# Conservative Front Tracking in Higher Space Dimensions 

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#### Abstract

We propose a fully conservative Front Tracking algorithm in two space dimension. The algorithm first uses the point shifted algorithm [12] on two adjacent time levels and then constructs space time hexahedra as computational units. We develope and prove a successful geometric construction under certain interface requirement. This algorithm has a first order local truncation error for cells near the tracked discontinuity, which is an improvement by one order of accuracy over most finite difference schemes, which have $\mathcal{O}(1)$ local truncation errors near discontinuities.


[^0]
## 1 Introduction

We propose and demonstrate a tracking finite difference algorithm which is (a) fully conservative and (b) improves local truncation error by one order (from $\mathcal{O}$ (1) to $\mathcal{O}(\Delta x)$ near tracked discontinuities.

Discontinuities in the solutions of systems of nonlinear hyperbolic conservation laws are widely recognized as a primary difficulty for numerical simulation. Surprisingly, the nonlinearities actually help, as they cause information, which flows along solution characteristics, to flow into the discontinuity, and disappear there. The nonlinear discontinuities (shock waves) function much as a black hole in this regard. Included in this flow of information are the solution errors generated by the nonlinear discontinuity. Because the nonlinear discontinuities absorb errors associated with their numerical approximation, these errors do not grow or spread with time.

Nonlinear equations also have linear discontinuities. In gas dynamics these are the contact discontinuities, across which temperature or shear velocities can be discontinuous. Errors in these modes are never forgiven and never dissipated. For this reason the linear modes are more difficult to control numerically. The dominant numerical solution error is typically associated with discontinuities in these linear modes and occurs as diffusion of mass, vorticity, and temperature. These errors increase with time.

Front Tracking was introduced to give special treatment to solution with discontinuities. Perceptions that Front Tracking software difficulties would be insurmountable were too pessimistic, and a robust, validated code has been developed and used in production simulation of fluid instabilities [5, 7, 6, 4]. See also the URL http://www.ams.sunysb.edu/~shock/FTdoc/FTmain.html. Here we address an algorithmic issue: formulation of a conservative tracking algorithm. In its original formulation, conservation was enforced only in regular grid cells, those not cut by the tracked front. The missing points of the computation stencil, in the case of a front cutting through the stencil, are filled in as ghost cells, with the state values obtained by extrapolation from nearby front states of the same component. Thus the state values are double valued near the front, with the left-component states extending by extrapolation for a small distance into the right component, and vice versa. The use of ghost cell states was introduced into Front Tracking in 1980 [9]. With the ghost states thus defined, the interior solver follows a conventional finite difference algorithm.

In the level set method [3] and the original Front Tracking, ghost cells constructed near the front (but using entropy extrapolation) allows a standard difference operation update. As with Front Tracking, the ghost cell extrapolation is non conservative and leads to $\mathcal{O}(1)$ local truncation error.

Here we propose an algorithm which is conservative for all grid cells, including the irregular ones cut by the front. The algorithm we propose is related to earlier work of Swartz and Wendroff [13], Harten and Hyman [11], and Colella and Chern [2], but differs from these works in several ways. We emphasize here tracking of a contact, rather than the shock tracking of [2]. Our support for fronts is fully general and can be used for unstable, convoluted, and bifurcating interfaces. The 1D version of this algorithm [8] is formally second order accurate in the $\mathrm{L}_{1}$ norm except for interactions of tracked waves. An important difference with [13] is our discussion of the extension to higher dimensions.

## 2 The Two Dimensional Algorithm

Consider the two space dimensional system of conservation laws

$$
\begin{equation*}
\frac{\partial u}{\partial t}+\frac{\partial f(u)}{\partial x}+\frac{\partial g(u)}{\partial y}=0 \tag{1}
\end{equation*}
$$

defined in a spatial domain $\Omega$. Assume that $\Omega$ can be partitioned by a uniform square grid and the boundary is along the grid lines. The side of a cell in the grid is of length $\Delta x$.

Our algorithm is organized into three main steps. The first is the construction of a spatial grid locally conforming to the old ( $\operatorname{time} t_{n}$ ) and new propagated (time $t_{n+1}$ ) INTERFACE. The second is construction of a space time grid joining these two, and the third is a finite volume discretization associated with the space time conforming grid.

In the following two sections, we emphasize the major results and leave the more technical lemmas and some proofs to the Appendix.

### 2.1 The Point Shifted Algorithm

In this section we describe a simple point shifted algorithm to achieve an INTERFACE conforming grid node displacement at a fixed time level. We begin with hypotheses which are requirements on the topology of the INTERFACE and the size of the cell. In the present study the INTERFACEs are topologically equivalent to a union of line segments or circles. Thus we postulate that triple or multiple CURVE intersection points do not occur. INTERFACEs that involve topological change during the time evolution can be resolved by premerging when the distance of the gap to be merged is within $\Delta x$. Analysis of this step is out of the scope of the present paper, but is supported in the numerical implementation of the point shifted construction.

The discretized INTERFACE [10] is a disjoint union of non intersecting CURVEs. Each CURVE is piecewise linear and connected, and composed of BONDs. Each $B O N D$ is a pair of INTERFACE POINTs or POINTs, and (conceptually) the straight line segment joining them. Each CURVE is assigned an orientation which remains unchanged during the propagation of the INTERFACE. If all the POINTs are on the interior of cell edges with at most one POINT occuring on the interior of any given grid cell edge, then the INTERFACE is called grid based [7]. Propagation of the POINTs of a grid based INTERFACE will yield a general INTERFACE, not grid based, as there is no reason for a propagated POINT to lie on a grid cell edge, just because it starts on one. According to the grid based construction of [10], we consider this propagated INTERFACE as a collection of polygonal CURVEs in $\Re^{2}$. Crossing points of the CURVE with grid cell edges are inserted as new POINTs. The propagated old POINTs will be deleted (named images of propagation in this sense), but their ordering along the CURVEs will be retained for later use in the construction of space time interface. The CURVE is then reconstructed, as straight line segments joining these new POINTs. In this process, the CURVE is displaced by an amount $\mathcal{O}\left(\Delta x^{2}\right)$, assuming that the CURVE is smooth, so that all angles between neighboring $B O N D$ s are $\mathcal{O}(\Delta x)$. Also all images of propagated POINTs on the original CURVE can be projected onto the grid based CURVE, with their ordering unchanged and a maximum displacement $\mathcal{O}\left(\Delta x^{2}\right)$.

This grid based INTERFACE is the starting point for the interface conforming volume grid which we construct here. An INTERFACE with a displaced rectangularly indexed volume grid is called a point shifted INTERFACE. It is point shifted if the grid corners have been displaced so that all POINTs are at displaced grid cell corners and all $B O N D$ s are either the edges or diagonals of displaced grid cells.

Here we construct an algorithm which yields a point shifted INTERFACE at each time step. In two space dimensions, we use the front propagation algorithm developed in $[5,7,6,4]$ to follow the INTERFACE evolution. We first projected the propagated INTERFACE to be grid based [7], by inserting new POINTs at cell edge crossings, and then removing old POINTs. To this INTERFACE we apply the point shifted algorithm [12] near the front on each time level to align the grid nodes nearest to the INTERFACE so that there is no intersection between the INTERFACE and the interior of cell edges (i.e., the INTERFACE passes through displaced grid cell corners only and thus lies on the diagonals and edges of displaced grid cells). Some POINTs are deleted in this construction; again with maximum displacement $\mathcal{O}\left(\Delta x^{2}\right)$ of the propagated old POINTs and the propagated CURVEs. We call the result an interface conforming grid node displacement.

Hypothesis 1 The INTERFACE is assumed to be grid based. Each CURVE is topologically equivalent to a line segment with its two end points on the boundary, or
a circle contained in the interior of $\Omega$. (Triple points where three or more CURVEs meet at a point are disallowed.) Each CURVE has at least three BONDs and the maximum angle between two adjacent BONDs is $\mathcal{O}(\Delta x)$. All interior POINTs of the CURVE must be interior to $\Omega$. There is no topological change of the INTERFACE during the time interval of computation.

To avoid consideration of degenerate cases, we assume that POINTs never lie exactly at center of a grid cell edge.

Hypothesis 2 At most one BOND intersects the interior of a given cell, and if this occurs, the CURVE separates the interior of the cell into two non-trivial domains.

This Hypothesis implies that at most two edges of a cell intersect the INTERFACE.

Hypothesis 3 No CURVE is totally contained within a square of side $2 \Delta x$ made up of four cells.

A grid node has grid distance $d$ to the INTERFACE if there is a grid line segment of length $d$ connecting the grid node to the INTERFACE. A grid node will have multiple grid distances. We call the smallest one the shortest grid distance. A grid node is called fixed if all its grid distances are greater than $\Delta x / 2$.

We call a grid node which is not on the boundary shiftable if one of the following three conditions holds: (See Fig. 1.)
(I1) Exactly one of its grid distances is less than $\Delta x / 2$. If this grid distance is on a grid line parallel to the $x$-axis, we call it $x$-shiftable, otherwise we call it $y$-shiftable.
(I2) Exactly two of its grid distances are less than $\Delta x / 2$ and they are not on the same grid line. If the shortest grid distance is on a grid line parallel to the $x$ axis, we call it $x$-shiftable; if the shortest grid distance is on a grid line parallel to the $y$-axis we call it $y$-shiftable. In the degenerate case of equal grid distance, the node is both $x$-shiftable and $y$-shiftable.
(I3) Exactly three of its grid distances are less than $\Delta x / 2$. In this case, the node is shiftable in the direction of the single grid distance.

We call a grid node which is on the boundary shiftable if it is not at a boundary corner, i.e. not at the intersection of two boundary lines, not on the INTERFACE and if the following condition holds: (See Fig. 1.)


Figure 1: Several cases for a shiftable node N
(B) Exactly one of its grid distances along a boundary grid line is less than $\Delta x / 2$. We call the grid node $x$-shiftable if the boundary grid line is parallel to $x$-axis; otherwise we call it $y$-shiftable.

If a node is neither fixed or shiftable, we call it unshiftable.
Hypothesis 4 At each time level during the computation, every grid node is either fixed or shiftable.

For each grid node, there is either one (I1, I3, B) or two (I2) interface POINTs to which it can be shifted. For each interface POINT, there is at most one grid node which can be shifted to it. Boundary grid nodes can be shifted only to boundary interface POINTs and interior (non boundary) grid nodes can be shifted to non boundary interface POINTs only.

The point shifted algorithm consists of shifting all shiftable nodes to interface POINTs with, for example, a choice of the x-direction in case of ambiguity. The POINTs to which no grid nodes are shifted are deleted. We state an obvious result concerning the algorithm.

Proposition 1 Assume Hypotheses 1-4. If a grid node is shifted to the INTERFACE, then it is shifted to a POINT located at an intersection of the INTERFACE with a grid line. The intersection point lies within an open circle of radius $\Delta x / 2$ centered at the original position of the node.

Theorem 1 Assume Hypotheses 1-4. Then the topology of the grid remains unchanged. Each cell area is between $0.5 \Delta x^{2}$ and $2.5 \Delta x^{2}$. Each POINT of the INTERFACE is a shifted grid cell corner and each BOND is a shifted grid cell edge or diagonal.

After the point shift algorithm, it is easily seen from Proposition 1 and Theorem 1 that the length of each $B O N D$ is no more than $\sqrt{5} \Delta x$. Because of the bound on $B O N D$ angles, the $B O N D$ is within a $\mathcal{O}\left(\Delta x^{2}\right)$ displacement to the grid based CURVE segment it approximates. We have the following corollary.

Corollary 1 Assume Hypotheses 1-4. If the grid based CURVE is within $\mathcal{O}\left(\Delta x^{2}\right)$ displacement to the smooth interface curve, then after point shift algorithm, the CURVE is still a $\mathcal{O}\left(\Delta x^{2}\right)$ approximation to it.

### 2.2 Construction of the Space-Time Hexahedra

We require a space-time triangulated interface surface joining the two spatial $I N$ TERFACEs at times $t_{n}$ and $t_{n+1}$. This construction is the major task of the present section.

The point shifted algorithm does not change the rectangular index structure of the mesh. Thus we connect the nodes of a cell $D_{i}^{n}$ at time $t=t_{n}$, to the nodes of its corresponding cell $D_{i}^{n+1}$ at time $t=t_{n+1}$ to form a space-time hexahedron. We call $D_{i}^{n+1}$ the top of the hexahedron and $D_{i}^{n}$ the bottom. If the both cells have not been affected by the point shifted algorithm we call the hexahedron regular, otherwise it is called irregular. We call a hexahedron mixed if the interface passes through its interior; otherwise it is pure. The mixed hexahedra are divided into pure partial hexahedra, and if necessary, these are combined with neighbors to form the finite volume space-time grid suitable for construction of a conservative difference algorithm in Sec. 2.3

Two hexahedra are adjacent if they share a non-trivial surface which is not on the space time interface. It is easy to see that two adjacent hexahedra must be on the same side of the space time interface. We consider INTERFACEs after the point shift algorithm. In this case, each POINT is a displaced, or shifted grid node, and all $B O N D \mathrm{~s}$, connecting adjacent POINTs are edges or diagonals of displaced or shifted grid cells. We also observe that the POINTs $P_{1}$ and $P_{2}$, defined at a common or adjacent time level, that is both at times $t_{n}$ or $t_{n+1}$, or one at time $t_{n}$ and the other at time $t_{n+1}$, share a common space time hexahedron if and only if they are identical, adjacent or diagonally adjacent as shifted grid nodes in space. We say that $P_{1}$ and $P_{2}$ are spatially nearest neighbors in this case. These points are strictly spatially nearest neighbors if they are identical or adjacent grid nodes (diagonal adjacency excluded) in space. Besides all the hypotheses in Section 2.1, we also assume that the CFL number is less than $1 / 2$ throughout this section so that each POINT of the INTERFACE can be propagated a distance less than $\Delta x / 2$.

Hypothesis 5 The CFL number is less than 1/2.

We introduce terminology for Proposition 2 and the supporting Lemmas 4-7. (See Appendix 2.)

Let $B^{n}$ be a BOND connecting adjacent POINTs $P_{1}$ and $P_{2}$ at the time level $t_{n}$. At the time level $t_{n+1} B^{n}$ has been propagated and point shifted to become the INTERFACE polygonal segment $b^{n+1}$ with a left end point $M_{1}$ as the image of $P_{1}$ and a right end point $M_{2}$ as the image of $P_{2}$.

Proposition 2 Assume Hypotheses 1-5. In the ordering of POINTs along $b^{n+1}$, those spatially nearest neighbor to $P_{1}$ only (if any) lie closest to $M_{1}$, followed by POINTs (if any) spatially nearest neighbor to both $P_{1}$ and $P_{2}$, followed by POINTs (if any) spatially nearest neighbor to $P_{2}$ only. The middle set of POINTs necessarily occurs if both of the other two are non empty.

Proposition 3 Assume Hypotheses 1-5. Let $B_{1}$ be a BOND at the time level $t_{n+1}$ connecting adjacent POINTs $P_{1}$ and $P_{2}$. Suppose that adjacent time level $t_{n}$ POINTs are propagated to $B_{1}$. Then those spatially nearest neighbor only to $P_{1}$ (if any) occur first in the INTERFACE order, followed by POINTs (if any) spatially nearest neighbor to both $P_{1}$ and $P_{2}$, followed by POINTs (if any) spatially nearest neighbor only to $P_{2}$. The middle set of POINTs necessarily occurs if both of the other two are non empty.

We construct the space time interface as triangles whose edges joint POINTs on successive time levels $t_{n}$ and $t_{n+1}$, and which are spatially nearest neighbor. Each edge of the triangle is then a surface edge or diagonal, or interior diagonal of a single space time hexahedron. The only further requirement, to avoid tangling of the space time interface, is that the edges preserve order along the interface, as a mapping from $t_{n}$ POINTs onto $t_{n+1}$ POINTs.

The correspondence between interface POINTs is nearly determined by Propositions 2 and 3. There is an alternation between expansion and contraction events in this pairing, as defined below;

Expansion: The INTERFACE segment $b^{n+1}$ defined by propagation and shifting of the $B O N D B$ contains one or more POINTs.

Contraction: A single time level $t_{n+1} B O N D$ contains the images defined by propagation and shifting of one or more consecutive time level $t_{n}$ POINTs.

For an Expansion event, following the terminology of Proposition 2, we connect (introducing triangle edges) $P_{1}$ to all POINTs on $\left[M_{1}, M_{2}\right]$ which are spatially nearest neighbor to $P_{1}$ but not $P_{2}$. We connect to $P_{2}$ all POINTs on $\left[M_{1}, M_{2}\right]$ spatially nearest neighbor to $P_{2}$ but not $P_{1}$. The POINTs, if any, spatially nearest neighbor to both


Figure 2: Construction of space time interface joining the time $t_{n}$ and $t_{n+1}$ interfaces. Upper three frames, expansion case; lower three frames, contraction case. In case 1, all POINTs are spatially nearest neighbor to $P_{1}$, in case 2 to $P_{2}$, and case 3 includes at least one mixed POINT. "x": interface POINT; " o ": image of propagation.


Figure 3: Adjacent events are necessarily of opposite type. "x": interface POINT; "o": image of propagation.


Figure 4: Either $P_{1}$ connects to $P_{4}$ or $P_{2}$ connects to $P_{3}$. "x": interface POINT; "o": image of propagation.
$P_{1}$ and $P_{2}$ are split; those before and including a reference POINT $P_{j}$ are connected to $P_{1}$, while those including $P_{j}$ and after are connected to $P_{2}$. See Fig. 2, upper frames.

For a Contraction event, following the terminology of Proposition 3, we connect $P_{1}$ to the points propagating onto $\left[P_{1}, P_{2}\right]$, which are spatially nearest neighbor to $P_{1}$ but not $P_{2}$. Similarly we connect to $P_{2}$ those spatially nearest neighbor to $P_{2}$ but not $P_{1}$. The remainder, spatially nearest neighbor to both $P_{1}$ and $P_{2}$ are split as above, with a separating point $P_{j}$ connected to both $P_{1}$ and $P_{2}$. See Fig. 2, lower frames.

Two events of the same type can not be adjacent to each other. See Fig. 3. Therefore it remains to show that the triangles from the Expansion and Contraction events join, and complete the space time interface. Assume that the Expansion is to the left of the Contraction. There is a gap between the two sets of triangles only if all Expansion triangles join to the left (case 1) and the contraction triangles join to the right (case 2) as we now assume. The gap is a quadrilateral and one of its diagonals must be added to complete the triangulation. According to Lemma 6 (see Appendix 2), either the right most time $t_{n+1}$ POINT of the Expansion event is spatially nearest neighbor to the left most time $t_{n}$ POINT of the Contraction event, or the right most time $t_{n}$ POINT of the Expansion event is spatially nearest neighbor to the left most time $t_{n+1}$ POINT of the Contraction event, so that we can connect one diagonal pair of POINTs of the quadrilateral to triangulate the gap. See Fig. 4. The case of a Contraction to the left of an adjacent Expansion is similar. In the case that a spatial interface CURVE is topologically equivalent to a circle, each space-time interface event is adjacent to a space-time interface event on the left, and another one on the right. Therefore the above process fully triangularizes the space time surface joining the two CURVEs at the two time levels. If a CURVE has its two end points on the boundary, each of the end points (which becomes a POINT after the point shift
algorithm) must be in the same space hexahedron at time levels $t_{n}$ and $t_{n+1}$ so that we can connect the end point at time levels $t_{n}$ to the end point at time level $t_{n+1}$ to complete the triangulation. This requires that end point should be propagated less than a distance $\Delta x / 2$ along the boundary grid line, which depends on the states near the end point, the angle at which the INTERFACE intersects the boundary and the time step size (or the CFL number) we choose. We formulate this requirement as the following hypothesis:

Hypothesis 6 The CFL number chosen during each time step must ensure that each intersection point of the CURVE with the boundary is propagated less than a distance $\Delta x / 2$ along the boundary grid line.

We summarize the above discussion with the following theorem:
Theorem 2 Assume Hypothesis 1-6. After the above triangulation process, each triangle on the space time interface will be on the face or interior of a point shifted space time hexahedron, with its base being an edge or diagonal of the top (or bottom) cell of the hexahedron and its other vertex being a grid node of the bottom (or top) cell of the hexahedron.

Proof: Each resulting triangle on the space-time interface has an edge, say the base, which corresponds to an interface BOND, i.e., an edge or a diagonal of a cell on $t_{n}$ (or $t_{n+1}$ ), while the other vertex of triangle corresponds to an interface POINT, i.e., a point shifted grid node on $t_{n+1}$ (or $t_{n}$ ). Since the vertex is spatially closest to the other two vertices of the triangle, it is easy to see that all of its three vertices will share a common space-time hexahedron. The proof is complete.

The mixed hexahedron is separated into two parts, each of which lies on one side of the space time interface. These parts are called pure partial hexahedra. We can similarly define a cell to be regular, irregular, pure, mixed or partial. Any partial hexahedra with a trivial top will be merged with an adjacent pure hexahedron or partial hexahedra having a nontrivial top. Recall that two adjacent hexahedra are on the same side of the interface. The merging process can be accomplished as follows:

Merge every pure or partial hexahedron having a nontrivial top with adjacent partial hexahedra having trivial tops which have not been merged elsewhere. Denote the resulting polyhedra the intermediate hexahedra. Merge every intermediate hexahedron repeatively with adjacent partial hexahedron having a POINT top which have not been merged elsewhere. Denote the resulting polyhedra the big hexahedra.

After the merging process, we also call the remaining pure and partial hexahedra big hexahedra for equivalence in the next computation. The following lemma ensures the success of the above algorithm.

Lemma 1 Assume Hypothesis 1-6. If a polyhedron is constructed by merging any number of adjacent partial hexahedra with trivial tops, then the polyhedron will be adjacent to a pure or partial hexahedron.

Proof: At least one non trivial piece (a triangle) of the side surface of the polyhedron is not on the boundary or the space time interface, otherwise the topological structure of the INTERFACE changes during this time step and Hypothesis 1 is violated. The proof is complete.

Theorem 3 Assume Hypothesis 1-6. After the above merging process,
(1) every partial hexahedron with a trivial top will be merged into a big hexahedron with non trivial top;
(2) every big hexahedron has a non trivial top which is a pure cell or a partial cell.

Proof: Let $H$ be a pure or partial hexahedron with trivial top, then it is adjacent to a pure or partial hexahedron from Lemma 1. We can separate it into several cases.
(a) The top of $H$ contains a BOND of the $t_{n+1}$ INTERFACE. Then the BOND must be the edge of a pure or partial cell (say $C$ ) on the same side of the space time interface since the BOND can not be on the boundary according to Hypothesis 1. Therefore $H$ must be adjacent to the pure or partial hexahedron with $C$ as its top and the algorithm will merge all such partial hexahedra into the intermediate hexahedra.
(b) The top of $H$ is a POINT and $H$ is adjacent to a partial hexahedron with a trivial top which consists of BONDs, or to a pure or partial hexahedron with a non trivial top. Then it will be merged either with an intermediate hexahedron (due to (a)) or with a pure or partial hexahedron with a non trivial top.
(c) The top of $H$ is a POINT $P$ and $H$ is adjacent only to partial hexahedra with a POINT top $P$. Let $M_{1}$ be the polyhedron resulting from merging all the adjacent partial hexahedra with a POINT top $P$. Then $M_{1}$ consists of at most four partial hexahedra with the POINT top $P$ since $P$ can belong to at most four pure or partial hexahedra on the same side of the space time interface. From Lemma 1 $M_{1}$ is adjacent to a pure or partial hexahedron with non trivial top or to a partial hexahedron with a trivial top which consists of $B O N D$ s. In the first case $M_{1}$ will be merged with the pure or partial hexahedron with non trivial top. The second case is similar to the previous case (b).

The first statement of the Theorem is proved. The second statement is from the observation that a pure or partial hexahedron with a non trivial top (a pure cell or a partial cell) will merge only with hexahedra with trivial tops. The proof is complete.

Note that the top of a big hexahedron (including the trivial parts) is within the union of the top cell (or partial cell) and its spatially closest neighbors, therefore the total number of pure or partial hexahedra in the big hexahedron is bounded. Actually in most cases of the computation the merging process yields the big hexahedron consisting of two pure or partial hexahedra. The number of pure or partial hexahedra in the big hexahedron could become larger if the radius of curvature of the moving $C U R V E$ is closer to the mesh size.

Note that the merging process does not increase the computational complexity since the net outflux of a big hexahedron is equal to the sum of the net outflux of each pure or partial hexahedron contained in it.

Since after the merging process the big hexahedron will have either a cell or a partial cell as its top, this construction does not change the reconstruction of state functions at the next time level discussed in the next section.

### 2.3 The Reconstruction, Limiter and the Numerical Scheme

Suppose at the time level $t=t_{n}$ we know the approximate state averages on each cell, regular, irregular or partial. We want to reconstruct a piecewise linear state function on these cells with 2 nd order accuracy. The reconstruction of the piece wise linear state function on irregular cells follows [1], with modifications to the limiter and some simplification. Let $D_{i}^{n}$ be a pure cell, regular, irregular, or partial with approximate state average $\mathcal{U}_{i}$ and cell center (centroid) $Y_{i}$, surrounded by any of these types of cells $D_{j}^{n}, D_{k}^{n}, D_{l}^{n}, D_{m}^{n}$ with approximate state averages $\mathcal{U}_{j}^{n}, \mathcal{U}_{k}^{n}, \mathcal{U}_{l}^{n}, \mathcal{U}_{m}^{n}$ and cell centers $Y_{j}, Y_{k}, Y_{l}, Y_{m}$ respectively, on the same side of the INTERFACE. Let $\tilde{\mathcal{U}}_{i}=\mathcal{U}_{i}+(a, b) \cdot\left(X-Y_{i}\right)$ be the 2 nd order accurate linear state function on $D_{i}^{n}$, where $a, b$ are two constants. Choose any two surrounding cells, say $D_{j}^{n}, D_{k}^{n}$ so that $Y_{i}, Y_{j}, Y_{k}$ are not colinear. We can determine $a, b$ by solving the following equation:

$$
\begin{align*}
& \tilde{\mathcal{U}}_{i}\left(Y_{j}\right)=\mathcal{U}_{j}^{n}, \\
& \tilde{\mathcal{U}}_{i}\left(Y_{k}\right)=\mathcal{U}_{k}^{n} . \tag{2}
\end{align*}
$$

Further, for the solution of the above equation to be well conditioned, we require the angle $\theta$ formed by line segment $\overline{Y_{i} Y_{j}}$ and $\overline{Y_{i} Y_{k}}$ to satisfy $0<\theta_{1}<\theta<\theta_{2}<$ $\pi$ where $\theta_{1}, \theta_{2}$ are two constants. We repeat the above procedure until we find all possible solutions, say, $a_{i}, b_{i}$, for all $0 \leq i \leq I$ where $I \leq 4$. Then we set $a=\operatorname{minmod}\left\{a_{1}, \cdots, a_{I}\right\}$ and $b=\operatorname{minmod}\left\{b_{1}, \cdots, b_{I}\right\}$. When there are not enough surrounding cells on the same side of the INTERFACE, we choose $a, b=0$ so that the reconstruction becomes first order.

When $D_{i}^{n}$ is a regular cell surrounded by regular cells, the reconstruction process is simpler. Let the cell center of $D_{i}^{n}$ be $\left(i_{1} \Delta x, i_{2} \Delta y\right)$ with neighboring cell centers

$$
\begin{align*}
& \left\{\left(\left(i_{1}+k_{1}\right) \Delta x,\left(i_{2}+k_{2}\right) \Delta y\right) \mid k_{1}, k_{2}=-1,0,1\right\} . \text { Let } \\
& \qquad \text { xslope }_{\mathrm{i}}=\operatorname{minmod}\left\{\left[\mathcal{U}\left(\left(i_{1}+k_{1}\right) \Delta x,\left(i_{2}+k_{2}\right) \Delta y\right)-\right.\right. \\
&  \tag{3}\\
& \left.\mathcal{U}\left(\left(i_{1}+k_{1}-1\right) \Delta x,\left(i_{2}+k_{2}\right) \Delta y\right)\right] / \Delta x \mid \\
& \left.\quad k_{1}=0,1 ; k_{2}=-1,0,1\right\}
\end{align*}
$$

and

$$
\begin{gather*}
\text { yslope }_{\mathrm{i}}=\operatorname{minmod}\left\{\left[\mathcal{U}\left(\left(i_{1}+k_{1}\right) \Delta x,\left(i_{2}+k_{2}\right) \Delta y\right)-\right.\right. \\
\left.\mathcal{U}\left(\left(i_{1}+k_{1}\right) \Delta x,\left(i_{2}+k_{2}-1\right) \Delta y\right)\right] / \Delta y \mid  \tag{4}\\
\left.k_{1}=-1,0,1 ; k_{2}=0,1\right\},
\end{gather*}
$$

and define

$$
\tilde{\mathcal{U}}_{i}=\mathcal{U}_{i}+\text { xslope }_{\mathbf{i}} \cdot\left(x-i_{1} \Delta x\right)+\text { yslope }_{\mathbf{i}} \cdot\left(y-i_{2} \Delta y\right)
$$

This is clearly a second order reconstruction which is better suited in multiple dimensional problem than operator splitting single line reconstruction (or limiter) for a uniform rectangular grid, because for example an untracked discontinuity in 2D may be in the form of a strip of width between $2 \Delta x$ and $3 \Delta x$. When the strip is almost parallel to and fully covers the line in which the single line reconstruction occurs, one cannot expect the limiter to choose any smooth solutions nearby.

Next we apply the technique in Section 2.2 to generate space-time hexahedra between time levels $t^{n}$ and $t^{n+1}$. Let $H$ be a big hexahedron with top $D^{n+1}$ and bottom $D^{n}$, and triangle sides $\left\{S_{i}\right\}$ with a unit outer normal $n_{i}$ and centroid $Z_{i}$. Notice that some elements of the $\left\{S_{i}\right\}$ may be on the approximate space time interface. Integrating (1) over $H$, we obtain

$$
\begin{equation*}
\left|D^{n+1}\right| U^{n+1}=\left|D^{n}\right| U^{n}-\sum_{i} \int_{S_{i}}(u, f, g) \cdot n_{i} d s \tag{5}
\end{equation*}
$$

Here $\left|D^{n}\right|$ represents the area of $D^{n}$ and similarly $\left|S_{i}\right|$ is the area of $S_{i}$. The numerical scheme can be written as

$$
\begin{equation*}
\left|D^{n+1}\right| \mathcal{U}^{n+1}=\left|D^{n}\right| \mathcal{U}^{n}-\sum_{i}\left|S_{i}\right|\left(\tilde{\mathcal{U}}_{i, m}, f\left(\tilde{\mathcal{U}}_{i, m}\right), g\left(\tilde{\mathcal{U}}_{i, m}\right)\right) \cdot n_{i}, \tag{6}
\end{equation*}
$$

where $\tilde{\mathcal{U}}_{i, m}$ can be calculated as follows: First use a Cauchy-Kowalewski procedure on the reconstructed state function on each side of $S_{i}$ to get 2nd order approximate states at $Z_{i}$ on the respective side of $S_{i}$, say $\mathcal{U}_{i, l}$ and $\mathcal{U}_{i, r}$. If $S_{i}$ is not on the tracked space time interface, we can simply use a Riemann solver, say $R$, to get the middle state on $S_{i}$, i.e.

$$
\tilde{\mathcal{U}}_{i, m}=R\left(\mathcal{U}_{i, l}, \mathcal{U}_{i, r}\right) .
$$

If $S_{i}$ is on the tracked space time interface, we use the Riemann solver to get the left and the right side states $\tilde{\mathcal{U}}_{i, l}$ and $\tilde{\mathcal{U}}_{i, r}$ on the wave we are supposed to track, and
the wave speed $\nu_{i}$. Then $\tilde{\mathcal{U}}_{i, m}$ in (6) can be replaced by either $\tilde{\mathcal{U}}_{i, l}$ or $\tilde{\mathcal{U}}_{i, r}$, depending on whether $l$ or $r$ is located within $H$ or not. Also the $n_{i}$ in (6) should be replaced by $\tilde{n}_{i} /\left|\tilde{n}_{i}\right|$, where $\tilde{n}_{i}=\left(-\theta_{i} \nu_{i} \sqrt{n_{i x}^{2}+n_{i y}^{2}}, n_{i x}, n_{i y}\right), n_{i}=\left(n_{i t}, n_{i x}, n_{i y}\right)$ and $\theta_{i}$ is a sign function which is 1 if the tracked wave from the Riemann solver is in the direction of $\left(n_{i x}, n_{i y}\right),-1$ otherwise. Note that $\tilde{n}_{i}$ is normal direction of the tracked space time wave from the Riemann solver, therefore this modification ensures that the Rankine-Hugoniot condition is satisfied.

The finite volume difference algorithm constitutes a flux through each boundary of the full, partial and big hexahedron. Since the flux through a boundary face of the hexahedron is identical when viewed from either side of the face, we have
Theorem $4 \sum_{\text {cells }}\left|D^{n}\right| \mathcal{U}^{n}$ in the finite volume difference scheme is conserved so that its increment over any time interval is equal to the net influx at the boundary.

Away from the INTERFACE the scheme is clearly a second order scheme. For the cells along the INTERFACE, its local truncation error is one order lower than in the 1 D case since we use a piece wise linear INTERFACE and its local displacement error is $\mathcal{O}\left(\Delta x^{2}\right)$. The scheme is one order better than untracked schemes, which typically have $\mathcal{O}(1)$ local truncation error at the untracked fronts.

Theorem 5 Suppose the exact space time interface and the solution on either side of it are smooth. Then the $L_{\infty}$ local truncation error is $\mathcal{O}(\Delta x)$ for cells adjacent to the INTERFACE.

Proof: Let the INTERFACE at $t_{n}$ be the interpolation of the exact interface and let $H$ be a big hexahedron adjacent to the approximate space time interface. We apply the finite volume scheme to obtain the approximate state average $\mathcal{U}_{i}^{n+1}$ at the time level $t_{n+1}$, with top $T$ and bottom $B$ and side boundaries $\left\{S_{i}\right\}$, where $S_{i}$ is a triangle. The INTERFACE at time $t_{n+1}$ has an $\mathcal{O}\left(\Delta x^{2}\right)$ displacement from the exact interface according to Corollary 1. The exact space time interface will cut $H$ into two pieces. Let $H_{1}$ be the piece on the same side of the interface with $H$. Let $T_{1}, B_{1}$, and $S^{1}$ be the top, bottom and side boundaries of $H_{1}$ respectively. Let $U_{T_{1}}^{n+1}, U_{B_{1}}^{n}$ be the exact state averages over $T_{1}$ and $B_{1}$ respectively. Choosing $\mathcal{U}_{B}^{n}=U_{B_{1}}^{n}$, we want to show that $U_{T_{1}}^{n+1}-\mathcal{U}_{T}^{n+1}=\mathcal{O}(\Delta x)$. In fact from (6),

$$
\begin{equation*}
\left.|T| \mathcal{U}_{T}^{n+1}=|B| \mathcal{U}_{B}^{n}-\sum_{i}\left|S_{i}\right| \mid \tilde{\mathcal{U}}_{i, m}, f\left(\tilde{\mathcal{U}}_{i, m}\right), g\left(\tilde{\mathcal{U}}_{i, m}\right)\right) \cdot n_{i} . \tag{7}
\end{equation*}
$$

The exact solution satisfies

$$
\begin{equation*}
\left|T_{1}\right| U_{T_{1}}^{n+1}=\left|B_{1}\right| U_{B_{1}}^{n}-\int_{S^{1}}(u, f(u), g(u)) \cdot n d s \tag{8}
\end{equation*}
$$

Note that $|B| \mathcal{U}_{B}^{n}-\left|B_{1}\right| U_{B_{1}}^{n}=\mathcal{O}\left(\Delta x^{3}\right)$ due to the interpolation error from the INTERFACE at time $t_{n}$. Also the numerical flux in (7) approximates the exact flux in (8) to at least $\mathcal{O}\left(\Delta x^{3}\right)$. In fact when $S_{i}$ is not on the approximate space time interface this is easily seen since $\int_{S_{i}}(u, f, g) \cdot n_{i} d s=\left|S_{i}\right|(u, f, g)\left(Z_{i}\right) \cdot n_{i}+\mathcal{O}\left(\Delta x^{4}\right)$. When $S_{i}$ is on the approximate space time interface. Because the approximate space time interface has an $\mathcal{O}\left(\Delta x^{2}\right)$ displacement error relative to the exact one, the difference between their respective areas is of $\mathcal{O}\left(\Delta x^{3}\right)$ due to the smoothness assumption of the exact space time interface and the area of $\bigcup S_{i}$ being $\mathcal{O}\left(\Delta x^{2}\right)$. Also the choices of $\tilde{\mathcal{U}}_{i, m}$ and $n_{i}$ in (7) ensure that $\left(\tilde{\mathcal{U}}_{i, m}, f\left(\tilde{\mathcal{U}}_{i, m}\right), g\left(\tilde{\mathcal{U}}_{i, m}\right)\right) \cdot n_{i}$ in (7) is a first order approximation to the integrand in (8) at any point within an $\mathcal{O}(\Delta x)$ distance from the centroid $Z_{i}$ of $S_{i}$. Therefore we have

$$
\begin{align*}
U_{T_{1}}^{n+1}-\mathcal{U}_{T}^{n+1} & =\left(\left|T_{1}\right| U_{T_{1}}^{n+1}-|T| \mathcal{U}_{T}^{n+1}\right) /|T|+U_{T_{1}}^{n+1}\left(\left(|T|-\left|T_{1}\right|\right) /|T|\right)  \tag{9}\\
& =\mathcal{O}(\Delta x)
\end{align*}
$$

The proof is complete.

## 3 Appendix 1: Proof of Theorem 1 (§2.1)

Lemma 2 Assume Hypotheses 1-4. If a cell edge connecting grid nodes $N_{1}$ and $N_{2}$ intersects a CURVE at a point $P$, then either $N_{1}$ or $N_{2}$ will be shifted to $P$, or to an intersection point of the CURVE with another edge adjacent to $P$ along the CURVE.

Proof: Without loss of generality, suppose the cell edge $l$ connecting $N_{1}$ and $N_{2}$ is parallel to the $x$-axis. At least one of the nodes must have grid distance less than or equal to $\Delta x / 2$, say node $N_{1}$ which is shiftable. We first suppose that $N_{1}$ is not on the boundary. If $N_{1}$ belongs to (I1) then $N_{1}$ is $x$-shiftable and therefore will be shifted to $P$.

If (I2) or (I3) is true for $N_{1}$, we consider two cases:
Case $1 N_{1}$ is $x$-shiftable, the result is the same as in (I1) during Step 1;
Case $2 N_{1}$ is only $y$-shiftable, then $N_{1}$ will be shifted to the intersection point adjacent to $P$ (along the CURVE).

If $N_{1}$ is on the boundary it will be shifted along the boundary to $P$ or to the POINT adjacent to $P$ along the CURVE. The proof is complete.

Lemma 3 Assume Hypotheses 1-4. Let $Q$ be the union of closed mesh cells with connected interior. Suppose a CURVE enters $Q$ at $P_{1} \in \partial Q$ and leaves $Q$ at $P_{2} \in \partial Q$ with the CURVE segment $\left[P_{1}, P_{2}\right] \subset Q$. Let $P_{0}$ be the intersection point between the CURVE and a cell edge just prior to $P_{1}$ if it exists; otherwise let $P_{0}=P_{1}$. Similarly let $P_{3}$ be the intersection point between the CURVE and a cell edge just after $P_{2}$ if it exists; otherwise let $P_{3}=P_{2}$. Then
(1) After the point shift algorithm, only the nodes originally in $Q$ are on the CURVE segment $\left[P_{1}, P_{2}\right]$.
(2) At least one such node will be on the CURVE segment $\left[P_{0}, P_{3}\right]$.
(3) If a grid node originally in the interior of $Q$ is shifted to the segment $\left(P_{1}, P_{2}\right)$, then after the point shift algorithm it will be adjacent (along the CURVE) only to the grid nodes originally in $Q$.

Proof: From Proposition 1 and Lemma 2 it follows that after the point shift algorithm, at least one grid node originally in $Q$ will be on the $C U R V E$ segment $\left[P_{0}, P_{3}\right]$ to prove statement (2). Also from Proposition 1 we know that only nodes originally in $Q$ will be on the CURVE segment [ $P_{1}, P_{2}$ ] proving statement (1). Statement (3) is obvious.

The proof is complete.
Proof of Theorem 1 By drawing an open circle of radius $\Delta x / 2$ centered at the original position of each grid node, according to Proposition 1 we find all the possible positions of each node after point shift algorithm. Note that these circles are disjoint. Because the original interface is grid based, and thus consists of straight line segments between its crossings with grid lines, the new grid does not introduce any new intersection between edges and thus the topology of the grid remains unchanged.

Proposition 1 combined with the fact that the shift is along grid lines only gives the upper bound of the cell area. Hypotheses 2 implies that at most two nodes of a cell can be shifted to the INTERFACE within its cell boundary by the point shift algorithm (in other words, at least two nodes of the cell will remain fixed or be shifted outside the cell), which gives the lower bound of the cell area.

The INTERFACE passes through displaced grid cell corners only, by direct construction. We further assert that if a point shifted interface $B O N D$ connects two nodes, then the two nodes must share a common cell. In fact, if an interior node $N$ is on a CURVE after the point shift algorithm, from the Hypothesis 3 the CURVE must intersect the boundary of the square of side $2 \Delta x$ centered at the original position of $N$. Therefore from Lemma 3, after the point shift algorithm, $N$ must be adjacent only to the nodes originally from the square, each of which share a common cell with $N$. If $N$ is a boundary node that is on a CURVE after point shift algorithm, then it follows from Hypothesis 1 that $N$ will be adjacent to an interior node along the CURVE, which returns to the previous case. The proof is complete.

## 4 Appendix 2: Proof of Propositions 2 and 3 (§2.2)

We introduce Lemmas 4-7 following the terminology of Proposition 2.

Lemma 4 All POINTs on $b^{n+1}$ are strictly spatially nearest neighbors with either $P_{1}$ or $P_{2}$.

Proof: The CFL number is less than $\frac{1}{2}$. Thus any point on $B^{n}$ can propagate at most a distance less than $\Delta x / 2$. Shifting at each of the time levels $t_{n}$ and $t_{n+1}$ moves the grid nodes at most $\Delta x / 2$, and only along grid lines. Therefore only grid nodes with a zero or unit lattice displacement from $P_{1}$ or $P_{2}$ in mesh index space can be shifted to lie on $b^{n+1}$. The lemma is proved.

Lemma 5 Let $M_{1}$ belong to a BOND connecting adjacent Points $P_{3}$ and $P_{4}$. Then either $P_{1}$ and $P_{3}$ are spatially nearest neighbors or $P_{1}$ and $P_{4}$ are.

Proof: Let $D$ be a closed square of side $2 \Delta x$ centered at the unshifted original position of grid node which is shifted at time $t_{n}$ to be the POINT $P_{1}$. Then $M_{1}$ is in the interior of $D$. According to the first two statements of Lemma 3, $M_{1}$ must be adjacent (along the INTERFACE) to at least one of the grid nodes originally contained in $D$ before being shifted. In other words, at least one of $P_{3}$ and $P_{4}$ is originally in $D$ before being shifted. The grid nodes originally in $D$ are all spatially nearest neighbor to $P_{1}$. Thus the proof is complete.

Lemma 6 Assume that at least one of two adjacent POINTs $P_{3}, P_{4}$ at time level $t_{n+1}$ lie on $b^{n+1}$. Suppose $\left[P_{1}, P_{2}\right]$ and $\left[P_{3}, P_{4}\right]$ have the same orientation relative to the INTERFACE, and that the pairs $P_{1}, P_{3}$ and $P_{2}, P_{4}$ are spatially nearest neighbors. Then either $P_{1}, P_{4}$ or $P_{2}, P_{3}$ are spatially nearest neighbors.

Proof: The points $P_{1}, P_{2}, P_{4}, P_{3}$ in cyclic order form a four sided loop of spatially nearest neighbor grid nodes, when projected to a common time, and viewed in grid index space. See Fig. 5. Moreover, the points $P_{3}$ and $P_{4}$ belonging to $b^{n+1}$ are strictly spatially nearest neighbor to either $P_{1}$ or $P_{2}$ by Lemma 4. This fact either completes the proof directly, or it forces one side of the loop to be a unit lattice distance (not a diagonal) at most. Each side of the loop is a single point, a unit lattice line or a unit lattice diagonal. The loop thus consists of a single line (multiply covered), a unit triangle (one side of the loop reduces to a point), a pair of adjacent unit triangles, a unit cell or a unit parallelogram (displaced by one lattice site from being a cell). The doubly displaced parallelogram of Fig. 6 is excluded. We are to prove that one of the diagonals of this loop is also spatially nearest neighbor. For the unit parallelogram, the shorter diagonal is a unit lattice diagonal, and its end points are thus spatially nearest neighbors. The other cases are elementary. The proof is complete.


Figure 5: Grid nodes spatially nearest neighbor to a pair of nearest neighbor nodes.

Lemma 7 Assume that $b^{n+1}$ is contained in a BOND connecting adjacent POINTs $P_{3}$ and $P_{4}$, Suppose $\left[P_{1}, P_{2}\right]$ and $\left[P_{3}, P_{4}\right]$ have the same orientation relative to the $I N$ TERFACE, and that the pairs $P_{1}, P_{3}$ and $P_{2}, P_{4}$ are both spatially nearest neighbors. Then either $P_{1}, P_{4}$ are spatially nearest neighbors or $P_{2}, P_{3}$ are.

Proof: As in the proof of Lemma $6, P_{1}, P_{2}, P_{4}, P_{3}$ form a loop of spatially nearest neighbor grid nodes. If the $\operatorname{BOND}\left[P_{1}, P_{2}\right]$ is the edge of a cell then one side of the loop is strictly spatially nearest neighbor, and the proof follows that of Lemma 6. If the $B O N D\left[P_{1}, P_{2}\right]$ is the diagonal of a cell, see the right frame of Fig. 5. Fig. 6 shows the only case in which neither $P_{1}, P_{4}$ nor $P_{2}, P_{3}$ are spatially nearest neighbors.

The two circles contain the possible ranges of $P_{1}$ and $P_{2}$ respectively after the point-shift algorithm at the time $t_{n}$ and propagation only at the time $t_{n+1} . A, B, C$, $D, E, F, G, H$ are the midpoints on the cell edges starting at $P_{3}$ and $P_{4}$. In order for $\left[P_{3}, P_{4}\right]$ to be an interface $B O N D$ as described in the proposition at $t_{n+1}$ after the point-shift algorithm, an INTERFACE segment $\gamma$ at time $t_{n+1}$ before the point-shift algorithm has to start from $\overline{A B} \bigcup \overline{C D}$, pass the circle centered at $P_{1}$, then pass the circle centered at $P_{2}$, and end up in $\overline{E F} \bigcup \overline{G H}$. Therefore $\gamma$ has to intersect the edges of some cells and cause the grid nodes of the cells to be shifted on $\gamma$ between POINTs $P_{3}$ and $P_{4}$. (See Lemma 2.) This fact violates the assumption that $P_{3}$ and $P_{4}$ are adjacent on the INTERFACE. The proof is complete.

Proof of Proposition 2 The INTERFACE, in mesh index space, traces a polygonal path joining nearest neighbor mesh nodes, and $b^{n+1}$, as a segment of this INTERFACE, does the same. The two extreme sets, of POINTs spatially nearest neighbor to $P_{1}$ but not to $P_{2}$ and the reverse set, while being spatially nearest neighbor to one of $P_{1}$ or $P_{2}$ are not adjacent. See Fig. 5. Thus $b^{n+1}$ cannot pass from the first


Figure 6: A configuration in which $P_{1}, P_{4}$ and $P_{2}, P_{3}$ are not spatially nearest neighbors.
set to the last without passing through the intermediate set. This proves the final statement of the Proposition.

The rest of the Proposition is an direct consequence of Lemma 4 provided that we can eliminate the following twisted nearest neighbor possibility: there are two POINTs $P_{3}, P_{4}$ on $b^{n+1},\left[P_{3}, P_{4}\right]$ having the same orientation with $\left[P_{1}, P_{2}\right]$, so that $P_{3}$ is a spatially nearest neighbor to $P_{2}$ but not with $P_{1}$ and $P_{4}$ is a spatially nearest neighbor to $P_{1}$ but not with $P_{2}$. In fact if it is true, according to Lemma $4, P_{3}$ must be a strictly spatially nearest neighbor to $P_{2}$ and $P_{4}$ be a strictly spatially nearest neighbor to $P_{1}$. Let us consider the case that $B$ is a diagonal of a cell, the other case being similar and simpler. Drawing a circle of radius $(1-\epsilon) \Delta x(0<\epsilon \ll 1)$ centered at the original positions of $P_{1}$ and $P_{2}$, say $O_{1}, O_{2}$ respectively, we find all the possible positions of $M_{1}, M_{2}$ respectively. See Fig. 7.

Let $N_{1}, N_{2}$ denote the nodes which are strictly spatially nearest neighbors to $P_{1}$ but not with $P_{2}$ and $N_{3}, N_{4}$ denote the nodes which are strictly spatially nearest neighbors to $P_{2}$ but not with $P_{1}$. Note that before the point shift algorithm at $t_{n+1}$, the propagation image of the $B O N D B$ is a straight line which has to start from area $O_{1}$, visit the $\Delta x / 2$ neighborhood of $N_{3}$ or $N_{4}$ (in order for it to be shifted to $b^{n+1}$ ), then visit the $\Delta x / 2$ neighborhood of $N_{1}$ or $N_{2}$, and finally end up in $O_{2}$, which is impossible. The proof is complete.


Figure 7: Nodes strictly spatially nearest neighbor to $P_{1}$ but not $P_{2}$ and the converse.

Lemma 8 Let $P_{1}, P_{2}$ be two (not necessarily adjacent) time level $t_{n}$ POINTs which are propagated and shifted to an interface BOND defined by adjacent time level $t_{n+1}$ POINTs $P_{3}$ and $P_{4}$. Assume that $\left[P_{1}, P_{2}\right]$ and $\left[P_{3}, P_{4}\right]$ have the same orientation relative to the INTERFACE. Then if $P_{3}$ is spatially nearest neighbor with $P_{2}$ but not $P_{1}, P_{4}$ can not have the reverse property of being spatially nearest neighbor with $P_{1}$ but not $P_{2}$.

Proof: We only consider the case that $\left[P_{3}, P_{4}\right]$ is the diagonal of cell, say $C$, after point-shifted algorithm. The other case that $\left[P_{3}, P_{4}\right]$ is the edge of a cell is similar and simpler. There are 5 grid nodes which are spatially nearest neighbor with $P_{3}$ but not with $P_{4}$. Also there are 5 nodes which are spatially nearest neighbor with $P_{4}$ but not with $P_{3}$. See Fig. 8. By drawing a circle of radius $(1-\epsilon) \Delta x(0<\epsilon \ll 1)$ centered at each of these 10 nodes we find all the possible positions of each these nodes after point-shift algorithm at time level $t_{n}$ and after propagation at time level $t_{n+1}$. Let the union of the first 5 circles be $O_{1}$ and the union of the second 5 circles be $O_{2}$. $A, B, C, D, E, F, G, H$ are the midpoints on the respective cell edges which limits the allowed time level $t_{n+1}$ shifting of $P_{3}$ and $P_{4}$. In order for the INTERFACE at the time $t_{n+1}$ before the point-shift algorithm to violate the above proposition the INTERFACE has to start from $\overline{A B} \bigcup \overline{C D}$, visit $O_{2}$, then $O_{1}$ and finally end up in $\overline{E F} \bigcup \overline{G H}$. The $t_{n+1}$ propagated INTERFACE will inevitablely intersect edges and cause grid nodes other than $P_{3}$ and $P_{4}$ to be shifted to the INTERFACE between


Figure 8: Nodes spatially nearest neighbor with one of $P_{3}, P_{4}$ but not the other.
them after the point-shift algorithm. (See Lemma 2.) This is a contradictory to the assumption that $P_{3}$ and $P_{4}$ are adjacent on the INTERFACE. The proof is complete.

Proof of Proposition 3 This is a corollary of Lemmas 5, 7 and 8.

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