



An introduction to Adaptive Mesh Refinement (AMR): Numerical Methods and Tools

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AMR - Introduction

- Solving Partial Differential Equations (PDEs)
 - PDEs solved using discrete domain
 - Algebraic equations estimate values of unknowns at the mesh points
 - Resolution/Spacing of mesh points determines error
 - Initial Solution and Boundary condition are needed
- Goal of grid adaptivity:
 - tracking features much smaller than overall scale of the problem providing adequate higher spatial and temporal resolution where needed.



AMR - Introduction



Uniform meshes

- High resolution required for handling difficult regions (discontinuities, steep gradients, shocks, etc.)
- <u>Computationally extremely costly</u>

Adaptive Mesh Refinement

- Start with a coarse grid
- Identify regions that need finer resolution
- Superimpose finer sub-grids only on those regions Increased computational savings over a static grid approach.
- Increased storage savings over a static grid approach.
- Complete control of grid resolution, compared to the fixed resolution of a static grid approach.

AMR makes it feasible to solve problems that are intractable on uniform grid

AMR - Applications



- CFD Astrophysics Climate Modeling Turbulence Mantle Convection Modeling
- Combustion
- Biophysics
- and many more

Demo of a Shock wave passing over a step function (wind tunnel with a step), rendered using the FLASH code. Courtesy of the Univ. of Chicago, Flash Code group



mesh distortion







mesh distortion







mesh distortion





mesh distortion
point-wise structured (tree-based)
refinement





mesh distortion
point-wise structured (tree-based)
refinement





 mesh distortion
 point-wise structured (tree-based) refinement





mesh distortion
 point-wise structured (tree-based)
 refinement
 block structured





mesh distortion
 point-wise structured (tree-based)
 refinement
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mesh distortion
 point-wise structured (tree-based)
 refinement
 block structured:





mesh distortion
 point-wise structured (tree-based)
 refinement
 block structured:



data blocks are created so

Courtesy of Dr. Andrea Mignone, University of Turin

that the same stencil can be

used for all points and no special treatment is required.

High level objects that encapsulate the functionality for AMR and its parallelization are independent of the details of the physics algorithms and the problem being solved.

Simplifies the process of adding/replacing physics modules as long as they adhere to the interface requirements.

Existing Frameworks

- PARAMESH <u>http://www.physics.drexel.edu/~olson/paramesh</u>
- SAMRAI <u>https://computation.llnl.gov/casc/SAMRAI/</u>
- p4est <u>http://www.p4est.org/</u>
- Chombo <u>https://commons.lbl.gov/display/chombo/Chombo</u>
- and many more



Block Numbering



•All the grid blocks are related to one another as the nodes of a tree.

•The starting block is called root block, and the blocks with an higher resolution are called leaf blocks.

•When a leaf block is designated for refinement, it spawns 2 child blocks in 1D, 4 child blocks in 2D or 8 child blocks in 3D, and the original block is called mother (or parent) block.

•These child blocks cover the same physical line, area or volume as their parent but with twice the spatial resolution.

•Usually it is helpful to use a particular numbering algorithm (see next slides).

Typical grid hierarchy



Each block has a fixed number of grid points
Each block can be divided into 2^{ndim} sub-blocks
Blocks are distributed between processes minimizing communications (see next slides)

An Example:

•6 x 4 grid is created on each block

•The numbers assigned to each block designate the blocks location in the quadtree

•The numbers assigned to each block designate the blocks location in the quadtree

Block ordering



•Usually, the most used block ordering algorithm is Morton (or Z) ordering.

It is particularly useful in order to:
Optimize the usage of cache memory;

•Optimize ghost cells communications between process (see next slide);



Block Structure



Usually, each block is composed by: •standard cells

• ghost cells

In Fortran, the indexes starts with 1 and ends with $N_{(X \text{ or } Y \text{ or } Z)} + 2^*$ (number of ghost cells) In C, the indexes starts whit 0 and ends $N_{(X \text{ or } Y \text{ or } Z)} + 2^*$ (number of ghost cells) -1

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- ghost zones values need to be filled before integration;
- Patches at the same level are synchronized;
- Physical boundaries are imposed externally;
- Fine-Coarse and Coarse-Fine interface need interpolation / averaging
- Integration proceeds as for the single-grid case



Ghost cells communications



Non-conservative linear interpolation. The pink histogram and red circles indicate data on the parent grid block. The dashed line and empty circles show how linear interpolation would prolong this data onto a child grid with an even number of grid cells. The yellow area indicates the misconservation. Conservative interpolation replaces the point labeled P with the green circle so that the area under the curve is now the same as the pink shaded area.



When we pass the ghost cells to the adjoining blocks, if these blocks have different resolutions we must modify the data.

The most simple (and used) method is the interpolation method:

•If we must pass the ghost cells to a block with higher resolution we can use the linear interpolation to artificially increase the resolution.

•If we must pass the ghost cells to a block with lower resolution we can average the data in order to have the same resolution.

Pros:

•Easy to implement

•It is possible to use many different kind of interpolation (linear, quadratic, and so on) increasing precision

Cons: •Non-conservative

Passing ghost cells to neighbors blocks



From Paramesh User Guide



Flux conservation: It is possible to ensure flux conservation after the interpolation checking the equation: f A + f A + f A - F = A

 $f_1\dot{A}_1 + f_2A_2 + f_3A_3 + f_4A_4 = F_{Tot}A_{Tot}$

Passing ghost cells to neighbors blocks



Circulation integral control: It is possible also to check the value of some physical quantity at the edges of the cells



NOTE: Both these three methods are usable in order to change the resolution of the blocks.

Particular Geometries



When we have a non symmetric computational domain many different approach can be used. For a rectangular domain:

We can have different number of points per block on x and y directions (dx = dy)
We can have different number of points on x and y directions (dx ≠ dy)
We can use more blocks on the x directions , and 1 block on x direction (same resolution on x and y, and more parallelizable)

If we have more complicate computational domains, we can always use more blocks in order to fully cover the whole domain.

fill data, level 0





- fill data, level 0
- find where refinement is needed;





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- group cells into patches according to the "grid efficiency"





- fill data, level 0
- find where refinement is needed;
- group cells into patches according to the "grid efficiency"
- refine and ensure proper nesting





- fill data, level 0
- find where refinement is needed;



Little more background on AMR



Refinement structure can be represented using a quad-tree (2D)/ oct-tree (3D)

An important condition in AMR

Refinement levels of neighboring blocks differ by ±1 Note: This is generally true, but Chombo library allow more than 1 refinement level discrepancy.

Traditional Approach - Parallel Implementations



A set of blocks assigned to a process Use space-filling curves for load balancing



Traditional Approach - Disadvantages

•Adaptive mesh restructuring:

Tree metadata replicated on each process

 ✓ Required memory increases with # of cores
 ✓ Memory can became a problem if we use more than 10⁵ cores (and more than 10⁶ boxes)

 Level-by-level restructuring

 Ripple propagation
 Step needed to propagate restructuring ∝ level of refinement (d)



•Load Balancing

•Memory needed \propto Number of blocks used

Time needed ∝ Number of blocks used

•Currently for 3D problems with less than 10⁶ boxes standard AMR library scales up to few tens of thousands of cores

•This is a serious problem considering that next generation supercomputers will require the use of many hundreds of thousands of cores



Improving AMR: Possible strategies

- 1. Compress tree metadata
 - •Already implemented in the last versions of CHOMBO, PARAMESH and SAMRAI libraries
- Rewrite the algorithm for coarse-fine interpolation in order to minimize communications
 •Already implemented in the last versions of CHOMBO, PARAMESH libraries
 •Using these first two methods it is possible to scale up to 2x10⁵ cores using 10⁷ grid cells
- Use a distributed memory version for tree metadata
 •Currently Langer at al are working on the implementation of this algorithm on CHARM++



Some additional information about PARAMESH

•Written in Fortran 90

•Easy to implement on a existing code

•Support many geometries (Cartesian, cylindrical, spherical, from 1D to 3D)

•Refinement levels of neighbouring blocks differ by ±1

•Compatible with hdf5 format

•Some simple routine are already written by the authors of the library in order to save the data and the grid structure into Fortran binary format, and hdf5 format.

•Easy visualization of the results using many external programs (e.g. visit)



Some additional information about CHOMBO

•Written in C

- •Easy to implement on a existing code
- •Support many geometries (Cartesian, cylindrical, spherical, from 2D to 3D)
- •Compatible with hdf5 format
- •Easy visualization of the results using many external programs (e.g. visit)



Example: 2D Blast Wave



Example: 3D Rayleigh-Taylor



Thank you for attention

