Adaptive Mesh Refinement for conservative systems: multi-dimensional efficiency evaluation

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Abstract

Obtainable computational efficiency is evaluated when using an Adaptive Mesh Refinement (AMR) strategy in time accurate simulations governed by sets of conservation laws. For a variety of 1D, 2D, and 3D hydro- and magnetohydrodynamic simulations, AMR is used in combination with several shock-capturing, conservative discretization schemes. Solution accuracy and execution times are compared with static grid simulations at the corresponding high resolution and time spent on AMR overhead is reported. Our examples reach corresponding efficiencies of 5 to 20 in multidimensional calculations and only 1.5-8~% overhead is observed. For AMR calculations of multi-dimensional magnetohydrodynamic problems, several strategies for controlling the $\nabla \cdot \mathbf{B} = 0$ constraint are examined. Three source term approaches suitable for cell-centered **B** representations are shown to be effective. For 2D and 3D calculations where a transition to a more globally turbulent state takes place, it is advocated to use an approximate Riemann solver based discretization at the highest allowed level(s), in combination with the robust Total Variation Diminishing Lax-Friedrichs method on the coarser levels. This level-dependent use of the spatial discretization acts as a computationally efficient, hybrid scheme.

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1 Introduction

Adaptive Mesh Refinement (AMR), an h-refinement method for solutionadaptive computations, is used in an ever increasing number of plasma physical and astrophysical simulations. The AMR algorithm and, specifically, all complications associated with ensuring global conservation properties when dealing with hyperbolic conservative systems, have been addressed in the work by Berger and Colella (7). In that pioneering paper, the authors stress that "...it seems possible to implement a general code where the number of dimensions is input". We make use of such a general code in all simulations presented here, where both the dimensionality and the particular system of conservation laws is selected in a pre-processing stage. The resulting versatility in the applications allows us to assess quantitatively how much can be gained from employing AMR in 1D, 2D, or 3D hydro- and magnetohydrodynamic (MHD) simulations, for a variety of spatial discretizations. Our coding effort benefits from the expertise gained through many general purpose and problem-tailored AMR software packages, which have been in active use and continued development for more than a decade. E.g., AMRCLAW (29) by LeVeque and coworkers combines AMR with a selection of time-dependent hyperbolic systems of partial differential equations, including both conservative and nonconservative systems. Berger and LeVeque (8) recently demonstrated the use of multi-dimensional wave propagation algorithms (30; 28) in 2D adaptive computations on Cartesian and logically rectangular, curvilinear grids. A related development effort by Walder and Folini (49; 50) is AMRCART, a software package which intends to encompass 3D adaptive MHD scenarios. Dimension-independent library approaches for AMR simulations include the publicly available PARAMESH toolkit by MacNeice et al. (31), as well as C++ class framework approaches such as the AmrLib extension to the BoxLib library (10). The code used here is the most recent extension to the Versatile Advection Code (see http://www.phys.uu.nl/~toth), briefly discussed in section 2.1.

Our AMR implementation differs somewhat from the original algorithm in (7) and these differences will be mentioned in section 2.2. For several multidimensional calculations, we report on computational efficiency and solution accuracy. We beneficially combine a fully upwind scheme on the highest AMR level with a Total Variation Diminishing Lax-Friedrichs discretization on all other levels. For 2D hydrodynamic and magnetohydrodynamic problems, we thereby typically gain a factor of 10 in execution time as compared to the equivalent high resolution static grid result. Our example simulations cover 1D Riemann problems (Section 3.1), a 2D advection problem (Section 3.2.1), and both shock-dominated and more turbulently evolving hydro- and magnetohydrodynamic multidimensional cases. Tables 3-4 collect our findings as presented in Section 3.

As our ultimate aim is to perform grid-adaptive, realistic 3D MHD simulations to investigate general plasma physical and astrophysical scenarios, we pay particular attention to strategies for controlling the solenoidal character of the magnetic field. In section 2.3, related efforts in AMR MHD simulations are discussed in relation to their specific treatment of this constraint. Recently, Balsara (2) and independently Tóth and Roe (47) introduced a divergence-free reconstruction and prolongation strategy for multi-dimensional AMR MHD simulations which involves staggered magnetic field components. In this paper, we evaluate three alternative, simple approaches with cell-centered $\bf B$ components where $\nabla \cdot \bf B = 0$ is maintained to truncation error by the addition of suitable source terms. Quantitative comparisons between AMR MHD calculations and high resolution static grid runs in section 3.2 demonstrate their effectiveness.

2 Algorithm

2.1 Versatile Advection Code, equations and solvers

The Versatile Advection Code (VAC), initiated by Tóth (44), is specifically designed for simulating dynamics governed by a system of (near-) conservation laws. It's versatility resides in the choices offered in the geometry of the grid, the dimensionality, the physical application (including Euler and MHD equation modules), the spatial (46) and temporal discretization (45; 24), and the computer platform (23).

VAC uses various second order shock-capturing numerical algorithms: the Flux Corrected Transport (FCT) scheme (11), the Lax-Wendroff type Total Variation Diminishing (TVD) (19) and the TVD-MUSCL (48) schemes (TVDMU) with Roe-type approximate Riemann solvers (38; 39), and the TVD Lax-Friedrichs (TVDLF) method (52). For the TVD type schemes different slope limiters are available, including the most robust *minmod*, and the sharper *monotonized central* (MC, also referred to as Woodward) limiter. For exact specifications of the algorithms, see (46) and references therein.

In multi-dimensional MHD simulations one needs to control the numerical value of $\nabla \cdot \mathbf{B}$. VAC provides several algorithms, such as the eight-wave scheme (36), the projection scheme (12), and several types of constrained transport and central difference type discretizations, which were recently described and evaluated for static grid simulations by Tóth (42). In section 2.3, we discuss three strategies which easily carry over to an adaptive grid. Until now, VAC had no automated means for dynamic grid adaptation. Our new AMRVAC software package combines the versatility of VAC with the advan-

tages of adaptive mesh refinement. In this first implementation, only Cartesian grids and explicit time stepping is allowed.

We demonstrate the flexibility of AMRVAC by using three equation modules in 1, 2 and 3 dimensions with different conservative high resolution spatial discretization schemes. In the example simulations from Section 3, the equation modules are subsets of the system of magnetohydrodynamics, given by

$$\frac{\partial \rho}{\partial t} + \nabla \cdot (\mathbf{v} \,\rho) = S_{\rho} \tag{1}$$

$$\frac{\partial \rho \mathbf{v}}{\partial t} + \nabla \cdot (\mathbf{v} \,\rho \mathbf{v} - \mathbf{B} \,\mathbf{B}) + \nabla p_{\text{total}} = \mathbf{S}_{\rho v}$$
 (2)

$$\frac{\partial e}{\partial t} + \nabla \cdot (\mathbf{v} \, e + \mathbf{v} \, p_{\text{total}} - \mathbf{B} \, \mathbf{B} \cdot \mathbf{v}) = S_e \tag{3}$$

$$\frac{\partial \mathbf{B}}{\partial t} + \nabla \cdot (\mathbf{v} \,\mathbf{B} - \mathbf{B} \,\mathbf{v}) = \mathbf{S}_B. \tag{4}$$

These are expressed in the conservative variables density ρ , momentum density $\rho \mathbf{v}$, total energy density e, and magnetic field \mathbf{B} , with all additional sources and sinks collected in the source terms S_{ρ} , $\mathbf{S}_{\rho v}$, S_{e} and \mathbf{S}_{B} . In these equations, the total kinetic and magnetic pressure is written as

$$p_{\text{total}} = (\gamma - 1) \left(e - \frac{1}{2} \rho \mathbf{v}^2 - \frac{1}{2} \mathbf{B}^2 \right) + \frac{1}{2} \mathbf{B}^2,$$
 (5)

for an ideal gas with adiabatic index γ . We stress that for all simulations done, we time advance the relevant subset in the selected dimensionality only, which is chosen *prior* to compilation of the code. This is possible because the entire AMR algorithm is implemented in the LASY-syntax (43), making it effectively dimension-independent. Similar to VAC, the AMRVAC package is configured to dimensionality using the VAC preprocessor (a Perl script), prior to compilation.

The time step is calculated to comply with the Courant-Friedrichs-Levy (CFL) stability criterion. If not stated otherwise, the Courant number is 0.9. The conservative high resolution spatial discretization schemes used are FCT, TVDLF, TVD, and TVDMU. These shock-capturing schemes differ in the numerical representation of the fluxes \mathbf{F}_i in the set of equations written as

$$\partial_t \mathbf{U} + \sum_i \partial_i \mathbf{F}_i(\mathbf{U}) = \mathbf{S}(\mathbf{U}, \partial_i \mathbf{U}, \partial_i \partial_j \mathbf{U}, \mathbf{x}, t). \tag{6}$$

Here, the conservative variables are collected in $\mathbf{U}(\mathbf{x},t)$, and i and j run over 1, 2 or 3 components of the spatial coordinate \mathbf{x} . If not stated otherwise, multidimensional scenarios add fluxes from the different orthogonal directions

simultaneously, in contrast to a dimensionally split (41) approach. The source terms S can also be unsplit, i.e. added at the same time as the fluxes, or split, i.e. added before and after the update with fluxes.

2.2 Adaptive Mesh Refinement

Detailed descriptions of the AMR algorithm can be found in Berger and Colella (7), Bell et al. (4), Friedel et al. (17), Steiner et al. (40), and many others. We summarize the essential steps and its user controlled features. Thereby, we pay specific attention to details which differ from the original Berger and Colella (7) approach.

2.2.1 Time stepping and conservation

In essence, AMR represents an automated procedure to generate or destruct—both controlled by the ensuing dynamics—hierarchically nested grids with subsequently finer mesh spacings Δx_i (index i = 1, ..., D for a D-dimensional problem). Up to a predefined maximal grid level l_{max} , consecutive levels l are characterized by even refinement ratios r_l with $l \in [2, l_{\text{max}}]$, such that

$$r_l \equiv \frac{\Delta x_{i,l-1}}{\Delta x_{i,l}} \equiv \frac{\Delta t_{l-1}}{\Delta t_l}.$$
 (7)

To make the global conservation property computationally tractable, we stick to Cartesian grids and the 'proper nesting' criterion (7): (i) level l > 1 grid boundaries coincide with grid lines of l - 1 meshes; and (ii) we insist on the telescoping hierarchy of l - 2, l - 1 and level l grids, except at (non-periodic) physical boundaries.

Consequently, two – related – issues for an AMR simulation are (1) how one time-advances on such a sequence of nested and thus overlapping grids, while (2) keeping the solutions consistent and conservative through a suitable 'update' process. The approach taken is illustrated in Figure 1, showing a hypothetical sequence of three time steps in a case where $l_{\text{max}} = 4$. The schema is traversed from left to right, bottom to top, and with horizontal 'update' arrows preceding vertical time 'advance' steps. Each time a level l > 1 has caught up in time with the underlying l - 1 grids, all level l - 1 cells that are covered by the finer grids are replaced by conservative averages. With notation defined in Figure 2 for a 2D example where $r_l = 2$, this is achieved by simple averaging since e.g. the density in the underlying coarse cell must be $\rho_{ij}\Delta x_{1,l-1}\Delta x_{2,l-1} = \sum_m \sum_n \rho_{mn}\Delta x_{1,l}\Delta x_{2,l}$ so that $\rho_{ij} = \sum_m \sum_n \rho_{mn}/r_l^2$. Also, those l - 1 cells immediately adjacent to a level l grid boundary are modified

to comply with the interface fluxes used on the finer level. This step is crucial for restoring the global conservation property across the AMR grid hierarchy. Following the notation used in Figure 2, this involves a change in the temporal update for the density in coarse cell indexed by ij on level l-1 bordered on the right hand side by level l cells. While originally

$$\rho_{ij}^{(n+1)} = \rho_{ij}^{(n)} - \frac{\Delta t}{\Delta x_{i,l-1}} \left(F_{i+\frac{1}{2}j}^{(n+\frac{1}{2})} - F_{i-\frac{1}{2}j}^{(n+\frac{1}{2})} \right) - \frac{\Delta t}{\Delta x_{i,l-1}} \left(F_{ij+\frac{1}{2}}^{(n+\frac{1}{2})} - F_{ij-\frac{1}{2}}^{(n+\frac{1}{2})} \right),$$

the time centered numerical flux across the right cell edge $F_{i+\frac{1}{2}j}^{(n+\frac{1}{2})}$ needs to be replaced by

$$F_{i+\frac{1}{2}j}^{(n+\frac{1}{2})} \Rightarrow \frac{1}{r_l^2} \left[F_{m-\frac{1}{2}n}^{(n+\frac{1}{4})} + F_{m-\frac{1}{2}n+1}^{(n+\frac{1}{4})} + F_{m-\frac{1}{2}n+1}^{(n+\frac{3}{4})} + F_{m-\frac{1}{2}n+1}^{(n+\frac{3}{4})} \right].$$

This flux fix restores the conservation. For more complete discussions of these update and flux fixing steps, we refer to (7).

In our AMR implementation, each level solves the same set of equations but can exploit a different discretization scheme per level. This in contrast with the more general Adaptive Mesh and Algorithm Refinement (AMAR) approach introduced by Garcia et al. (18), who coupled a continuum description with a particle method on the finest level of the AMR hierarchy. AMAR is possible because the only communication between different level solutions is through (i) the update process and (ii) the filling of 'ghost' cells as used for imposing boundary conditions.

The time advancing is further complicated by the automated regridding operation, as indicated by the grey circles in Figure 1. This regrid action is controlled by a simple criterion: when at least k time steps are taken on a certain level, this level is evaluated for refinement (k=2 in the figure). This affects all higher level grids which may either suddenly appear (in Figure 1 after the first and halfway in the second time step), disappear (after the second time step), or simply get rearranged or be left unchanged (halfway in the third time step). Of course, all level 1 grids remain unaltered, while the maximal allowed level $l_{\rm max}$ is never evaluated for further refinement.

2.2.2 Refinement criteria and regridding

Conceptually, regridding consists of two steps: in a first step, cells on level l which need to be refined are identified - 'flagged' - through physically and numerically controlled criteria. The second step ensures that a properly nested new grid level l+1 is formed which encompasses most, preferably all, of the

flagged cells. As pointed out above, the entire grid tree above a certain fixed base level l_{base} may change in that process. With the current finest level in existence being l_{fine} , one descends in levels from $l = \min(l_{\text{fine}}, l_{\text{max}} - 1)$ down to $l = l_{\text{base}}$.

In the first step – the flagging procedure – a great variety of refinement criteria can be adopted. Powell et al. (37) use local measures of compressibility, rotationality, and current density in their 3D MHD simulations and a region is refined when chosen treshold values are exceeded. Ziegler (53) uses similarly motivated, though more abstract, 'gradient value based' criteria where the slopes of functionals of the fluid variables, like the sound speed c_s , control refinement. In cases where an (approximate) Riemann solver based method is exploited, one may flag cells in a 'wave-affected region'. The extent of this region can be computed from the wave strengths and speeds as calculated in the Riemann solver. Such criterion was successfully exploited for unsteady Euler flows in (14).

Our refinement criterion to 'flag' cells is a variant of the error estimation procedure from Berger and Colella (7), which is generally applicable to any system of equations and integration method used. The idea is to rely on Richardson extrapolation, by comparing a solution for a level l obtained by coarsening and integrating with one which is integrated first, and subsequently coarsened. Bell et al. (4) improved on this basic idea by additionally exploiting the uncoarsened solution vector as well, to avoid missing features that get lost in the averaging. Moreover, they allowed for a certain amount of user-forced refinement or derefinement. In our implementation, the integration method in the error estimation can be selected as any one-step, dimensionally unsplit method, in combination with source terms. The criterion to flag cells can also exploit uncoarsened, high-resolution data, since we keep two full solution vectors \mathbf{U}_l^{n-1} and \mathbf{U}_l^n on each level $l < l_{\text{max}}$, separated in time by the corresponding Δt_l^{n-1} . For a fair variety of test problems (including the examples described below and those in Nool and Keppens (33), Keppens et al. (26; 27)), we found it sufficient to employ a first order version of the TVDLF method in the error estimation procedure, while integrating with a level-dependent mixture of TVD, TVDLF, or TVDMU on the full AMR hierarchy. The most crucial matter is to select the right combination of physics variables and mutual weighting factors to flag for. This is done as soon as a critical tolerance level $\epsilon_{\rm tol}$ is exceeded. In the examples from section 3, we always list the chosen variables and tolerances.

Given any procedure to identify cells which need refining, the AMR scheme must ultimately arrange them into properly nested 'rectangular' grids. Note that 'rectangle' is to be interpreted according to dimensionality. The basic steps to create this grid hierarchy were laid out in Berger and Colella (7). In the original algorithm, each flagged cell was surrounded with a buffer zone of width

 n_{buff} cells in each direction. The collection of flagged cells was subsequently made compatible with the nesting criterion. Some of the subtleties involved with ensuring this proper nesting are illustrated in Figure 3. Eventually, the collection of properly nested level l cells from which to form the level l+1grids need to be processed into 'rectangles'. This latter step was based on a succession of bisections of the encompassing 'rectangle', followed by a merge step. Bisections were repeated till all grids had their ratio of flagged to total points within the rectangle greater than a predetermined efficiency $E \in [0, 1]$. The merging was meant to balance the possible creation of many small grids. In its original form, this clustering algorithm created acceptable, but nonoptimal new candidate grids which would often overlap and could end up with efficiencies below E, due to the merging process. Berger and Rigoutsos (9) designed a more sophisticated algorithm to overcome these problems, where the grid partitioning was done along edges where transitions from flagged to unflagged regions occured. The edges were detected using pattern recognition techniques exploiting signature arrays. This clustering algorithm has been used on realistic 3D problems by Bell et al. (4), Balsara (2), and various others. We chose to modify the original merge step from (7) to avoid the creation of overlapping grids at the same nesting level. At the same time, we introduced an enforced minimal efficiency $E_{\min} \in [0,1]$ on all resulting new grids. It could be of interest to study how our modifications of the clustering algorithm compare with the Berger and Rigoutsos (9) approach on a variety of 2D and 3D synthetic input clusters. In all simulations reported in what follows, we took $E_{\min} = 0.5$.

The resulting new grid structure is then initialized from the solutions available on levels l_{base} and up. For those cells which can not be copied directly from the same level on the original grid hierarchy, we use a conservative, minmod limited linear interpolation from the coarser level underneath. Due to the nesting property, this is always possible. In a notation similar to the one used in Figure 2, this limited linear interpolation in a 2D case would read

$$\rho_{mn} = \rho_{ij} + \frac{x_{1,m} - x_{1,i}}{\Delta x_{1,i}} \Delta_i \rho^{(j)} + \frac{x_{2,m} - x_{2,i}}{\Delta x_{2,i}} \Delta_j \rho^{(i)}$$

where $\Delta_i \rho^{(j)}$ denotes the minmod limited slope in direction *i* involving leftsided, right-sided, and centered slopes determined from $[\rho_{i-1j}, \rho_{ij}, \rho_{i+1j}]$. Note that this is a conservative operation and second order accurate in smoothly varying density regions. Multidimensional MHD computations that make use of AMR have only been emerging over the last few years. Such computations must handle the additional complication of maintaining a solenoidal magnetic field. Two-dimensional adaptive calculations for incompressible MHD flows were presented by Friedel et al. (17). Those authors borrowed projection method techniques from incompressible Navier-Stokes simulations (5). Details of these projection methods and how to handle them on adaptive grid hierarchies have been discussed in Almgren et al. (1).

Steiner et al. (40) performed compressible MHD calculations using Flux Corrected Transport (11) in two space dimensions. They combined AMR with a staggered representation where the magnetic field components are defined at cell interfaces while all other conserved quantities are cell-centered. Balsara (2) has presented and applied a divergence-free AMR MHD scheme where the same staggering was used. The key elements in his scheme are a divergence free prolongation strategy to transfer coarse solution information to finer grids, and a divergence free restriction step to update coarse grids from overlying finer levels. Similar to the flux fix-up step inherent in the Berger and Colella (7) method to ensure the global conservation property, a similar electric field correction step is needed at mesh level interfaces to keep the overall solutions consistent.

Powell et al. (37) presented full 3D ideal MHD scenarios where solutionadaptive refinement is combined with an eight-wave Roe-type approximate Riemann solver method. Representative calculations of the magnetized solar wind impacting on a planetary magnetosphere have been reported by those authors. They advocated the use of an eight-wave formulation, which maintains the $\nabla \cdot \mathbf{B} = 0$ constraint to truncation error. The approach modifies the approximate Riemann solver to 'propagate' $\nabla \cdot \mathbf{B}$ errors with the plasma velocity and adds non-conservative source terms to the MHD equations proportional to $\nabla \cdot \mathbf{B}$. The latter was found to work well for non-Riemann solver based methods as well (46), and is trivial to carry over to an adaptive grid. Recently, Janhunen (20) and Dellar (15) formulated various arguments to only add the $\nabla \cdot \mathbf{B}$ related source term to the induction equation, which restores momentum and energy conservation. Yet another approach has been forwarded by Marder (32), which relies on diffusing the numerically generated divergence at its maximal rate. Dedner et al. (16) revisited this 'parabolic' approach and introduced various variants for hyperbolic divergence cleaning. In summary, three 'source term' approaches to handle the $\nabla \cdot \mathbf{B}$ constraint have respective

sources $(S_{\rho}, \mathbf{S}_{\rho v}, S_{e}, \mathbf{S}_{B})$ given by

$$\begin{pmatrix} 0 \\ -(\nabla \cdot \mathbf{B}) \mathbf{B} \\ -(\nabla \cdot \mathbf{B}) \mathbf{B} \cdot \mathbf{v} \\ -(\nabla \cdot \mathbf{B}) \mathbf{v} \end{pmatrix}_{\text{pow}}, \begin{pmatrix} 0 \\ 0 \\ 0 \\ -(\nabla \cdot \mathbf{B}) \mathbf{v} \end{pmatrix}_{\text{jan}}, \begin{pmatrix} 0 \\ 0 \\ \mathbf{B} \cdot C_{\text{d}} \Delta x^{2} \nabla (\nabla \cdot \mathbf{B}) \\ C_{\text{d}} \Delta x^{2} \nabla (\nabla \cdot \mathbf{B}) \end{pmatrix}_{\text{dif}}. (8)$$

In this paper, we apply these three simple approaches described by (8) in our cell-centered magnetic field representation. We cross-validate them on several 2D MHD problems using AMR. The coefficient $C_{\rm d}=0.2$ in all cases reported. This diffusive approach could also be altered to keep the energy equation fully conservative.

3 Results and efficiency evaluation

In this section, we present simulations covering 1D to 3D evolutions. Our findings are discussed for each problem in what follows. Refinement is based on the density, if not stated otherwise. Tables 3-4 collect some relevant statistics, including execution times. All 1D and 2D problems report timings done on a single 270 MHz MIPS R12000 CPU of an SGI Origin 200. The 3D cases have been obtained on single or multiple processors of the SGI Origin 3800, with 500 Mhz MIPS R14000 CPUs. The efficiency of an AMR calculation is defined as the ratio between its execution time and the time needed to run the same problem on a static grid at the resolution of the highest level allowed l_{max} . We also report on the percentage of computing time spent on AMR-specific overhead. This overhead includes the total time spent on the combination of regridding operations (including the error estimation procedure and the steps needed for creating and initializing properly nested new grid hierarchies) and update operations, i.e. the replacement of underlying coarse cell values and the flux fixing of coarse cells neighbouring a finer grid level. This overhead measure leaves out two factors which are arguably AMR related, namely the time spent on storing fluxes across grid boundaries – which is negligible – and the time spent on enforcing boundary conditions by means of ghost cells. On AMR grid hierarchies, the latter involves spatio-temporal interpolation procedures from underlying coarser levels for filling internal boundary regions surrounding level l > 1 grids.

3.1 1D simulations

3.1.1 Hydrodynamic shock tube

Our first example considers a Riemann problem for the 1D Euler system from Harten (19). With a ratio of specific heats $\gamma = 1.4$, the left and right states are

$$(\rho, \rho v_x, e)_L = (0.445, 0.311, 8.928)$$
 for $x < 8$
 $(\rho, \rho v_x, e)_R = (0.5, 0, 1.4275)$ for $x \ge 8$

in the $x \in [0, 14]$ domain with open (zero gradient) boundaries on both sides. Using 140 cells at level 1 and allowing for 5 levels with refinement ratios $r_l = 2$, we strive to efficiently achieve the accuracy of a static grid simulation exploiting 2240 cells. In Figure 4, we show two solutions at t = 2: the top panel shows the density as obtained with FCT. In the bottom panel, the TVDMU scheme with MC/Woodward limiter was used on all levels. The extent of the level $l \geq 2$ grids at this time is indicated. We set the tolerance $\epsilon_{\text{tol}} = 0.005$ and regrid parameters k = 4, E = 0.6.

As soon as the resulting rarefaction wave, contact discontinuity and shock are well-separated, higher level grids are only activated around the contact and the shock discontinuities. Locally, the solutions compare favorably with static, high-resolution results exploiting the same method: an equal amount of grid points resolving the contact and shock are found. We overplotted the high resolution static grid result in the bottom panel of Figure 4. Note that the FCT simulation does well in representing the discontinuities at the cost of introducing spurious oscillations, while the TVDMU scheme yields oscillation-free results. This may well be specific for the ETBFCT variant used here and described in Tóth and Odstrčil (46). Similar conclusions were reached there comparing both methods for static grid simulations and in Steiner et al. (40) where AMR calculations used FCT.

The execution times reported in Table 3 are for simulations up to time t = 2, using a fixed time step $\Delta t_5 = 0.000625$. They demonstrate an obtained efficiency of 2.43 when using FCT, which improves to 2.95 when using TVDLF and up to 3.76 for the TVDMU scheme. This increase in AMR efficiency is reflected in a decrease of the AMR overhead from 25.7 % down to 19.8 %.

3.1.2 Magnetohydrodynamic Riemann problem

Our second test problem is the 1.5D MHD Riemann problem introduced by Brio and Wu (13). With fixed $B_x = 0.75$ and $\gamma = 2$, the constant states are

$$(\rho, \rho v_x, \rho v_y, e, B_y)_L = (1, 0, 0, 1.78125, 1)$$
 for $x < 0.5$
 $(\rho, \rho v_x, \rho v_y, e, B_y)_R = (0.125, 0, 0, 0.88125, -1)$ for $x \ge 0.5$

in the $x \in [0, 1]$ domain with open boundaries. We use 40 base cells and compare two means to achieve locally the resolution of a 1280 cell static grid. Using the TVDMU scheme and minmod limiting on all levels, we compare in Figure 5 the density at t = 0.1 from a calculation with $l_{\text{max}} = 6$ and $r_l = 2$ versus one where $l_{\text{max}} = 4$ and $[r_2, r_3, r_4] = [4, 4, 2]$. The tolerance was set to $\epsilon_{\text{tol}} = 0.001$ and regrid parameters k = 4, E = 0.8. Both solutions accurately capture the slow compound shock, the contact discontinuity, and the slow shock. Locally, the obtained solution essentially coincides with a high resolution static case, again overplotted in the bottom panel. The timings in Table 3 indicate only little differences in efficiency between the AMR simulations exploiting 4 and those exploiting 6 levels. This is true for the TVDLF and the TVDMU scheme. For the latter discretization, the AMR overhead drops to about 15 %.

3.2 2D simulations

3.2.1 Advection tests

Our first 2D examples consider pure advection problems. On a doubly-periodic domain $[0,1] \times [0,1]$, a prescribed velocity field $\mathbf{v} = [v_x, v_y] = [1,1]$ diagonally advects a two-dimensional density profile. We perform two tests, one for a discontinuity dominated profile, and one for a smoothly varying density pulse.

The discontinuous profile is a two-dimensional density field representing the VAC-logo, see Figure 6. Within the logo, the density is set to 2, while it is 0.5 outside. We use a 50×50 level 1 grid, and allow for three levels with $r_l = 2$. The AMR control parameters are taken k = 4, E = 0.8, and $\epsilon_{\text{tol}} = 0.01$.

Figure 6 shows the initial t=0 grid structure, along with two snapshots at times t=0.6 and t=2, time-advanced with TVDLF and a MC/Woodward slope limiter. To demonstrate the maintained accuracy, cuts along the x-direction at y=0.5 from times t=0,1,2 are plotted as well. At each integer time, one should regain the t=0 situation. The numerical diffusion immediately smears the discontinuities, but the resulting density field is in exact agreement with a static grid result using 200×200 cells. As shown in Table 3,

the AMR simulation is twice as fast as a static grid run, with only 10 % of the execution time spent on regridding operations.

For the timings reported using the FCT method, we used a dimensionally split approach. As pointed out by Bell $et\ al.$ (4), directional sweeping can be less favorable to the AMR calculations as additional boundary work is needed (extra top and bottom layers in the x-sweep must be taken along to start the y-sweep). The stencil of the FCT discretization for directionally unsplit multidimensional simulations requires filling the corner ghost cells, which we avoid with the split approach.

Since the exact solution for this advection problem is known, we illustrate quantitatively that the AMR method reaches the same level of accuracy as a corresponding high resolution simulation. Table 1 summarizes the relative numerical errors obtained for a variety of AMR runs of different effective resolution. In general, we define the relative error for a variable u with respect to a reference solution $u^{\rm ref}$ on a $N \times M$ grid as

$$\delta u = \frac{\sum_{i=1}^{N} \sum_{j=1}^{M} | u_{i,j} - u_{i,j}^{\text{ref}} |}{\sum_{i=1}^{N} \sum_{j=1}^{M} | u_{i,j}^{\text{ref}} |}.$$
 (9)

In this equation, the $u_{i,j}$ values are to be obtained by extrapolating the AMR solution to a uniform grid of size $N \times M$ corresponding to its (highest level) effective resolution. When more than one variable is advanced, the average $\bar{\delta}$ is taken over all primitive variables u.

Table 1 demonstrates that (1) the AMR runs achieve the same (first order) convergence behaviour on this discontinuity dominated problem as the static grid results; (2) the AMR runs are essentially identical to their high resolution static grid counterparts, with minute improvements noticable when refining more often (lowering k) and/or lowering the tolerance ϵ_{tol} . Note that it is advisable to take k consistent with the buffer zone width n_{buff} used in the regridding process, as pointed out by Berger and LeVeque (8). As could be expected, the AMR efficiency is better for more grid levels than the 3-level one reported in Table 3: an efficiency of 6 is reached for the 800 × 800 AMR run exploiting 5 refinement levels.

To demonstrate that we obtain second order accuracy on smooth solutions, we similarly advected a Gaussian bell profile, initially set to

$$\rho(x, y, t = 0) = 2 \exp\left(-100\left[(x - 0.5)^2 + (y - 0.5)^2\right]\right)$$

within a radius 0.2 from the center of the square domain and constant elsewhere. Again, we compare the obtained numerical solution after two full advection cycles at time t=2 with the exact solution. Table 2 shows that we get

second order convergence behaviour to the exact solution, on static and corresponding AMR runs. The AMR simulations refined every second timestep (k=2), while the tolerance was fixed at $\epsilon_{\text{tol}} = 0.001$. At this tolerance, the 5 level AMR result achieving an 800×800 effective resolution did not trigger level 5 grids on the entire bell profile. Note that we did obtain a solution with the specified accuracy 0.001, which shows the accuracy and efficiency of our error estimation procedure. Hence, this explains the apparent loss in convergence rate for this AMR run.

Taken together with the 1D test cases from above, these AMR convergence studies of 2D advection problems demonstrate some general trends. First, the effective resolution of the AMR grid is setting the overall accuracy of the obtained solution, as desired. Note that it is controlled by the combination of (1) the base grid resolution; (2) the number of refinement levels l_{max} ; and (3) the refinement ratio's in between consecutive levels r_l . AMR runs of similar effective resolution which differ in the exact combinations taken of these three controlling elements should behave identical (cfr. the test problem shown in section 3.1.2). These conclusions are valid provided that the refinement criterion successfully tracks all flow features. The latter is influenced by the choice of $\epsilon_{\rm tol}$ and k. Table 1 demonstrates that only minor differences ensue when varying these parameters within reasonable bounds. A final set of AMR parameters, consisting of the efficiencies E, E_{\min} , and n_{buff} plays a role in determining the (highly varying) number and the structure of the grids on a certain level. There seems to be little reason to alter their values dramatically from run to run.

3.2.2 Double Mach Reflection of a Shock

A well-known shock dominated hydrodynamical test is the double Mach reflection of a shock on a wedge, introduced by Woodward and Colella (51). On a domain $[0,4] \times [0,1]$, a planar Mach 10 shock with post- and pre-shock states given by

$$(\rho, \rho v_x, \rho v_y, e)_{\text{post}} = (8, 8 \times 8.25 \sin 60^{\circ}, -8 \times 8.25 \cos 60^{\circ}, 563.5)$$

 $(\rho, \rho v_x, \rho v_y, e)_{\text{pre}} = (1.4, 0, 0, 2.5).$

makes an angle of 60° with the wedge, which is represented as a reflective boundary located at $x \in [1/6, 4]$ and y = 0. The adiabatic index is $\gamma = 1.4$. A self-similar pattern develops as the shock reflects off the wedge. The boundary conditions are fixed to the post-shock state at the left edge, open at the right edge, a partly fixed, partly reflective bottom, and a time- and space-dependent boundary value prescription along the top. In the AMR simulation, we use TVDLF on a maximum of 4 levels, with an 80×20 base resolution and refinement ratios $r_l = 2$. Furthermore, we took k = 6, E = 0.8, and a tolerance of $\epsilon_{\text{tol}} = 0.01$ in the regrid process. Figure 7 shows the density structure at time t = 0.2. We had to use a Courant number of 0.4 for the TVDLF discretization, since the x = 1/6 bottom boundary transition was not coincident with a cell edge at this resolution. As a result, a TVDMU simulation developed a carbuncle-type error at the location of the leading Mach shock intersecting with the wall. A parametric study of the AMR efficiency in terms of both execution time and memory needs for a large variety of effective grid resolutions is reported in Nool and Keppens (33).

Our timings indicate a significant gain in execution times – by a factor of 11 – between the corresponding 640×160 static grid simulation and the adaptive one. This high efficiency reflects the fact that the AMR process nicely succeeds in tracing the localized fronts, so that the domain coverage by level l > 1 grids remains low. In particular, level 4 grids initially cover only 5 % of the entire computational domain, with a modest increase to 15 % at the end of the calculation at t = 0.2. When raising the base resolution to 160×40 and again allowing for 4 levels with otherwise identical parameter settings, we reach an efficiency of 19.8 with still only 8% computing time spent on regridding.

3.2.3 Hydrodynamic Rayleigh-Taylor Instability

While the previous example represents a typical usage of AMR in hydrodynamical shock governed evolutions, we now turn to a simulation where a transition to a fairly global, turbulent state takes place. In such cases, it is not a priori evident that we can benefit from the use of dynamically triggered mesh refinement.

We simulate a 2D Rayleigh-Taylor instability on a $[0, 1] \times [0, 1]$ domain, with gravity $\mathbf{g} = -\hat{e}_y$ pointing downwards. As in Keppens and Tóth (22), we start from a dense fluid with $\rho_{\text{dense}} = 1$ above the interface $y_{\text{int}} = 0.8 + 0.05 \sin 2\pi x$, resting on top of a light fluid with $\rho_{\text{light}} = 0.1$. Both fluids are at rest at t = 0 and $\gamma = 5/3$. The pressure field is set from a centered differenced hydrostatic balance $dp/dy = -\rho$, ensuring that the pressure about y_{int} equals unity. This configuration is inherently unstable, while linear stability analysis in incompressible hydrodynamics predicts the shortest wavelengths growing the fastest. The staircased representation of the interface y_{int} initiates such short wavelength perturbations, while its sinusoidal variation and the periodic side boundaries select a preferential longer wavelength variation. Top and bottom are impenetrable to the fluids.

In (22), the fixed 100×100 resolution and the (small) numerical diffusion associated with the TVDMU scheme suppressed the development of much

of the small-scale structure. Here, we repeat that simulation using AMR to locally achieve a resolution of 800×800 . As we anticipate a need for higher resolution spreading across the entire computational domain, we want to make the cost associated with all intermediate levels $l \in [2, l_{\text{max}} - 1]$ as low as possible, without sacrificing accuracy on the highest level l_{max} . This can be done by using a level-dependent discretization scheme, such as the cheap but robust TVDLF method on all levels $l \leq l_{\text{max}} - 1$, in combination with a Riemann-solver based method, here TVDMU, on the highest level $l_{\text{max}} = 5$ only. The base grid is 50×50 and all refine ratios are $r_l = 2$. Other AMR parameters are as in the Woodward and Colella simulation. The snapshot of the logarithm of the density at time t = 1.9 shown in Figure 8 clearly demonstrates that we reached a very high effective Reynolds number regime.

In Figure 9 (left panel) we plot the domain coverage for each grid level in the Rayleigh-Taylor simulation as a function of time. This figure confirms the visual impression of fine structure spreading across much of the computational volume, such that at time t=1.9, more than 50 % of the area is at the highest resolution allowed. Also, the level l=2 grids cover the full domain at this time, making the l=1 grid calculation completely obsolete. Clearly, since the physical process cascades to smaller scales, the strategy of minimizing computational cost on lower lying levels through the use of a cheaper discretization pays off. In Table 3, we give an estimate of the execution time needed for a static 800×800 calculation from performing 20 time steps at this resolution. As the CFL-limited time step is observed to decrease monotonically in the AMR run, this allows to deduce both a lower and upper bound, and we report a mean value in the table. The corresponding efficiency of the AMR calculation is about 8.3. Note also the low value of 3.9 % of adaptive grid related overhead.

3.2.4 Orszag-Tang Vortex System

The compressible evolution of the Orszag-Tang vortex system (34; 35) is a well-studied model problem where a shock-governed transition to an MHD turbulent state takes place. On a doubly periodic domain of size $[0, 2\pi] \times [0, 2\pi]$, velocity "convection cells" described by $\mathbf{v} = [-\sin y, \sin x]$ are out of phase with magnetic islands resulting from $\mathbf{B} = [-\sin y, \sin 2x]$. At time t = 0, the density and pressure take the uniform values $\rho = 25/9$ and $\rho = 5/3$ with $\gamma = 5/3$. We evolve this system till time t = 3.14, using four levels with a 50×50 resolution at level 1. With $r_l = 2$, we aim to find the 'reference' high resolution 400×400 solution shown in Fig. 16 from (42). As in that paper, our Figure 10 shows the temperature distribution, and the fine details agree closely. Other parameters used in the AMR simulation are k = 6, E = 0.7, and $\epsilon_{\text{tol}} = 0.025$.

To demonstrate the advantages of the mixed strategy where TVDMU is used on the higher levels only, we report in Table 3 timings for 3 AMR simulations with $l_{\text{max}} = 4$: one where we use TVDMU on all levels, one where we use it on levels l = 3 and l = 4, mixed with TVDLF on l = 1, 2, and the advocated choice where TVDMU is active on level l = 4 alone. The efficiency increases from 5.85, to 6.28, to 7.38, respectively. In the right panel of Figure 9, the coverage of the domain by the four grid levels is shown for the most efficient case. In agreement with the initially smooth evolution of the density (uniform at t = 0), higher level grids are only activated from about t = 0.5 onwards. This represents an immediate gain in the obtained efficiency. As in the Rayleigh-Taylor simulation from above, the level l = 2 grids cover up the entire domain after some time. At the end of the simulation, this is about to happen for level 3 as well: a restart with a higher base level resolution should then be performed.

This 2D MHD problem is also an excellent test for comparing the three different strategies (referred to as 'Powell', 'Janhunen' or 'diffusive' and defined by their respective sources in equation 8) for controlling $\nabla \cdot \mathbf{B}$ in AMR simulations. Since the exact solution to this problem is not available analytically, our best reference is to use the same high resolution solution used in Tóth (42) for quantitative comparisons. As described in that paper, using a dimensionally split one-step TVD scheme with MC limiting, different approaches for controlling $\nabla \cdot \mathbf{B}$ still differ slightly at resolutions of 400×400 . Static grid MHD simulations exploiting a projection scheme (12) were found to yield almost always the most accurate solutions. Repeating the Orszag-Tang simulation with 4 AMR levels all exploiting the dimensionally split TVD scheme (for a fair comparison), the obtained relative errors as defined above at time t=3.14 with respect to a reference projection scheme solution are

$$\bar{\delta}_{\text{pow}} = 0.0546 \ \bar{\delta}_{\text{jan}} = 0.0808 \ \bar{\delta}_{\text{dif}} = 0.0465.$$
 (10)

These values should be contrasted to remaining differences among two high resolution static grid simulations: between a constrained transport solution and a projection scheme approach, a relative error of $\bar{\delta} = 0.021$ could be measured. Again, by lowering the tolerance $\epsilon_{\rm tol}$ to 0.01, improvements up to $\bar{\delta}_{\rm dif} = 0.02768$ can be obtained. These values for the relative errors are also consistent with the values reported in Table 7 of Tóth (42). Given the fact that for a considerable amount of time (for times t < 1), the AMR simulations could suffice with lower effective resolutions, the obtained relative errors are truly optimal.

3.2.5 Rotor

A second ideal MHD problem frequently used for comparing static grid MHD simulations is the rotor problem introduced by Balsara and Spicer (3). We perform the 'first' rotor test as mentioned in Tóth (42) where the setup is as follows. On the domain $[0,1] \times [0,1]$, thermal pressure at t=0 is p=1, and a uniform magnetic field has $B_x=5/\sqrt{4\pi}$ and $B_y=0$. We set $\gamma=1.4$, and create a dense disk within $r\equiv [(x-0.5)^2+(y-0.5)^2]^{1/2} < r_0=0.1$ with $\rho=10$ and $v_x=-2(y-0.5)/r_0$, $v_y=2(x-0.5)/r_0$. For $r>r_1=0.115$ the fluid is at rest with $\rho=1$. In between $r_0< r< r_1$, linear profiles for density and angular speed connect both states. We run the simulation till t=0.15. We strive for a 400×400 resolution, and use AMR exploiting 4 levels with k=6, E=0.7 and $\epsilon_{\rm tol}=0.025$. Flagging is done on density and both magnetic field components with weighting factors 0.8, 0.1 and 0.1, respectively. A snapshot of the thermal pressure is shown in Figure 10.

Since all important features are captured on the highest grid level at all times, adaptive simulations with 'source term' treatments to keep the magnetic field solenoidal yield effectively similar solutions as static runs. We quantify this latter statement as follows. We perform several AMR runs with a base grid of 50×50 and a varying number of grid levels $l_{\text{max}} = 2, 3, 4$. In each case, we use TVDLF on the lowest level(s), with TVDMU and MC limiting on the highest level. The latter scheme was used for the reference 400^2 solution of Figure 18 from (42). This static grid solution again used a projection scheme for $\nabla \cdot \mathbf{B}$. As in the previous test case, even static grid results yield mutual differences of order $\bar{\delta} = 0.013$ for a flux constrained transport versus a projection algorithm at this high resolution. When we then calculate the relative errors for AMR runs with effective resolutions increasing from 100^2 to 400^2 , these turn out to be

$$l_{\text{max}} = 2 \quad \bar{\delta}_{\text{pow}} = 0.0760 \quad \bar{\delta}_{\text{jan}} = 0.0794 \quad \bar{\delta}_{\text{dif}} = 0.0753.$$

$$l_{\text{max}} = 3 \quad \bar{\delta}_{\text{pow}} = 0.0358 \quad \bar{\delta}_{\text{jan}} = 0.0412 \quad \bar{\delta}_{\text{dif}} = 0.0373.$$

$$l_{\text{max}} = 4 \quad \bar{\delta}_{\text{pow}} = 0.0157 \quad \bar{\delta}_{\text{jan}} = 0.0248 \quad \bar{\delta}_{\text{dif}} = 0.0107.$$
(11)

These experiments demonstrate that each source term approach achieves first order convergence behaviour, as expected for this discontinuity dominated problem. Furthermore, the errors at effective resolutions of 400^2 are comparable to remaining differences between static grid solutions. As a final note, the advocated mixed discretization approach does not significantly sacrifice accuracy by the use of the more diffusive scheme on lower levels. Indeed, when we repeat the simulation with $l_{\rm max}=4$ and exploit TVDMU on all levels, the relative error for a diffusive approach becomes $\bar{\delta}_{\rm dif}=0.0096$. From this and the previous test case, it seems that the diffusive approach is preferable over

both other source term treatments.

3.2.6 MHD Kelvin-Helmholtz Instability

Since the Orszag-Tang and the rotor system represented ideal MHD evolutions, we include a resistive MHD simulation taken from Keppens *et al.* (25). The initial condition has a uniform density $\rho = 1$ and pressure p = 1, a sheared and perturbed velocity field given by

$$\mathbf{v} = [0.645 \tanh 20y, 0.01 \sin(2\pi x) \exp(-25y^2)]$$

and a discontinuously varying magnetic field $\mathbf{B} = \pm 0.129 \hat{e}_x$. The magnetic field is thus of uniform strength, but changes sign at y=0 in the domain $[0,1] \times [-1,1]$. With a resistivity $\eta = 10^{-5}$, high resolution simulations on static 400×800 grids demonstrated in (25) how the nonlinear evolution is characterized by Kelvin-Helmholtz roll-up triggering tearing behavior. As the Kelvin-Helmholtz instability warps and amplifies the initial current sheet at y = 0, several magnetic islands develop due to local reconnection events, which are clearly visible in the density pattern (see Fig. 9 in reference (25)). Shown in Figure 11 is a snapshot of the density pattern at t = 4.4 as obtained with an AMR simulation using 4 levels. Since it is vital to capture the initial dynamics at the y=0 interface accurately, we base the refinement criterion on the horizontal field component B_x so that all levels are activated at t=0. Other than that, parameters are k = 6, E = 0.8, $\epsilon_{tol} = 0.01$. Instead of the one-step TVD method used in (25), here we use the combined TVDLF (on levels l < 3) and TVDMU (on l = 4) approach. The timing reported in Table 3 is for a simulation up to t=5, at which time the transition to a turbulent state has occurred. Note the very high efficiency of 13.9 and the marginal regridding overhead. We used the Powell source terms in these calculations.

3.3 3D Simulations

3.3.1 3D Advection

We made an efficiency assessment of a 3D advection problem by diagonally advecting a sphere of radius 0.2 in the unit cube where the density was set to 2, while it was 0.5 external to the sphere. With triple periodic boundaries, the velocity $\mathbf{v} = (1, 1, 1)$ brings the sphere back to the center at t = 1. Similar to the 2D advection problem evaluated in Table 1, accuracy and convergence properties are equivalent to corresponding high resolution simulations. As reported in Table 4, a case where AMR exploits 3 levels, with a 40^3 base grid and refine ratios 2 and 4, demonstrates an efficiency of 19.9. The AMR overhead

has dropped to a rather negligible 4.14~%. This problem was run on a single CPU of the SGI Origin 3800.

3.3.2 3D MHD problems

As a first 3D MHD simulation, we recovered the evolution of a magnetized Rayleigh-Taylor instability as in Keppens and Tóth (22). 3D AMR MHD simulations recovering previous high resolution Kelvin-Helmholtz unstable jet evolutions are documented in Nool and Keppens (33). In those simulations, the diffusive source terms for controlling $\nabla \cdot \mathbf{B}$ were exploited. The problem setup for the Rayleigh-Taylor study straightforwardly augments the 2D HD simulation in section 3.2.3 with a uniform horizontal magnetic field $\mathbf{B} = 0.1\hat{e}_x$ and a 3D displacement of the interface separating the heavy from the light fluid according to $y_{\text{int}} = 0.8 + 0.05 \sin 2\pi x \sin 2\pi z$. The magnetic field suppresses the formation of much of the fine scale structure. Here, the magnetic field constraint is handled by the Powell source method.

Using AMR settings with k = 6, E = 0.7, $\epsilon_{\text{tol}} = 0.05$, the timings for two AMR simulations with $l_{\text{max}} = 2$ and $l_{\text{max}} = 3$ recovering a $80 \times 160 \times 80$ static grid case are reported in Table 4. In both simulations, TVDMU is used on the highest level only, and a roughly fivefold efficiency is reached. Since the base grid is only $20 \times 40 \times 20$, the problem setup immediately triggers level 2 grids on 20 % and 3 grids on 10 % of the total domain. These values increase to 50 and 20 %, respectively, at the end of the calculation. Using the average coverage values of 15 % and 35 %, and taking account of the relative refinement ratios, we can thus expect that the CPU time is at best reduced to 15 + 35/16 + 100/256 = 17.57 %, giving an optimal efficiency of 5.7, to be contrasted with the fivefold efficiency actually obtained. Since the runtime is dominated by the two highest levels, the obtained efficiency is truly optimal. The time spent on the regridding process is fully negligible: less than 2 % overhead was observed for this problem. Since one would typically use a higher base grid resolution, much better efficiencies can be expected to hold in practical 3D MHD simulations. Indeed, Ziegler (53) demonstrated an efficiency of 45 on a purely kinematic 3D magnetic 'reconnection' problem (where only the induction equation (4) was solved) starting from a base grid of $100 \times 120 \times 60$ while allowing for 3 grid levels.

To demonstrate improved efficiencies at higher base grid resolutions, we set up a 3D MHD implosion problem on the cube $[-0.5, 0.5]^3$ with base grid resolution 60^3 . We again only allow 3 grid levels, with $r_l = 2$, so that an effective resolution of 240^3 is realized. At time t = 0, the problem setup has a uniform magnetic field $\mathbf{B} = \sqrt{3/5}\hat{e}_y$ pervading a static medium where the pressure is 0.6 external to the sphere $x^2 + y^2 + z^2 = 0.2^2$ and 0.06 inside this sphere. The density is $\rho = 1$ except for a small off-center sphere within the

low pressure zone: where $(x+0.1)^2+y^2+z^2<0.04^2$ the density is increased tenfold. We run this simulation using triple periodic boundaries up to time t=0.4, corresponding to 125 level 1 discrete time steps Δt_1 . The solution at this time is shown in Figure 12, showing several quantities in cross-cuts at x=0, y=0, and z=0. We used TVDLF on all levels and the diffusive source term treatment for the solenoidal constraint. Other AMR parameters were k=2, E=0.8, $\epsilon_{\rm tol}=0.01$. This simulation was run on 16 processors of the SGI Origin 3800 and exploited OpenMP parallelization as explained in reference (27).

During the entire run, the level 3 grids covered on average 7.3 % of the entire domain, reaching a maximum of 10.2 %, while the level l=2 covered an average of 17.8 %, and a maximal coverage of 26.6 %. This already implies that this AMR run can at best achieve efficiencies in the range 8-11. For a fair comparison, we measured the wall clock time needed for performing one timestep at a fixed 240³ resolution, while again exploiting 16 processors and using AMRVAC as a domain decompositioner. This high resolution static grid run needed then 78.9 seconds per time step. The average wall clock time needed for setting one Δt_1 in the AMR calculation is 37.8 seconds. Taking into account the factor 4 in refinement from level 1 to level 3, the obtained efficiency is 8.3, exactly in its optimal range and indeed improved from the fivefold efficiency in the low base grid Rayleigh-Taylor problem. It is also worthwhile mentioning that the memory requirements for the 3-level AMR versus the static grid run are reduced by a factor of 6. Similarly, the space needed for storing one snapshot (in binary format where the three coordinates plus 8 unknowns per cell are saved in double precision) is a factor of 9 smaller in the AMR calculation. This also improves the time needed for IO operations: saving one snapshot from the AMR run on average took about 8 seconds, while that took roughly 51 seconds in the high resolution run. Hence, the overall advantages of performing grid-adaptive 3D MHD simulations are found in optimally reduced CPU times, drastically reduced memory requirements, and significantly smaller data volumes for eventual visualization purposes.

4 Overview and outlook

A dimension-independent implementation of the AMR algorithm, in combination with the modular structure of the Versatile Advection Code, has allowed us to assess the gain in computing time for a variety of 1D, 2D and 3D hydrodynamic and MHD problems. The overhead associated with the automated regridding process is fully negligible for realistic 2D and 3D problems. A tenfold efficiency is typical for the 2D cases shown, while the 3D timings indicated that very high efficiencies are indeed achievable: all 3D test problems demonstrated that the optimal efficiency reachable was in fact obtained. This will be

exploited in future AMR studies of correspondingly high resolution 3D MHD simulations of different, interacting fundamental plasma instabilities and stellar wind studies. It will be computationally beneficial to use Riemann solver based methods like TVDMU on the highest level only, in combination with the robust, more diffusive, but sufficiently accurate TVDLF discretization on lower levels.

The AMR algorithm introduces a fair amount of control parameters, which do affect run strategies and the obtainable efficiency. As a general rule, two important parameter sets can be distinguished: (1) those controlling the effective resolution (level l = 1 grid resolution, maximal allowed levels l_{max} , and consecutive refinement ratios r_l); and (2) those determining the refinement criterion (the choice of variable(s) to refine on, the number k of time steps taken before regridding, and the tolerance in the criterion $\epsilon_{\rm tol}$). The physics problem at hand usually suggests suitable variables to use in the latter. Refining every second timestep k=2 is a safe strategy, while the range for $\epsilon_{\rm tol}$ in all problems studied above was $\epsilon_{\text{tol}} \in [0.001, 0.05]$, where the largest values were used in the higher dimensional problems. As expected, the effective resolution plays a decisive role in the eventual efficiency. In cases where HD or MHD instabilities are simulated, their length scales will put constraints on the base grid resolution to use: it must be sufficiently high to allow for mode development. The localized nature of the problem will then help to determine what is a sufficient number of AMR levels l_{max} : this may require some trial and error while analyzing the first few time steps to find an optimal value. Generally, raising the number of levels will beneficially influence the efficiency.

For multi-dimensional MHD simulations exploiting AMR, three 'source term' approaches listed in equation (8) for controlling $\nabla \cdot \mathbf{B}$ to truncation error were investigated. Quantitative comparisons with reference high resolution simulations showed that all three work effectively for cell-centered \mathbf{B} representations. For the tests included here, the diffusive approach was found to yield the smallest relative errors. As previously demonstrated by Tóth (42) for static grid MHD simulations, we conclude that enforcing the solenoidal character of the magnetic field to machine precision in a particularly favoured discretization is not an absolute necessity for obtaining accurate multi-D MHD simulations exploiting AMR. The simple source term treatments offer a viable alternative to staggered field representations with considerable complications associated with constrained transport implementations on AMR grid hierarchies (2). Projection scheme approaches would work accurately and reliably for AMR MHD simulations as well, but can be expected to be more complicated algorithmically and computationally costly.

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Table 1 Convergence study for the 2D advection of the VAC logo. The resolution indicates the effective resolution for AMR runs, refining every k (sub-)time steps.

resolution	$l_{ m max}$	k	$\epsilon_{ m tol}$	$\delta_{ m static}$	δ_{exact}
100^{2}	1	_	_	0	0.12866
	2	4	0.01	0.000179	0.12867
	2	2	0.01	0.000215	0.12867
	2	2	0.005	0.000132	0.12866
200^{2}	1	_	_	0	0.07646
	3	4	0.01	0.000301	0.07656
	3	2	0.01	0.000172	0.07648
	3	2	0.005	0.000102	0.07648
400^{2}	1	_	_	0	0.04612
	4	4	0.01	0.001018	0.04676
	4	2	0.01	0.000222	0.04618
	4	2	0.005	0.000129	0.04617
800^{2}	1	_	_	0	0.02781
	5	2	0.01	0.000506	0.02811

Table 2 Convergence study for the 2D advection of the Gaussian bell profile. The resolution indicates the effective resolution for AMR runs ($l_{\rm max}>1$), refining every k=2 (sub-)time steps, at a tolerance $\epsilon_{\rm tol}=0.001$.

resolution	$l_{ m max}$	$\delta_{ m exact}$
100^{2}	1	0.0238102
	2	0.0238104
200^{2}	1	0.0058374
	3	0.0058379
400^{2}	1	0.001548
	4	0.001571
800^{2}	1	0.000422
	5	0.001271

Table 3 Overview of all 1D and 2D Simulations and Timings (in seconds).

problem	method	$l_{ m max}$	timing	eff.	$\mathrm{AMR}\%$
1D HD	FCT	1	18.35	_	_
Harten		5	7.55	2.43	25.7
$2240\times[0,2]$	TVDLF	1	24.12	_	_
		5	8.17	2.95	22.5
	TVDMU	1	35.39	_	_
		5	9.41	3.76	19.8
1.5D MHD	TVDLF	1	4.36	_	_
$Brio ext{-}Wu$		4	3.44	1.27	17.45
$1280\times[0,0.1]$		6	3.28	1.33	17.79
	TVDMU	1	7.38	_	_
		4	4.02	1.84	14.30
		6	3.83	1.93	15.65
2D Advection	FCT	1	47.91	_	
VAC- $logo$		3	33.70	1.42	7.9
$200\times200\times[0,2]$	TVDLF	1	107.36	_	_
		3	49.32	2.18	10.1
2D HD Shock reflection	TVDLF	1	4827.04	_	_
$640\times160\times[0,0.2]$		4	428.72	11.26	7.13
$1280\times320\times[0,0.2]$		1	33283.7	_	_
		4	1679.3	19.82	8.08
2D HD Rayleigh-Taylor	TVDMU	1	303715	_	
$800\times800\times[0,1.9]$	TVDLF-MU (4-1)	5	36504	8.32	3.91
2D MHD	TVDMU	1	8624.62	_	
$Orszag ext{-} Tang$	TVDMU	4	1474.58	5.85	3.43
$400\times400\times[0,3.1]$	TVDLF-MU (2-2)	4	1372.82	6.28	3.49
	TVDLF-MU (3-1)	4	1167.91	7.38	3.94
2D MHD Kelvin-Helmholtz	TVDMU	1	173572	_	_
$400\times800\times[0,5]$	TVDLF-MU (3-1)	4	12477	13.91	2.36

Table 4 Overview of 3D Simulations and Timings (in seconds). Note that the 3D Implosion problem mentions wall clock times for OpenMP parallelized code running on 16 processors of the SGI Origin 3800.

problem	method	$l_{ m max}$	$_{ m timing}$	eff.
3D Advection	TVDLF	1	172197	_
$320^3\times[0,1]$		3	8653	19.90
3D MHD	TVDMU	1	218650	_
$Rayleigh\hbox{-} Taylor$	TVDLF-MU $(1-1)$	2	46659	4.69
$80 \times 160 \times 80 \times [0,1]$	TVDLF-MU $(2-1)$	3	40726	5.37
3D MHD Implosion	TVDLF	1	39437	_
$240^3 \times [0, 0.4]$		3	4733	8.3

Fig. 1. A hypothetical sequence of three time steps in an AMR simulation allowing for 4 nested refinement levels. Vertical arrows indicate a temporal advance, horizontal arrows correspond to update operations ensuring conservation and consistency, while the automated regridding operation is called at the position of the grey circles. Regridding is invoked when at least k=2 time steps are taken.

Fig. 2. Illustration of the update and flux fix step. An underlying coarse (level l-1) cell value is merely replaced by a conservative average. A coarse cell bordering a finer level l grid needs its numerical flux across the edge $x_{i+\frac{1}{2}}$ replaced by the more accurate fluxes used on the level l.

Fig. 3. The regridding operation illustrated in 2D. Top left: 5 cells are flagged for refinement on 2 adjacent level l grids. The edge of the computational domain is indicated by thick solid lines. Each flagged cell is surrounded by a 2-cell buffer zone. Top right: Buffer cells not located on level l grids are discarded. A previously formed, enlarged level l+2 grid (small rectangle) is to be projected for proper nesting in the level l+1 grids being formed. Bottom left: discard cells which would violate the proper nesting criterion. Note the flagged cell which is lost! Bottom right: A possible grid structure for level l+1, whereby two extra level l cells (hashed squares) are taken along. Note that the level l+1 grids may overlap multiple level l grids.

Fig. 4. The density structure at time t=2 for the 1D hydrodynamic Harten shock tube. Using AMR with 5 grid levels, we confront a solution obtained with FCT (top panel) with a TVDMU result (bottom panel). The extent of the level l>1 grids is indicated by the solid bars. A high resolution static grid result is overplotted as a solid line in the bottom panel.

Fig. 5. The density variation at t=0.1 in the Brio-Wu shocktube problem. A solution using 6 refinement levels (top panel) is compared with one where 4 levels and correspondingly higher refinement ratios were used (bottom panel). The extent of the level l>1 grids is indicated by the solid bars and the bottom panel shows a high resolution reference result.

Fig. 6. Diagonal advection of the VAC-logo on a doubly periodic domain. We show the initial 3-level grid structure at top left, and snapshots at t = 0.6 (bottom left) and after two full advection cycles (bottom right). Cuts at y = 0.5 for times t = 0, 1, 2 (top right) demonstrate the maintained accuracy.

Fig. 7. The density structure at t=0.2 for the Woodward and Colella reflected shock problem, obtained with an AMR simulation using 4 grid levels and a 80×20 base grid.

Fig. 8. The logarithm of the density at time t = 1.9 shows the fine scale mixing process of a heavy fluid (dark) into a lighter one underneath under the influence of gravity. Only the level l = 5 grids are indicated.

Fig. 9. For the Rayleigh-Taylor simulation from Figure 8 (left) and for the Orszag-Tang vortex simulation from Figure 10 (right), the domain coverage of the grid levels used in the AMR simulation as a function of time.

Fig. 10. Left: the temperature at t = 3.14 for the Orszag-Tang magnetized vortex system as obtained with a 4 level AMR simulation. Right: the thermal pressure at t = 0.15 for the rotor problem as obtained with a 4 level AMR simulation.

Fig. 11. The density structure at time t=4.4 for a Kelvin-Helmholtz unstable shear flow configuration containing a current sheet shows the triggering of magnetic islands through tearing instabilities. Only the central part of the computational domain is shown.

Fig. 12. 3D MHD implosion problem at time t = 0.4. **Top left:** in the plane z = 0, we show a schlieren plot of the pressure, the magnetic field lines, and the velocity field. **Top right:** schlieren plot of the density in y = 0. **Bottom panels:** density (left) and thermal pressure (right) variation in the plane x = 0.