state we must track for each region, we have not yet shown how the state evolves to account for delays in synchronization. The details will be explored more below, but for now consider that algorithm 1 will be run for each region, whenever every cell in that region has undergone an update. Lines 3-5 contracts the interior region's boundaries while simultaneously expanding the boundaries of each adjacent region. Lines 6-13 contracts the various corner regions, while expanding their neighbor regions, and lines 14-25 performs the same complimentary contraction and expansion for border regions. Note finally that if algorithm 1 is called for every region $a_x a_y$ for the same global timestep, the region boundaries will all return to where they have started and remain fixed.

¹ This isn't proven formally at the moment, but is part of the research plan

Algorithm 1 is verbose for clarity, but for all other state manipulation we'll introduce a more compact notation. To that end, we build intersection points from the variables introduced. For example, where line $y = x_{-1}$ intersects with line $x = y_{-1}$ is the point $[x_{-1}, y_{-1}]$. In general we'll write,

$$p_{ap} \equiv [x_a, y_p]. \tag{3}$$

For B terms,

$$b_{qp} \equiv [B_x^q, B_y^p]. \tag{4}$$

Finally, for vectors of the subscripts themselves,

$$u_{qp} \equiv [q, p]. \tag{5}$$

We'll operationalize these terms later, and state them in matrix form now ². For boundary intersections,

$$\mathbf{B} \equiv \begin{bmatrix} [B_x^{-1}, B_y^{-1}] & [B_x^{1}, B_y^{-1}] \\ [B_x^{-1}, B_y^{1}] & [B_x^{1}, B_y^{1}] \end{bmatrix} \equiv \begin{bmatrix} b_{-1-1} & b_{1-1} \\ b_{-11} & b_{11} \end{bmatrix}$$
(6)

For interior point intersections,

$$\mathbf{P} \equiv \begin{bmatrix} [x_{-1}, y_{-1}] & [x_1, y_{-1}] \\ [x_{-1}, y_1] & [x_1, y_1] \end{bmatrix} \equiv \begin{bmatrix} p_{-1-1} & p_{1-1} \\ p_{-11} & p_{11} \end{bmatrix}.$$
 (7)

For a subset of labeling terms which will be explored more in the next section,

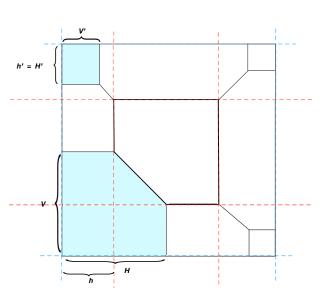
$$\mathbf{U} \equiv \begin{bmatrix} [-1, -1] & [1, -1] \\ [-1, 1] & [1, 1] \end{bmatrix} \equiv \begin{bmatrix} u_{-1-1} & u_{1-1} \\ u_{-11} & u_{11} \end{bmatrix}. \tag{8}$$

Finally for delta terms, we'll write

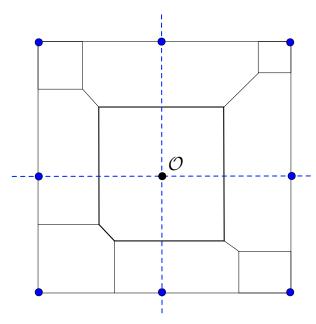
$$\mathbf{\Delta} \equiv \begin{bmatrix} \delta_{-1-1} & \delta_{1-1} \\ \delta_{-11} & \delta_{11} \end{bmatrix}. \tag{9}$$

With this summary of state for the new boundaries of a process' grid, we have the tools to update each grid cell while ensuring valid dependency structures. In the next section, a strategy for updating each region described by this state will be presented.

3 Uniform Region Updates



(a) Illustration of the fundamental dimensions that define each region's boundary: h, H, V and h', H', V'. We'll show in this section how to derive both bounds from the same function



(b) A subdivision of all regions by drawing a horizontal and vertical line through the origin $\mathcal O$ - note how each subdivision of each region looks very similar to the regions annotated in 4a

Figure 4: An illustration of the minimal set of dimensions needed for each region to be described by the same bounds.

While the 9 regions above do not share the same dependency structure with respect to neighbor process cells, they do share the same geometric forms which are captured by the corner regions: The pentagon of region -1-1 and the squares of regions 1-1, 11,

 $^{^{2}}$ Defining equations 3–5 allows us to ignore the inner vectors and manipulate the following terms with matrix operations. A more robust formulation may be to use tensors which is a current topic of investigation.

and -11 in figure 4a for example. The figure further illustrates dimensions H, h and V for regions -1^-1 and H', h' and V' for -11, which uniformly capture both the pentagon and square shapes. We'll refer to the geometric forms described by these examples as tiles or unit tiles.

The uniformity of the tiles illustrated in figure 4a, suggests an approach to iterating over the cells of each region in a uniform way which is summarized in the pseudocode of algorithm 2. In line 1 of the pseudo code, we assume that the unit tiles illustrated in figure 4a can be derived as a function of the region $a_x a_y$, and then in lines 2–3, each tile dimension (H, h, and V) can be derived as a function of the tile specific to that region. This pseudocode is a very rough outline of what we develop formally below, but it serves as useful checkpoint in our intuition.

Figure 4b provides an illustration of how a uniform tile structure can be extracted from every region. This figure rescales figure 4a slightly and draws a subdivision of each region by intersecting perpendicular lines through an origin term (black center dot) given by,

$$\mathcal{O} \equiv \left[x_{-1} + \frac{x_1 - x_{-1}}{2}, \quad y_{-1} + \frac{y_1 - y_{-1}}{2} \right]. \tag{10}$$

The intersecting lines through this point, cut across the center of 00 horizontally and vertically, and cut across the other side regions $(01,0^-1,10,-10)$ either horizontally or vertically. The blue dots illustrated alongside the origin term represent the origins of each unit geometry which can be specified by H, h, and V as illustrated in figure 4a.

With this in mind, as long as we can concisely shift our origin and bounds for each region, we can use the uniform iterates provided by algorithm 2 for every region and its subdivision. We can do this via simple coordinate transformations, and rotations. Transformations are defined by

$$T_x(a_x a_y) = B_x^1 \cdot \max(a_x, 0) + (1 - |a_x|) \cdot \mathcal{O}_x$$

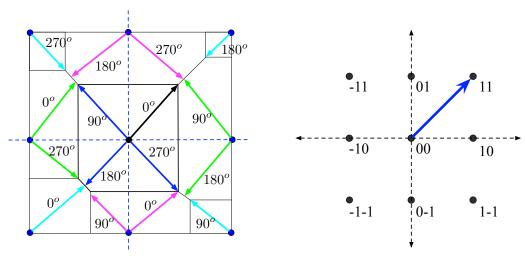
$$\tag{11}$$

$$T_y(a_x a_y) = B_y^1 \cdot \max(a_y, 0) + (1 - |a_y|) \cdot \mathcal{O}_y,$$
 (12)

where \mathcal{O}_x is the first term of definition 10, and \mathcal{O}_y the second term. To rotate our orientation in each region so that each region may share the same origin, we can employ the standard rotation matrix in 2 dimensions,

$$R_{\theta} \equiv \begin{bmatrix} \cos(\theta) & -\sin(\theta) \\ \sin(\theta) & \cos(\theta) \end{bmatrix}. \tag{13}$$

To employ rotations properly, we'll need to know what angles of rotation are required in each region and for each subdivision. To relate these rotations to the bounds of figure 4a, we'll develop a function that maps these angles to the state matrices of section 2.



(a) Vectors and angles corresponding to each region, with color coded sets of angles from $0^{o} - 270^{o}$. The angle is calculated relative to the unit vector of figure 5b

(b) Unit vector overlaid on coordinates corresponding to region labels

Figure 5: Region angles, and the unit vector

3.1 Regional Coordinate Rotation Angles

Figure 5a illustrates the number of rotations present in each region, along with their rotation angles and the vectors they're derived from. In practice, we'll want a procedure to derive these angles using only the region vector, $a_x a_y$. To do so, it's necessary to understand how all of the vectors drawn in figure 5a acquire their angle and magnitude. Figure 5b explains this in part by illustrating the labels of each region overlaid on a cartesian plain, where the labels assemble to form a unit square centered on [0, 0]. Further, the *unit vector* of figure 5b whose tail to head goes from [0, 0] to [1, 1] provides an orientation from which to measure the angles in figure 5a. Our procedure for uncovering angles for each region can now proceed as follows:

- 1. Find the vectors which map to our unit tiles
- 2. Find the angle between each vector and the unit vector
- 3. return the set of angles.

Given that we know the angles we're looking for—enumerated in figure 5a— it is straightforward to verify the derived procedure. To

ALGORITHM 3: Procedure, AngleSearch which takes as input $a_x a_y \in \mathcal{R}$, and returns the set Θ of all the angles associated with it.

```
procedure Angle Search(a_x a_y \in \mathcal{R}) is
            \mathcal{S}' = \{\};
2
             for s \in \mathcal{S}_u do
3
              | \mathcal{S}' = \mathcal{S}' \cup \{ \mathbf{s} + [a_x, \ a_y] \}
4
             \mathcal{S}' = \mathcal{S}' \cap \mathcal{R};
5
             \Theta = \{\};
6
             for \mathbf{s} \in \mathcal{S}' do
7
                    \mathbf{u} = \mathbf{s} - [a_x, \ a_y];
                    \Theta = \Theta \cup atan2(\mathbf{u} \cdot \mathbf{v}^u, |\mathbf{u}^T||\mathbf{v}^{uT}|);
9
            return \Theta;
```

find the angles enumerated in figure 5a, we formalize some notions of our unit labelings of each region:

$$\mathcal{R} \equiv \{-1, 0, 1, 1, 1, -1, 0, 00, 10, -11, 01, 11\}$$

$$(14)$$

$$\mathbf{v}^u \equiv [1, 1]. \tag{16}$$

 \mathcal{R} is just the set of all possible regions where $a_x a_y \in \mathcal{R}$ always. \mathcal{S}_u is a subset of \mathcal{R} , and provides a way to derive the angles of figure 5a when taken together with \mathbf{v}^u , the unit vector. Finally, recall that our notation for $a_x a_y$, can also be thought of as indicating a vector $[a_x, a_y]$, and we will use the notations interchangeably.

Algorithm 3 formalizes the procedure outlined in steps 1-3 above, and proceeds as follows. Line 4 adds every region vector $a_x a_y$ to each element of S_u , before finding the intersection of the resulting vectors with the set of valid labels in line 5. With these valid label's in hand, line 8 reverses the operation of line 5, and the resulting \mathbf{u} are in fact the vectors displayed and color coded for each region in figure 5a. Finally, the angle between the vector \mathbf{u} and the unit vector \mathbf{v}^u are calculated in line 8. The trigonometric function atan2 is used in order to return an angle between 0 and 360 to conform to our target rotations (figure 5a). The inputs to this function are the dot product of \mathbf{u} and \mathbf{v}^u followed by the determinant of the same vectors concatenated together—concatenation denoted by ||, and the determinant denoted with vertical bars on each side of the expression, $|\cdot|$.

3.2 Indexing State By Region Label and Rotation Angles

In order to write an analytic expression for H, h, and V, one more relationship is required: The state defined in section 2 as a function of the region $a_x a_y$ and the elements of its set $\theta \in \Theta$, described in the prior section. This relationship has two parts: a method of indexing 2x2 matrices by an angle $\theta \in \{0, 90, 180, 270\}$, and operations to flip the rows and columns of a 2x2 matrix. We'll start with an intuition of the indexing scheme. Recall the variables of state 7 and 9:

$$\mathbf{P} \equiv \begin{bmatrix} [x_{-1}, y_{-1}] & [x_1, y_{-1}] \\ [x_{-1}, y_1] & [x_1, y_1] \end{bmatrix},$$

and

$$\mathbf{\Delta} \equiv \begin{bmatrix} \delta_{-1-1} & \delta_{1-1} \\ \delta_{-11} & \delta_{11} \end{bmatrix}.$$

Note that for region 00, H, V, and h can all be written in terms of each term in these matrices for each angle θ by moving clockwise starting in the bottom right corner of each matrix (i.e. \mathbf{P}_{11} and $\boldsymbol{\Delta}_{11}$). For example at $\theta = 0^{\circ}$ and for region 00, H, V, and h will

all be bounded by $[x_1, y_1]$ and δ_{11} (i.e. the terms of \mathbf{P}_{11} and $\boldsymbol{\Delta}_{11}$). This observation on the pattern of matrix indexing according to angle rotations can be generalized for rows (denoted r) and columns (denoted c) of matrices as

$$c \equiv \min(1 + [R_{-\theta} \cdot \mathbf{v}^u]_u, 1) \tag{17}$$

$$\mathbf{r} \equiv \min(1 + [R_{-\theta} \cdot \mathbf{v}^u]_x, 1). \tag{18}$$

To simplify notation however, for any 2x2 matrix M and angle $\theta \in \{0, 90, 180, 360\}$, we write

$$M[\theta] \equiv M_{\rm rc},$$
 (19)

which allows us to directly index variables of state by angle θ . However this function will remain limited to region 00 without first providing regional transformations for the matrices of state presented in section 2.

The matrices of state given by equations (6)–(9) can be indexed by θ for all regions $a_x a_y \in \mathcal{R}$ as in equation (19) if they are first transformed. This transformation is given by the following function, where for the state matrix M, the region transformation $M_{a_x a_y}$, is defined as:

$$M_{a_x a_y} \equiv \begin{pmatrix} 0 & 1 \\ 1 & 0 \end{pmatrix}^{2-|a_y|} \cdot M \cdot \begin{pmatrix} 0 & 1 \\ 1 & 0 \end{pmatrix}^{2-|a_x|}, \tag{20}$$

where the pre-multiplication of matrix $\begin{pmatrix} 0 & 1 \\ 1 & 0 \end{pmatrix}$ flips the rows of matrix M, and the post-multiplication of matrix M by $\begin{pmatrix} 0 & 1 \\ 1 & 0 \end{pmatrix}$ flips its columns. Given our relatively small state space, this can be verified as yielding appropriate matrices of state for each region through an exhaustion of cases.

3.3 Uniform Region Bounds

Taken together, equations 19 and 20, allow us to retrieve the variables of state given θ and $a_x a_y$,

$$\mathbf{p} = \mathbf{P}_{a_r a_n}[\theta] \tag{21}$$

$$\mathbf{b} = \mathbf{B}_{a_n a_n}[\theta] \tag{22}$$

$$\delta = \Delta_{a_n a_n}[\theta] \tag{23}$$

$$\mathbf{u} = \mathbf{U}_{a_n a_n}[\theta]. \tag{24}$$

From equations 21–24, we can write $h, H, V : (a_x a_y, \theta) \to \mathbb{R}^+$,

$$h(\theta, \mathbf{a}) = [R_{-\theta} \cdot \mathbf{p}^T]_x + [R_{-\theta} \cdot \mathbf{a}^T]_x \cdot [R_{-\theta} \cdot \mathbf{b}^T]_x - [R_{-\theta} \cdot \mathcal{O}^T]_x \cdot (1 - |R_{-\theta} \cdot \mathbf{a}^T|_x) + (1 - |R_{-\theta} \cdot \mathbf{a}^T|_y) \cdot \min_{x, y} (R_{-\theta} \cdot \mathbf{u}^T) \cdot \delta \quad (25)$$

$$H(\theta, \mathbf{a}) = [R_{-\theta} \cdot \mathbf{p}^T]_x + [R_{-\theta} \cdot \mathbf{a}^T]_x \cdot [R_{-\theta} \cdot \mathbf{b}^T]_x - [R_{-\theta} \cdot \mathcal{O}^T]_x \cdot (1 - |R_{-\theta} \cdot \mathbf{a}^T|_x) + |R_{-\theta} \cdot \mathbf{a}^T|_y \cdot [R_{-\theta} \cdot \mathbf{u}^T]_x \cdot \delta$$
(26)

$$V(\theta, \mathbf{a}) = [R_{-\theta} \cdot \mathbf{p}^T]_y + [R_{-\theta} \cdot \mathbf{a}^T]_y \cdot [R_{-\theta} \cdot \mathbf{b}^T]_y - [R_{-\theta} \cdot \mathcal{O}^T]_y \cdot (1 - |R_{-\theta} \cdot \mathbf{a}^T|_y) + |R_{-\theta} \cdot \mathbf{a}^T|_x \cdot [R_{-\theta} \cdot \mathbf{u}^T]_y \cdot \delta$$
(27)

Making these functions discrete (i.e. $h, H, V : (a_x a_y, \theta) \to \mathbb{Z}^+$) requires reworking equation 10 with the ceiling function as follows,

$$\mathcal{O} \equiv \left[x_{-1} + \left\lceil \frac{x_1 - x_{-1}}{2} \right\rceil, \quad y_{-1} + \left\lceil \frac{y_1 - y_{-1}}{2} \right\rceil \right], \tag{28}$$

and using using it in equations (25)–(27).

Algorithm 4 puts these equations together and gives precise analytic form to the pseudocode developed in algorithm 2. In the interest of concision, line 6 omits a precise definition of a 5 point stencil, but simply states that g(t, x, y) = f(t, x, y) meaning at time t, we can compute the value of a stencil, g(t, x, y), given some function f of the same point x, y at the prior time t - 1. Line 1 executes the procedure of algorithm 3, and illustrates the parallel between our notion of a tile and it's apparent encoding as an angle. The potential for an intersection in the mathematical research on tiling is intriguing, but has yet to bear analytic fruit beyond the broad observations made in this section.

ALGORITHM 4: Uniform Region Update \forall input $a_x a_y = \mathbf{a} \in \mathcal{R}$

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\begin{array}{lll} \mathbf{1} & \Theta = AngleSearch(a_xa_y); \\ \mathbf{2} & \mathbf{for} \ \theta \in \Theta \ \mathbf{do} \\ \mathbf{3} & & \mathbf{for} \ j = 0 \ \mathbf{to} \ j = H(\theta, \mathbf{a}) \ \mathbf{do} \\ \mathbf{4} & & & \mathbf{for} \ k = 0 \ \mathbf{to} \ k = \min\{V(\theta, \mathbf{a}), V(\theta, \mathbf{a}) - j + h(\theta, \mathbf{a}) \ \mathbf{do} \\ \mathbf{5} & & & & \mathbf{i} = R_\theta \cdot [j, \ k]^T; \\ \mathbf{6} & & & & \mathbf{g}(t, T_x(\mathbf{a}) + \mathbf{i}_x T_y(\mathbf{a}) + \mathbf{i}_y)) = f(t - 1, T_x(\mathbf{a}) + \mathbf{i}_x, T_y(\mathbf{a}) + \mathbf{i}_y) \end{array}
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4 Update Constraints for the Multi-Process Problem

Up until now we've been considering updates on a single process using the regions decomposed by the state laid out in section 2, but we still need to consider coordination between processes. Recall in our discussion of algorithm 1 that if the procedure is called once for every region (i.e. $\forall a_x a_y \in \mathcal{R}$), then the borders introduced in the static region decomposition in section 1 remain unchanged. However when this requirement is relaxed, then a few things immediately follow. In particular, certain conditions must be met for algorithm 4 to be invoked by a process region at all. This section enumerates those conditions.

4.1 Nonzero Region Area

This first such condition is that for every region $a_x a_y \in \mathcal{R}$, and for all $\theta \in AngleSearch(a_x a_y)$, it must be the case that $h(\theta, a_x a_y) > 0$, $H(\theta, a_x a_y) > 0$, and $V(\theta, a_x a_y) > 0$. This can be summarized by the boolean function,

$$\mathcal{G}^{S}(a_{x}a_{y},\Theta) \equiv \underset{\theta \in \Theta}{\wedge} H(\theta,\mathbf{a}) > 2 \wedge h(\theta,\mathbf{a}) > 2 \wedge V(\theta,\mathbf{a}) > 2, \tag{29}$$

where $\Theta = Angle Search(a_x a_y)$. Here, 0 is replaced by 2 given that in discrete space, a 5 point stencil must have at minimum a 3x3 grid of cells to update a single point (or at least 5 cells, with its 4 corners missing). We'll refer to formula 29 as the *spatial update quard*.

4.2 Interprocess Communication Constraints

Any process wishing to update its border cells must receive ghost cells from one or more neighbor processes. Therefore, there must be a means of ensuring that the requisite *communication* (sending and receiving of data) is performed and completed. Given our assumption that communication completion is nondeterministic, the abstraction we will use will mimic MPI's non-blocking communication. Let every process have a unique id $\operatorname{pid} \in \mathbb{Z}^+$. Let the $\mathcal{N}(a_x a_y, \operatorname{pid}) : (a_x a_y, \operatorname{pid}) \to \{\mathbb{Z}^+\}$ be a mapping from a region $a_x a_y$ and a process id pid to a set of positive integers which themselves are process ids of the processes that pid must communicate with (these ids are referred to as *neighbors*).

Next, we define functions analogous to MPI's ISend and IRecv respectively. $\operatorname{send}_{a_x a_y}[n] \leftarrow \operatorname{postSend}(n \in \mathbb{Z}^+, \mathsf{d})$ is the signature for a function which takes a neighbor process id as its first argument, and some arbitrary data structure d as its second argument, and returns a boolean value in the array $\operatorname{send}_{a_x a_y}[0..|\mathcal{N}(a_x a_y, \mathsf{pid})|]$, which is initially set to false, and then nondeterministically set to true when the neighbor process n receives the value d , and copies it to a local data structure. Likewise, $\operatorname{rcv}_{a_x a_y}[n] \leftarrow \operatorname{postRecv}(n \in \mathbb{Z}^+, \mathsf{d})$ is the signature for a function which takes a neighbor process id as its first argument, and some arbitrary local data structure d as its second argument, and returns a boolean value in the array $\operatorname{rcv}_{a_x a_y}[0..|\mathcal{N}(a_x a_y, \mathsf{pid})|]$, which is initially set to false, and then nondeterministically set to true when the neighbor process pid receives a new value in its local data structure d from process n, and can perform a validly access the memory for d .

Given these definitions for communication between neighbor processes, it follows for all $a_x a_y \in \mathcal{R} - \{00\}$ that the following boolean function must be true for any valid update to be performed:

$$\mathcal{G}^C(a_x a_y, \mathsf{pid}) \equiv \underset{n \in \mathcal{N}(a_x a_y, \mathsf{pid})}{\wedge} \mathsf{send}_{a_x a_y}[n] \wedge \mathsf{rcv}_{a_x a_y}[n]. \tag{30}$$

We'll call this the *communication update guard*.

4.3 Mirror Process Misalignment

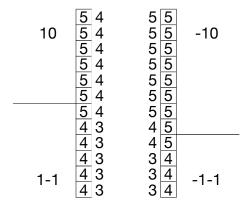


Figure 6: A motivating example for the utility of $\mathcal{G}_{\mathcal{M}}$. A subset of the 2D data arrays of a left and right process are illustrated—ghost cells drawn without a border, and border cells given a border. A delimiter between regions on each process perpendicular To B_x^1 and B_x^{-1} is drawn for the left and right process, respectively. Region labels are shown for the 4 regions of interest to the example, and the rest are omitted

Given a global spatial decomposition as described in section 1, 2 neighbor processes are referred to as mirror processes of each other. In figure 1, process 2 and 4 are process 1's mirror processes, process 1 and 3 are process 2's mirror processes, and so on. Further, the state variables outlined in section 2 are referred to as the *mirror state* of the processes adjacent to them.

Figure 6 illustrates an example how the mirror state of 2 processes can become misaligned. This discussion will not show how such a state is reachable, but what must be done when it occurs in order to respect the dependency constraints of a cell update at the border. Drawn is a left process with regions -10 and 10 shown, and its mirror process on the right with regions -10 and -1-1. Border cells of the processes are bordered by a square boxes, and a timesteps are displayed within them. Ghost cell timesteps are displayed with no border. Observe that the border cells of the left process are exactly reflected in the ghost cells of the right process

ALGORITHM 5: Procedure to check mirror state for each $a_x a_y \in \mathcal{R}$

```
1 procedure \mathcal{G}^M(a_x a_y \in \mathcal{R} - \{00\}, \Theta) is
          \theta_1 = \Theta[0]; \quad \theta_2 = \Theta[1] ? |\Theta| = 2 : \theta_1;
          return H[\theta_1] \leq V^M[\theta_1 - 90] \wedge V[\theta_2] \leq V^M[\theta_2 + 90];
```

and vice versa. This in fact corresponds to the state in each process, \mathbf{P} and Δ . We assume further that this state is exchanged between processes, and stored as locally in mirror variables, where

$$\Delta_n^M \equiv \Delta \text{ matrix of neighbor process } n$$
(31)
$$\mathbf{P}_n^M \equiv \mathbf{P} \text{ matrix of neighbor process } n.$$
(32)

$$\mathbf{P}_n^M \equiv \mathbf{P} \text{ matrix of neighbor process } n.$$
 (32)

It follows, that any calculation made with these terms will receive a superscript M to denote that it is in fact measuring the term for the mirror process (e.g. $V^M(\theta, \mathbf{a}), h^M(\theta, \mathbf{a})$).

To ensure that a region $a_x a_y$ is not updated when its appropriate ghost cells are not present as in region -10 in figure 6, the boolean function \mathcal{G}^M shown in algorithm 5 can be used. The function returns true if the region is an interior region, and then checks alignment with mirror quantities from its neighbor process(es). $H(\theta, \mathbf{a})$ is mirrored by $V^M(\theta - 90^o, \mathbf{a})$, and $V(\theta, \mathbf{a})$ is mirrored by $V^M(\theta+90^o,\mathbf{a})$. This is a relationship that holds up for all $a_x a_y \in \mathcal{R} - \{00\}$, and can be checked by drawing out the rotated bounds V and H for all border and corner regions, along side their region bounds.

5 Full Algorithm

The full description for a parallel decomposition of a 5 point stencil is provided in algorithm 6. Some details that haven't been explored are provided here for completeness. To start with, how to handle discrete space indexing given rotations within a region $a_x a_y$. The modulo terms below account for this (where $r = x \mod y$, and r is the remainder from dividing x into y). Given its frequency, the shorthand $\overline{x} = x \mod 2$ is used as well when y = 2. Additionally, a termination condition has not been introduced for the algorithm. Recall that the modeling problem initially set out in section 1 is that of PDEs, which evolves a spatially dependent quantity over time. Therefore the stopping condition will be the number of timesteps to evolve the spatial property over, nsteps.

Finally, we will actually introduce operations on an array of cells being used to model the PDE. data $[0..1][0..B_u^1+1][0..B_x^1+1]$ will hold all the grid cells regardless of region, including the ghost cells. It is a 3D array with a 'double buffering' strategy for updating in time in the first dimension, and the height and length of the local portion of the process' PDE stored in the rows and columns, respectively. For a set of shorthands for defining the bounds to send and receive in each spatial dimension of data we define

$$\mathbb{I}_x(q,\theta,\mathbf{a}) = T_x + |a_x| \cdot (a_x - q) \cdot R_\theta \cdot [H(\theta,\mathbf{a}), V(\theta,\mathbf{a})]_x^T$$
(33)

$$\mathbb{I}_{y}(q,\theta,\mathbf{a}) = T_{y} + |a_{y}| \cdot (a_{y} - q) \cdot R_{\theta} \cdot [H(\theta,\mathbf{a}), V(\theta,\mathbf{a})]_{y}^{T}, \tag{34}$$

where $q \in \{-1,1\}$ indicating the left bound and right bound respectively. This is a discrete function which works for the receive buffers (ghost cells), but given we're also accessing send buffers (border cells), this won't be sufficient. This combined with the need to account for the case where $\mathbb{I}_x(-1,\theta,\mathbf{a}) = \mathbb{I}_x(+1,\theta,\mathbf{a})$ motivates the additional helper function

$$\mathbb{I}_x(q,\theta,\mathbf{a})^s = \min(\max(\mathbb{I}_x(q,\theta,\mathbf{a}) - q, 1), B_x^1)$$
(35)

$$\mathbb{I}_{y}(q,\theta,\mathbf{a})^{s} = \min(\max(\mathbb{I}_{y}(q,\theta,\mathbf{a}) - q, 1), B_{y}^{1}). \tag{36}$$

Finally, note that we do not define the entire stencil update method for a 5 point stencil, but continue to denote it by f', which no longer needs a variable of time (as in algorithm 4), as our data buffer has made this access precise. Further, q has been dropped from the original formulation of algorithm 4 as the left hand side of the update need only access the single appropriate center cell of the 5-point dependency structure under study.

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```
/* Initialize Global Variables.
  1 t_{max} \leftarrow 1;
  2 x_{-1} \leftarrow 1; x_1 \leftarrow B_x^1; y_{-1} \leftarrow 1; y_1 \leftarrow B_y^1;
  3 \delta_{-11} \leftarrow 0; \delta_{-1-1} \leftarrow 0; \delta_{11} \leftarrow 0; \delta_{1-1} \leftarrow 0;
  4 for a_x a_y \in \mathcal{R} - \{00\} do
                \Theta = \text{AngleSearch}(a_x a_y);
               for \theta \in \Theta do
                       for n \in \mathcal{N}(a_x a_y, \mathsf{pid}) do
                                send_{a_x a_y}[n] \leftarrow postSend(n, \Delta); \quad rcv_{a_x a_y}[n] \leftarrow postRecv(n, \Delta_n^M);
  8
                                send_{a_x a_y}[n] \leftarrow postSend(n, \mathbf{P}); \quad rcv_{a_x a_y}[n] \leftarrow postRecv(n, \mathbf{P}_n^M);
  9
                                \mathsf{send}_{a_x a_y}[n] \leftarrow \mathit{postSend}(n, \mathsf{data}[1][\mathbb{I}_y(-1, \theta, \mathbf{a})^s..\mathbb{I}_y(+1, \theta, \mathbf{a})^s][\mathbb{I}_x(-1, \theta, \mathbf{a})^s..\mathbb{I}_x(+1, \theta, \mathbf{a})^s]);
10
                                  \mathsf{rcv}_{a_x a_y}[n] \leftarrow postRecv(n, \mathsf{data}[0][\mathbb{I}_y(-1, \theta, \mathbf{a})..\mathbb{I}_y(+1, \theta, \mathbf{a})][\mathbb{I}_x(-1, \theta, \mathbf{a})..\mathbb{I}_x(+1, \theta, \mathbf{a})]);
11 while t_{max} < nsteps \lor x_{-1} > 0 \lor y_{-1} > 0 \lor x_1 < B_x^1 \lor y_{-1} > B_y^1 do
               for a_x a_y \in \mathcal{R} do
12
                        \Theta = \text{AngleSearch}(a_x a_y);
13
                        when \mathcal{G}^S(a_x a_y, \Theta) \wedge ((\mathcal{G}^C(a_x a_y, \mathsf{pid}) \wedge \mathcal{G}^M(a_x a_y, \Theta)) \vee a_x a_y = 00)
14
                                \theta_c = 0;
15
                                for \theta \in \Theta do
16
                                        \mathbf{p} = \mathbf{P}_{a_x a_y}[\theta];
17
                                        \mathbf{b} = \mathbf{B}_{a_x a_y}[\theta];
18
                                        \delta = \mathbf{\Delta}_{a_x a_y}[\theta];
19
                                        \mathbf{u} = \mathbf{U}_{a_x a_y}[\theta];
20
                                        h(\theta, \mathbf{a}) = [R_{-\theta} \cdot \mathbf{p}^T]_x + [R_{-\theta} \cdot \mathbf{a}^T]_x \cdot [R_{-\theta} \cdot \mathbf{b}^T]_x - [R_{-\theta} \cdot \mathcal{O}]_x \cdot (1 - |R_{-\theta} \cdot \mathbf{a}^T|_x) + (1 - |R_{-\theta} \cdot \mathbf{a}^T|_y) \cdot \min_{x, y} (R_{-\theta} \cdot \mathbf{u}) \cdot \delta;
21
                                        H(\theta, \mathbf{a}) = [R_{-\theta} \cdot \mathbf{p}^T]_x + [R_{-\theta} \cdot \mathbf{a}^T]_x \cdot [R_{-\theta} \cdot \mathbf{b}^T]_x - [R_{-\theta} \cdot \mathcal{O}]_x \cdot (1 - |R_{-\theta} \cdot \mathbf{a}^T|_x) + |R_{-\theta} \cdot \mathbf{a}^T|_y \cdot [R_{-\theta} \cdot \mathbf{u}]_x \cdot \delta;
22
                                        V(\theta, \mathbf{a}) = [R_{-\theta} \cdot \mathbf{p}^T]_y + [R_{-\theta} \cdot \mathbf{a}^T]_y \cdot [R_{-\theta} \cdot \mathbf{b}^T]_y - [R_{-\theta} \cdot \mathcal{O}]_y \cdot (1 - |R_{-\theta} \cdot \mathbf{a}^T|_y) + |R_{-\theta} \cdot \mathbf{a}^T|_x \cdot [R_{-\theta} \cdot \mathbf{u}]_y \cdot \delta;
23
                                        \mathbf{q} = R_{\theta} \cdot [1, 1]^T;
\mathbf{24}
                                        for j = 0 to j = H(\theta, \mathbf{a}) - \mathbf{q}_x do
25
                                                for k = 0 to k = \min\{V(\theta, \mathbf{a}) - \mathbf{q}_y, V(\theta, \mathbf{a}) - j + h(\theta, \mathbf{a}) - \mathbf{q}_y\} do
26
                                                        \mathbf{i} = R_{\theta} \cdot [j, \ k]^T;
27
                                                        \mathsf{data}[1 - \overline{t(a_x a_y, j, k)}][T_y(\mathbf{a}) + \mathbf{i}_y) + \mathbf{q}_y \cdot (1 - |a_y|) \cdot \min((\theta_c \bmod 3), 1)][T_x(\mathbf{a}) + \mathbf{i}_x + \mathbf{q}_x \cdot (1 - |a_x|) \cdot \overline{\theta_c}] = 0
28
                                                           f'(\mathsf{data}[t(a_xa_y,j,k)][T_y(\mathbf{a}) + \mathbf{i}_y + \mathbf{q}_y \cdot (1-|a_y|) \cdot \min((\theta_c \bmod 3),1)][T_x(\mathbf{a}) + \mathbf{i}_x + \mathbf{q}_x \cdot (1-|a_x|) \cdot \overline{\theta_c}])
                                        \theta_c = \theta_c + 1;
29
                                when a_x a_y \neq 00
30
                                        for \theta \in \Theta do
31
                                                for n \in \mathcal{N}(a_x a_y, \mathsf{pid}) do
32
                                                        send_{a_x a_y}[n] \leftarrow postSend(n, \Delta); \quad rcv_{a_x a_y}[n] \leftarrow postRecv(n, \Delta_n^M);
33
                                                        send_{a_x a_y}[n] \leftarrow postSend(n, \mathbf{P}); \quad rcv_{a_x a_y}[n] \leftarrow postRecv(n, \mathbf{P}_n^M);
34
                                                         \mathsf{send}_{a_x a_y}[n] \leftarrow postSend(n, \mathsf{data}[1 - \overline{t(a_x a_y, j, k)}][\mathbb{I}_y(-1, \theta, \mathbf{a})^s..\mathbb{I}_y(+1, \theta, \mathbf{a})^s][\mathbb{I}_x(-1, \theta, \mathbf{a})^s..\mathbb{I}_x(+1, \theta, \mathbf{a})^s]);
35
                                                           \mathsf{rcv}_{a_x a_y}[n] \leftarrow postRecv(n, \mathsf{data}[\overline{t(a_x a_y, j, k)}][\mathbb{I}_y(-1, \theta, \mathbf{a})..\mathbb{I}_y(+1, \theta, \mathbf{a})][\mathbb{I}_x(-1, \theta, \mathbf{a})..\mathbb{I}_x(+1, \theta, \mathbf{a})]);
                                when a_x a_y = 00
36
                                   t_{max} \leftarrow t_{max} + 1;
37
                                update\_state(a_x a_y);
```