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Algorithm refinement for the stochastic Burgers' equation

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Abstract

In this paper, we develop an algorithm refinement (AR) scheme for an excluded random walk model whose mean field behavior is given by the viscous Burgers' equation. AR hybrids use the adaptive mesh refinement framework to model a system using a molecular algorithm where desired while allowing a computationally faster continuum representation to be used in the remainder of the domain. The focus in this paper is the role of fluctuations on the dynamics. In particular, we demonstrate that it is necessary to include a stochastic forcing term in Burgers' equation to accurately capture the correct behavior of the system. The conclusion we draw from this study is that the fidelity of multiscale methods that couple disparate algorithms depends on the consistent modeling of fluctuations in each algorithm and on a coupling, such as algorithm refinement, that preserves this consistency.

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1. Introduction

Algorithm refinement (AR) is an emerging paradigm in the modeling and simulation of multiscale problems. Mathematical models use distinctly different representations for microscopic and macroscopic scales with the corresponding algorithms echoing this disparity. Particle-based algorithms are a class of methods, typically used to model the microscopic scale, that represent the physical system by discrete, interacting entities. These "particles" represent anything from individual atoms to parcels of fluid to bacteria to automobiles. Field-based algorithms, typically used to model the macroscopic scale, are derived from models based primarily on partial differential equations with the physical system represented by continuum fields.

Algorithm refinement schemes (sometimes called "multi-algorithm hybrids") couple structurally different computational schemes such as particle-based molecular simulations with continuum partial differential equation (PDE) solvers.¹ The general idea is to perform detailed calculations using an accurate but expensive

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¹ Note that other types of AR hybrids exist (e.g., coupling spectral and discrete algorithms [33]).

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algorithm in a small region (or for a short time), and couple this computation to a simpler, less expensive method applied to the rest. The formulation of an AR scheme requires: projecting from the microscopic model to macroscopic; refining from macroscopic to microscopic; and handshaking between the two representations where they are coupled. A related issue is the establishment of "refinement criteria" that specify when a microscopic representation is needed and when a macroscopic representation is sufficient. Examples of algorithm refinement applied to fluid dynamics may be found in [15,17,29,34,35,38]; AR hybrids for interfacial propagation are discussed in [27,28,30].

One aspect of multiscale modeling that has received insufficient attention is the presence of spontaneous fluctuations at microscopic scales and their effect on the macroscopic scale. Accurate modeling of many phenomena require the correct representation of the variances and correlations of fluctuations, specifically when studying systems where the microscopic stochastics drive a macroscopic phenomenon. For physical systems, the correct treatment of fluctuations is especially important for stochastic, nonlinear systems, such as those undergoing phase transitions, nucleation, noise-driven instabilities and combustive ignition. In these and related applications, the nonlinearities can exponentially amplify the influence of the fluctuations.

Stochastic fluctuations in AR schemes have been investigated for two simple diffusive systems: linear diffusion [3,37] and the quasi-linear train model [5]. For those parabolic problems, one finds that when a particle algorithm is coupled to a deterministic continuum algorithm the variance of fluctuations is reduced in the particle regime near the interface. The variance of fluctuations within the continuum regime falls quickly away from the interface; however, variables, such as fluid velocity in the train model, that have long-range correlations retain these correlations of fluctuations (though at reduced magnitude) within the deterministic continuum region. Finally, stochastic continuum algorithms may be formulated such that when coupled to particle schemes they correctly duplicate the physical fluctuations throughout the computational domain.

Our longer term goal is to extend the development of AR methods with fluctuations to an adaptive mesh and algorithm method for the fluctuating compressible Navier–Stokes using a framework analogous to the non-fluctuating CNS solver discussed in [17]. As a prelude to that extension, in the present work we develop an AR method for Burgers' equation that couples nonlinear hyperbolic waves and diffusion. For the particle (microscopic) model we consider the asymmetric excluded random walk (AERW) [25,32]. The hydrodynamic (macroscopic) model for the AERW model is the viscous Burgers' equation with stochastic forcing [8,9,12,20].

In the following section, we describe in detail the AERW model. In the following section, we introduce the form of Burgers' equation that represents the hydrodynamic limit of the AERW model and describe a discretization of that equation based on a second-order Godunov scheme. In Section 4, we discuss the construction of the hybrid method that uses an overall adaptive mesh refinement framework to design the coupling between microscopic and macroscopic models. Section 5 contains computational examples that illustrate and validate the hybrid algorithm. As discussed in the concluding section, the numerical results demonstrate the importance and challenge of accurately modeling fluctuations to simulate and resolve both microscopic and macroscopic phenomena.

2. Asymmetric excluded random walk

2.1. Theory

The microscopic model for our system is an asymmetric excluded random walk. This model was selected since it has been studied extensively by the statistical mechanics community [25,32], shown to be equivalent, in the hydrodynamic limit, to the stochastic Burgers' equation and found to exhibit a variety of interesting phenomena (e.g., long-ranged correlations of non-equilibrium fluctuations). The AERW model is a system of N random walker particles on a two-dimensional rectangular lattice of dimensions $M_x \times M_y$. Each site is denoted by a coordinate pair (x_j, y_k) where $j = 1, ..., M_x$ and $k = 1, ..., M_y$.

Only one particle may occupy a site; the occupation number n(x, y) = 1 (or =0) if a site is occupied (or unoccupied). We choose the horizontal, or x-dimension of the lattice to correspond to the spatial domain of the PDE and define the corresponding density,

$$u_p(x_j) = \frac{1}{M_y} \sum_{k=1}^{M_y} n(x_j, y_k)$$
(1)

so $0 \le u_p(x) \le 1$. At equilibrium u_p is homogeneous and binomially distributed as the sum of M_y Bernoulli random variables, each with probability $U = N/M_x M_y$ of occupation. The mean and variance are

$$\langle u_p \rangle = U, \quad \langle \delta u_p^2 \rangle = \langle u_p^2 \rangle - \langle u_p \rangle^2 = \frac{U(1-U)}{M_y}.$$
 (2)

In the local equilibrium approximation the equal-time correlation of fluctuations is [32]

$$\langle \delta u_p(x_i) \delta u_p(x_j) \rangle = \frac{1}{M_y} \langle u_p(x_i) \rangle (1 - \langle u_p(x_i) \rangle) \delta_{i,j}.$$
(3)

At a non-equilibrium steady state the variance is more complicated due to long-ranged correlations of fluctuations [31].

Particles on the lattice move between adjacent sites according to the asymmetric exclusion process. Each particle waits a random time between moves with a mean free time of τ . The next particle to move is drawn at random by choosing a random site (x_j, y_k) ; if the site is occupied then its particle is selected otherwise another random site is chosen.

The selected particle may move up, down, left or right to an adjacent site, according to the probabilities assigned to the system. We take the particles to move horizontally or vertically with equal probability $(p_{\uparrow} = p_{\leftrightarrow} = 1/2)$ and the probabilities of moving up or down conditioned on vertical movement as equal $(p_{\uparrow} = p_{\downarrow} = 1/2)$. Asymmetry is introducing by taking unequal conditional probabilities for attempting to move left or right, that is, $p_{\leftarrow} \neq p_{\rightarrow}$ with $p_{\leftarrow} + p_{\rightarrow} = 1$. Once the particle and move direction are chosen, the particle moves to the destination site, if unoccupied; if the destination site is occupied then the particle remains in place. In either case the time is advanced and the entire process repeats.

2.2. Numerics

Given an initial density distribution u(x), the lattice is initialized by randomly filling sites. The dynamics is advanced by randomly choosing particles and move directions, as described above. In particle simulations the physical time may be advanced continuously (e.g., event-driven dynamics) or in time increments (e.g., molecular dynamics) and either approach may be used for the AERW. For the former, the time between moves is chosen as an exponential random variable with mean τ/N , where N is the number of particles. For the latter, the number of moves that occur during a time increment Δt_p is a Poisson distributed random value with mean $\mu = N\Delta t_p/\tau$; if $\Delta t_p \ll \tau/N$ then the probability of a move occurring during a particle time step Δt_p is $\mu + O(\mu^2)$.

The lattice is periodic in y so that particles attempting to move up from row M_y move to the bottom (first) row, provided it is unoccupied, with a similar definition for particles at the bottom row attempting to move downward. If the x-direction is also periodic, then its treatment is analogous to the treatment of periodicity in y.

The other type of boundary condition we consider is the imposition of Dirichlet conditions in x; in particular, fixing particle densities, $u_{\rm L}$ and $u_{\rm R}$ at the left and right boundaries, respectively. These boundary conditions represent the occupation probabilities for each site on the boundary. We view the system as being augmented with fictitious columns at j = 0 and $j = M_x + 1$ and with an effective total number of particles

$$N_e = N + u_{\rm L}M_y + u_{\rm R}M_y. \tag{4}$$

We then view the AERW as occurring on the enlarged lattice $((M_x + 2) \times M_y)$ with probabilistic "virtual" particles in the two boundary columns. We note that with Dirichlet boundary conditions, the number of particles in the system, N varies in time. Operationally these virtual particles enter the algorithm in two ways. First, suppose the selected particle location for the next move is in the left boundary column, say (x_0, y_k) ; with probability u_L that site is considered occupied by a virtual particle. If the adjacent site, (x_1, y_k) , is unoccupied then with probability $p_{\leftrightarrow}p_{\rightarrow}$ a virtual particle moves to that destination, becoming a real particle. Similarly, if a particle attempts to jump into the left boundary from an interior position, the destination is unoccupied with

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probability $1 - u_L$ in which case the jump is accepted and the particle removed. Analogous rules apply to the right boundary.

3. Burgers' equation and continuum method

3.1. Theory

The AERW model of the previous section is defined entirely in terms of a discrete lattice. In order to define a macroscopic model, we spatially embed the AERW model by assigning a spatial width Δx_p to the lattice sites in the x direction. With this definition the hydrodynamic limit of the asymmetric excluded random walk described in the previous section is the stochastic Burgers' equation [20]:

$$\frac{\partial}{\partial t}u(x,t) = -\frac{\partial}{\partial x}\{f(u) + d(u) + g(x,t)\},\tag{5}$$

where g is a stochastic flux and

$$f(u) = cu(1-u); \quad d(u) = -\epsilon \frac{\partial u}{\partial x}$$
(6)

are the nonlinear advective flux (sound speed, c) and the diffusion flux (diffusion constant, ϵ). With the change of variable u' = (1 - 2u)c this may be written in the more traditional form,

$$u_t' + u'u_x' = \epsilon u_{xx}' + 2cg_x \tag{7}$$

Note that variants of the stochastic Burgers' equation, with different types of stochastic forcing, are common in the literature (e.g. [8]). Also note that there are other particle models, such as the Boghosian–Levermore cellular automaton, that also converge to a stochastic Burgers' PDE in the hydrodynamic limit [11,24].

The wave speed and diffusion constant are determined from the AERW parameters as

$$\boldsymbol{c} = \frac{2p_{\rightarrow}\Delta x_p}{\tau} \left(p_{\rightarrow} - \frac{1}{2} \right) = \boldsymbol{c}_0 (2p_{\rightarrow} - 1) \tag{8}$$

and

$$\boldsymbol{\epsilon} = \frac{2p_{\leftrightarrow}\Delta x_{\rm p}^2}{\tau} p_{\rightarrow} p_{\leftarrow} = 2c_0 \Delta x_{\rm p} p_{\rightarrow} (1 - p_{\rightarrow}), \tag{9}$$

where $c_0 = p_{\leftrightarrow} \Delta x_p / \tau$ is the wave speed for the completely asymmetric walk ($p_{\rightarrow} = 0$ or 1). Since the wave speed f'(u) varies between +c and -c on the range of u, we define a dimensionless cell Reynolds number as

$$Re_{\rm c} = \frac{|c\Delta x_{\rm p}|}{\epsilon} = \frac{|p_{\rightarrow} - \frac{1}{2}|}{p_{\rightarrow}(1 - p_{\rightarrow})} \tag{10}$$

that characterizes the relative importance of diffusion and advection for the dynamics at a given mesh spacing. Note that for $p_{\rightarrow} = 1/2$ the random walk is symmetric (pure diffusion) and $Re_c = 0$; as p_{\rightarrow} approaches 0 or 1 the random walk is unidirectional (pure advection) and Re_c goes to infinity.

The stochastic flux is a Gaussian white noise with zero mean and correlation

$$\langle g(x,t)g(x',t')\rangle = A(x,t)\delta(x-x')\delta(t-t'),\tag{11}$$

where the brackets denote ensemble average. The noise amplitude, A(x, t), is related the correlation of density fluctuations; in the local equilibrium approximation,

$$\langle \delta u(x,t) \delta u(x',t) \rangle = \hat{u}(x,t)(1-\hat{u}(x,t))\delta(x-x'), \tag{12}$$

where \hat{u} is the solution to the deterministic Burgers' equation, that is,

$$\hat{u}_t = -f(\hat{u})_x + \epsilon \hat{u}_{xx}. \tag{13}$$

From the above one finds,

$$A(x,t) = 2\epsilon \hat{u}(1-\hat{u}). \tag{14}$$

The noise amplitude may also be obtained from the continuum limit of the master equation for the AERW [20].

3.2. Numerics

The stochastic Burgers' equation may be simulated numerically by a variety of CFD algorithms, with the choice guided by the application. For example, spectral methods have been developed for homogeneous, isotropic turbulence (see [21,26] and references therein). For our AR hybrid we choose a cell-centered finite difference method, specifically a second-order Godunov scheme to calculate the hyperbolic flux and a simple explicit predictor-corrector centered difference scheme to compute the diffusion term. See, for example, Colella [14] for a detailed discussion of second-order Godunov methods. We denote the spatial and temporal grid sizes as Δx and Δt and denote by u_j^n the average of u in cell-j at time n. We note that for the construction of a hybrid algorithm discussed later we will require that Δx be a multiple of Δx_p .

The second-order Godunov scheme constructs a linear profile within each cell with the slopes estimated by a higher-order finite difference approximation

$$u_{x,j} = \frac{-u_{j+2}^n + 8u_{j+1}^n - 8u_{j-1}^n + u_{j-2}^n}{12\Delta x}.$$
(15)

For advection-dominated problems a limiter is typically applied to these slopes; however, in the present context, we are resolving at the viscous length scale so no limiting is performed. These slopes are used to predict values at cell interfaces at the half-time level $t^{n+1/2}$. In particular, we define

$$u_{j+1/2,\ell}^{n+1/2} = u_j^n + \frac{1}{2} [\Delta x - \Delta t \max(f'(u_j^n), 0)] u_{x,j}$$
(16)

and

$$u_{j+1/2,r}^{n+1/2} = u_{j+1}^n - \frac{1}{2} [\Delta x + \Delta t \min(f'(u_j^n), 0)] u_{x,j},$$
(17)

where f' = df/du. We then define the hyperbolic flux $f_{j+1/2}^{n+1/2} = f(u_{j+1/2}^{n+1/2})$ where $u_{j+1/2}^{n+1/2}$ is the solution of the Riemann problem for $u_t + f_x = 0$ along the ray x/t = 0 with left and right states $u_{j+1/2,\ell}^{n+1/2}$ and $u_{j+1/2,r}^{n+1/2}$, respectively.

The diffusion and stochastic flux terms are evaluated using a predictor-corrector scheme, treating the hyperbolic flux terms as source terms. In particular, we first compute predicted values

$$u_{j}^{p} = u_{j}^{n} - \frac{\Delta t}{\Delta x} (f_{j+1/2}^{n+1/2} - f_{j-1/2}^{n+1/2}) + \frac{\Delta t}{(\Delta x)^{2}} \epsilon (u_{j-1}^{n} - 2u_{j}^{n} + u_{j+1}^{n}) + \frac{\sqrt{2\Delta t}}{\Delta x} (g_{j+1/2}^{n} - g_{j-1/2}^{n}).$$
(18)

We then compute corrected values

$$u_{j}^{n+1} = \frac{u_{j}^{p}}{2} + \frac{1}{2} \left(u_{j}^{p} - \frac{\Delta t}{\Delta x} (f_{j+1/2}^{n+1/2} - f_{j-1/2}^{n+1/2}) + \frac{\Delta t}{(\Delta x)^{2}} \epsilon (u_{j-1}^{p} - 2u_{j}^{p} + u_{j+1}^{p}) + \frac{\sqrt{2}\Delta t}{\Delta x} (g_{j+1/2}^{p} - g_{j-1/2}^{p}) \right).$$
(19)

This can be rewritten as

$$u_j^{n+1} = u_j^n - \frac{\Delta t}{\Delta x} \left(F_{j+1/2}^n - F_{j-1/2}^n \right), \tag{20}$$

where

$$F_{j+1/2}^{n} = f_{j+1/2}^{n+1/2} - \frac{1}{2} \left(\epsilon \frac{u_{j+1}^{n} - u_{j}^{n}}{\Delta x} + \epsilon \frac{u_{j+1}^{p} - u_{j}^{p}}{\Delta x} \right) - \frac{1}{\sqrt{2}} \left(g_{j+1/2}^{n} + g_{j+1/2}^{p} \right)$$
(21)

is the total flux.

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The stochastic flux for the asymmetric excluded random walk on an $M_x \times M_y$ lattice is discretized as

$$g_{j+1/2}^{n,p} = \sqrt{\frac{A_j^{n,p} + A_{j+1}^{n,p}}{2\Delta t M_y}} \Re.$$
(22)

where

$$A_j^{n,p} = 2\epsilon \tilde{u}_j^{n,p} (1 - \tilde{u}_j^{n,p}), \quad \tilde{u} = \min(\max(u, 0), 1)$$
(23)

and \Re is a Gaussian (normal) distributed random variable with zero mean and unit variance. Note that the instantaneous fluctuating values are used in place of the deterministic value (see Eq. (14)), which is accurate as long as the fluctuations remain small [18].

The scheme outlined above is stable provided it satisfies time step limits for both the hyperbolic and diffusive terms. In particular, we require

$$\frac{\Delta t \max |f'(u)|}{\Delta x} \leqslant 1 \quad \text{and} \quad \frac{\epsilon \Delta t}{\Delta x^2} \leqslant \frac{1}{2}.$$
(24)

We also note that it is possible to use a simpler version of the scheme based on an explicit first-order treatment of the diffusion term. For the most part, the simpler version provides reasonable predictions; however, at larger Δt the first-order scheme over-predicts the variation in the equilibrium solution by approximately 5%, suggesting that the temporal truncation error terms suppress the smoothing effect of the diffusion.

4. Algorithm refinement hybrid

In this section, we develop a hybrid algorithm refinement method that couples the AERW model introduced in Section 2 with the stochastic Burgers' equation algorithm in Section 3.

4.1. Basic construction

Philosophically, the construction of the hybrid is based on the notion that the particle description provides a more accurate representation of the solution than the stochastic PDE. Thus, the basic idea is to represent the dynamics with the continuum model except in a localized region where higher-fidelity particle representation is required.

Our perspective in designing the algorithm follows the adaptive mesh and algorithm (AMAR) approach introduced in [17]. In contrast to other AR approaches (see, e.g. [15]), the AMAR approach maintains a solution of the macroscopic model over the entire domain (see Fig. 1). An error estimation criterion is used to estimate where the improved-representation of the particle method is required. That region, which can change dynamically, is then "covered" with a particle patch. In this hierarchical representation, the solution is given by the particle solution on the region covered by the particle patches and the continuum solution on the remainder of the domain.

The coupling between the particle and continuum regions uses the analog of constructs used in developing hierarchical adaptive mesh refinement algorithms. For simplicity, we will assume that there is a single refined patch and that the mesh spacing for the continuum solver is equal to the lattice spacing Δx_p . Generalization of the approach to include multiple patches (e.g. [39]) and allowing the continuum mesh to be an integral multiple of Δx_p (e.g. [4]) is fairly straightforward.

Integration on the hierarchy is a three step process. First, we integrate the continuum algorithm from t^n to t^{n+1} , i.e., for a continuum step Δt . The old and new states, u_j^n and u_j^{n+1} , are retained until the particle time step is complete. Continuum data at the edge of the particle patch is interpolated in time to provide Dirichlet boundary conditions for the particle method.

We have considered both of the time-evolution schemes discussed in Section 2. For the equal time step version, we choose Δt_p so that $\Delta t_p = \Delta t/M_t$ for a specified integer M_t . We then advance the particle method by M_t steps until the particle and the continuum solver are at the same time. For the random time version of the algorithm, the particle method is advanced by moves, each with a random time increment, until the next transition would advance the particle time beyond t^{n+1} at which point the two solutions are synchronized. We note that at the synchronization juncture the particle and continuum solutions are not quite at the same time level.

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Fig. 1. Schematic of AERW/PDE hybrid. Advance: (A) advance continuum solution; (B) set boundary conditions for AERW from timeinterpolated PDE solution; (C) advance particle system by AERW. Synchronize: (D) replace overlaying continuum values with particle values; (E) reset PDE interface cells by refluxing.

For the most part, this has little effect on the computational results; however, it does lead to errors of approximately 1% in the mean solution at equilibrium, which are not observed with the temporally synchronized version.

4.2. Synchronization

The initial stage of the integration process essentially advances the macroscopic model separately with a one-way coupling to the microscopic model by way of the Dirichlet boundary conditions. The macroscopic model is not influenced by the microscopic model; the goal of the synchronization process is to correct the macroscopic solution to reflect the effect of the microscopic model as though the integration were tightly coupled.

There are two components to the synchronization process. First, on the region covered by the particle representation we replace the continuum solution obtained from the SPDE discretization by the more accurate particle representation, i.e, set

$$u_j^{n+1} = \frac{1}{M_y} \sum_{k=1}^{M_y} n(x_j, y_k)$$
(25)

for each cell covered by the particle patch. Second, the continuum cells immediately adjacent to the particle region, which supplied boundary data for the particle region during its advance, are corrected by "refluxing". Specifically, suppose the left boundary of the particle patch occurs in cell J + 1. The value in continuum cell J was updated with the continuum scheme

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$$u_J^{n+1} = u_J^n - \frac{\Delta t}{\Delta x} \left(F_{J+1/2}^n - F_{J-1/2}^n \right)$$
(26)

with the fluxes F^n computed from the continuum values (see Eq. (20)). The microscopically correct flux is given by the net number of particles moving across edge J + 1/2 rather than by the continuum flux $F^n_{J+1/2}$. To perform the refluxing correction we monitor the number of particles, $N^{\rightarrow}_{J+1/2}$ and $N^{\leftarrow}_{J+1/2}$, that move into and out of the particle region, respectively, across the continuum/particle interface at edge J + 1/2. We then correct the continuum solution as

$$u_{J}^{\prime n+1} = u_{J}^{n+1} + \frac{\Delta t}{\Delta x} F_{J+1/2}^{n} - \frac{N_{J+1/2}^{\rightarrow} - N_{J+1/2}^{\rightarrow}}{\Delta x M_{y}}.$$
(27)

This update effectively replaces the continuum flux component of the update to u_J^{n+1} on edge J + 1/2 by the flux of particles through the edge. The use of particle fluxes is seamless at the interface because the continuum flux is already stochastic (see Eqs. (21) and (22)). An analogous refluxing step occurs in the cell adjacent to the right-hand boundary of the particle region. Finally, note that this synchronization procedure guarantees exact conservation of integrated density. The technical details of refluxing in higher dimensions (e.g., the treatment of corners) are discussed in Garcia et al. [17].

4.3. Refinement criterion and regridding

The AMAR framework allows us to dynamically change the location of the particle region. There are several possible strategies for designing refinement criteria. For the examples described in Section 5.3, we will focus on criteria that identify cells where the solution has a large gradient characteristic of a viscous shock profile. A straightforward measurement of the local gradient of the solution (e.g., $(u_{j+1} - u_j)/\Delta x)$ is not adequate since the inherent fluctuations could trigger refinement even at equilibrium. What is needed is a robust measure that identifies viscous shocks without generating substantial "false positives" leading to unnecessary refinement. To this end we define a regional gradient using

$$D_{j} = \frac{1}{S\Delta x} \left[\frac{1}{S} \sum_{i=1}^{S} u_{j+i} - \frac{1}{S} \sum_{i=1}^{S} u_{j-(i-1)} \right],$$
(28)

where the stencil size S is specified; we take S = 4 in the computations in the following section. From Eq. (3) one may easily estimate the expected standard deviation σ of D resulting from equilibrium fluctuations and set a tolerance of $C\sigma$, where C is a constant. To estimate where to place the particle region in the adaptive code we compute the regional gradient at each point; if $|D_j|$ exceeds the tolerance level, then cells j and j + 1 are tagged. Since we restrict ourselves to a single particle patch, the largest interval containing all tagged cells is then the new particle region. If multiple patches were allowed then tagged cells would be collected to form particle regions; techniques for collecting tagged cells in an optimal manner are well established in the mesh refinement literature [7].

Once the new particle region has been identified, it must be initialized. For continuum cells that were already in the particle region, we simply retain the distribution of occupied sites. For cells that were not in the previous particle region, we use the continuum density to compute N_f , the desired number of particles for filling a column. The simple way to do this is to take $N_f = u_j^n M_y$ randomly rounded to the nearest integer; an alternative approach would be to fill randomly each site with probability u_j^n . We use the former approach since it preserves conservation of total density (to within quantization rounding). We note the regridding algorithm does not need to be done every step. Simple estimates based either on CFL considerations or estimates of discrete traveling wave velocities can be used to determine how often to regrid [6].

5. Computational examples

This section presents a series of computational examples, of progressively increasing sophistication, that demonstrate the accuracy and effectiveness of the algorithm refinement hybrid. We consider four numerical schemes: the asymmetric excluded random walk (AERW) from Section 2; the stochastic PDE (SPDE) for Bur-

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gers' equation from Section 3 and two algorithm refinement hybrids from Section 4. The first hybrid couples the AERW and SPDE, with the particle scheme in a single patch within the system. The second hybrid is similar but without a stochastic flux, that is, using a deterministic PDE (DPDE). Both fixed-patch and adaptive hybrids are considered as well as a handful of minor variants, discussed below.



Fig. 2. Mean $\langle u \rangle$, variance $\langle \delta u^2(x) \rangle$ and center point correlation $\langle \delta u(x) \delta u(0) \rangle$ versus x for a system at equilibrium ($u_L = u_R = 0.5$). Lines are: SPDE/AERW hybrid with $\Delta t = 0.05$ (red dashed dot); SPDE/AERW hybrid with $\Delta t = 0.1$ (solid red); SPDE/AERW hybrid with $\Delta t = 0.2$ (red dotted); AERW (green); SPDE with $\Delta t = 0.05$ (solid blue); predicted variance $\langle u \rangle (1 - \langle u \rangle) / M_y$ (dashed black). Note that for the correlation $\langle \delta u(x) \delta u(0) \rangle$ the large spike at x = 0 is omitted from the plot.

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The following parameter values are common to the simulations in all the examples: $M_y = 150$, $\Delta x = \Delta x_p = 0.01$, $\tau = 1$, $p_{\leftrightarrow} = 1/2$, $p_{\uparrow} = p_{\downarrow} = 1/2$, and $c_0 = 0.01$. The particle time step is chosen to be small ($\Delta t_p \approx \tau/(75N)$) so the probability of a move occurring during a time step is taken as $N\Delta t_p/\tau$. For a system length of L the viscous relaxation time is $T_{\epsilon} = L^2/\epsilon$; in our simulations $T_{\epsilon} = O(10^4)$. For the simulations of steady states the system is initialized near the final state and allowed to relax for a time that is long compared to the relaxation time (typically for >100 T_{ϵ}) before taking samples.

5.1. Equilibrium state

First, we consider the simplest scenario, a system at the equilibrium state with equal, fixed density at x = -0.5 and 0.5. The probability of moving to the right is $p_{\rightarrow} = 0.55$, corresponding to a cell Reynolds number of $Re_c = 0.20$ (weakly hyperbolic). The single-algorithm simulations (AERW and Burger's SPDE) have 100 sites or grid points in the x-direction. The AR hybrids introduce a fixed particle patch at the center of the system between x = -0.1 and x = 0.1 with $M_x = 20$. The hybrid simulation is performed at three continuum time step sizes: $\Delta t = 0.05$, 0.1, and 0.2. Since incremental time stepping is used in the particle algorithm, to keep $\Delta t_p = \Delta t/M_t \ll \Delta t$ we take three corresponding values: $M_t = 8000$, 14,000, and 28,000. Each simulation is run to a final time of $T = 2 \times 10^7$, which corresponds to $N_t = T/\Delta t$ continuum time steps (e.g., $N_t = 4 \times 10^8$ for the smallest Δt).

Typical results from the various numerical schemes are shown in Fig. 2 where the mean, $\langle u \rangle$; variance, $\langle \delta u^2 \rangle$; and correlation, $\langle \delta u \delta u' \rangle$ of density are plotted versus position. These three quantities are estimated from samples as



Fig. 3. Center point correlation versus x for a system at equilibrium ($u_L = u_R = \langle u \rangle = 0.5$) with the SPDE method using mean and instantaneous solution for the noise amplitude. Lines are SPDE: 1 using the mean (blue); SPDE: 2 using the instantaneous solution (magenta); AERW (green). Time step is $\Delta t = 0.05$.

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$$\langle u(x_i)\rangle = \frac{1}{N_s} \sum_{n=N_r+1}^{N_r} u_i^n;$$
⁽²⁹⁾

$$\langle \delta u(x_i)^2 \rangle = \langle \delta u(x_i) \delta u(x_i) \rangle; \tag{30}$$

$$\langle \delta u(x_i) \delta u(x_j) \rangle = \frac{1}{N_s} \sum_{n=N_r+1}^{N_r} u_i^n u_j^n - \langle u(x_i) \rangle \langle u(x_j) \rangle, \tag{31}$$

where $N_s = N_t - N_r$ is the number of samples and N_r is the number of continuum time steps that the system is allowed to relax before sampling begins. Typical values for the steady-state runs were $N_s = 10^9$ and $N_r = 10^5$.

Even at equilibrium the stochastic PDE scheme does not exactly match the AERW results, for example the SPDE method has an error of about 1.8% in the variance and a relative error in the correlation of $\langle \delta u \delta u' \rangle / \langle \delta u^2 \rangle \approx 1\%$. This discrepancy is expected since the numerical scheme uses the instantaneous solution to compute the amplitude of the noise instead of the mean (see Eqs. (14) and (23)). This effect is illustrated in Fig. 3, which compares simulations where the mean and instantaneous state are used to calculate the noise amplitude and verifies that the former is in agreement with the AERW results. Obviously the mean is known at equilibrium but for time-dependent, non-equilibrium problems the SPDE method needs to use instantaneous values of the state to evaluate the noise. In any case, Eq. (14) is only rigorously valid at equilibrium.

In general, the SPDE/AERW hybrid gives good results with a small discrepancy due to the effect just described regarding the noise amplitude. For the hybrid simulations, there is a small error in the mean of approximately 0.3% of the solution which decreases to about 0.2% when Δt is decreased to 0.05. This error is due to the buildup of the discrepancy between SPDE and AERW models at the boundaries of the particle patch in the hybrid. In the variance, we see an error of about 0.6% in the SPDE part of the domain, due to the same effect. In addition, there are spikes at the edges of the particle patch, representing an error of 3% at



Fig. 4. Variance $\langle \delta u^2(x) \rangle$ and center-point correlation $\langle \delta u(x) \delta u(0) \rangle$ versus x for a system at equilibrium ($u_L = u_R = 0.5$). Lines are SPDE/AERW hybrid (red); DPDE/AERW hybrid (blue) (compare with Fig. 2). For both cases $Re_c = 0.20$ ($p_{\rightarrow} = 0.55$) and $\Delta t = 0.05$.

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 $\Delta t = 0.2$ decreasing to about 0.3% at $\Delta t = 0.05$, evidently due to the temporal truncation error in the SPDE solver. These errors neither increase nor decrease with further sampling.

Fig. 4 illustrates the effect on fluctuations when the continuum PDE scheme does not include a stochastic flux. Clearly, the variance drops to near zero inside the DPDE regions, left and right of the particle patch, yet the variance within the patch remains nearly correct except near the interface. As discussed in Section 1, this general result was observed in previous studies of AR hybrids for parabolic systems [4,5] but it was not obvious that hyperbolic systems would be similar. Even more interesting is the appearance of a large correlation of



Fig. 5. Mean $\langle u \rangle$, variance $\langle \delta u^2(x) \rangle$ and center point correlation $\langle \delta u(x) \delta u(0) \rangle$ versus x for a rarefaction steady state ($u_L = 0.9$, $u_R = 0.1$). Lines are Hybrid AERW/SPDE (red); Hybrid AERW/DPDE (dashed dot red); AERW (green); SPDE (blue).

fluctuations in the particle region of the DPDE/AERW hybrid, an effect that will be discussed in the following subsection.

One final note. Using periodic boundary conditions gives similar results to those presented above for Dirichlet boundary conditions. Furthermore, using periodic boundary conditions we confirmed that the AR hybrids conserve total density, $\sum_i u_i$, exactly. (When the grids move dynamically this exact conservation is lost because of quantization effects in defining a particle distribution from the continuum data.)

5.2. Rarefaction steady state

Next, we consider non-equilibrium states by taking different densities at the left and right boundaries. For c > 0 the steady solutions to Burgers' equation are a shock wave if $u_L < u_R$ and a rarefaction wave if $u_L > u_R$. In this subsection we examine the latter, turning to shocks in the last two examples.

The parameters used for the rarefaction steady state are $u_L = 0.9$, $u_R = 0.1$, L = 1, $Re_c = 0.20$ ($p_{\rightarrow} = 0.55$), $T = 4 \times 10^7$, and $\Delta t = 0.05$. In the hybrid, $M_t = 8000$ and the particle patch is fixed between x = -0.1 and 0.1.

Fig. 5 shows typical results from the simulations. All methods give good agreement with the AERW method in the mean. As in the equilibrium case, the hybrid AERW/SPDE gives good agreement in the variance with both the AERW and SPDE solvers. However, when the AERW is coupled to the deterministic PDE solve, the variance falls to zero outside the particle patch while remaining close to the correct value inside the particle patch (similar to the equilibrium result of Fig. 4).

Fig. 5 shows that a long-range correlation, predicted by [13,19,31], is observed in the AERW, SPDE, and AERW/SPDE solvers and the three simulations are in good agreement with each other. This figure also shows that the AR hybrid using a deterministic PDE solver erroneously *enhances* this correlation; Fig. 2 shows a sim-



Fig. 6. Instantaneous density u(x, t) versus position x for a moving shock ($u_L = 0.1, u_R = 0.8$) for $Re_c = 0.20$ ($p_{\rightarrow} = 0.55$), which has shock speed $\sigma = 5 \times 10^{-5}$. Methods used are AERW/SPDE hybrid (red); DPDE (blue); vertical green lines delineate the AERW particle region of the hybrid.

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ilar effect appearing even at equilibrium, where no correlation is expected. It is not clear why this occurs since the correlation in a related AR hybrid of the "train" model is diminished [5]. One possible explanation is the induction of spurious correlations, even at equilibrium, when a reservoir does not generate the correct fluctuations spectrum [36].

5.3. Shock tracking

Unlike the rarefaction case, when $u_L < u_R$ the deterministic solution develops a shock wave in finite time propagating with speed $\sigma = c(1 - u_L - u_R)$. When viscous terms are added, the solution forms a smooth traveling wave moving at speed σ . Figs. 6–8 show examples of propagating shock waves for increasing cell Reynolds number. For these examples we have used the automatic criterion discussed in Section 4.3 to localize the particle region around the shock. Note that in each case the refinement criteria does a good job of localizing the particle region near the shock. For the most diffuse case, $p_{\rightarrow} = 0.55$, the solution looks essentially like the deterministic solution with superimposed noise. On the other hand, for the stronger shocks the fluctuations introduce sufficient perturbations to noticeably shift the shock location; this drift in the shock position is investigated further in the next subsection.



Fig. 7. Instantaneous density u(x, t) versus position x for a moving shock ($u_L = 0.1, u_R = 0.8$) for $Re_c = 0.95$ ($p_{\rightarrow} = 0.7$), which has shock speed $\sigma = 2 \times 10^{-4}$. Methods used are AERW/SPDE hybrid (red); DPDE (blue); vertical green lines delineate the AERW particle region of the hybrid.

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Fig. 8. Instantaneous density u(x, t) versus position x for a moving shock ($u_L = 0.1, u_R = 0.8$) for $Re_c = 1.87$ ($p_{\rightarrow} = 0.8$), which has shock speed $\sigma = 3 \times 10^{-4}$. Methods used are AERW/SPDE hybrid (red); DPDE (blue); vertical green lines delineate the AERW particle region of the hybrid.

5.4. Shock diffusion

Motivated by the observations in the previous section regarding the variation in the shock position, we consider the diffusion of the position of a stationary shock. Shock drift is well known in other particle simulations, such as in shock tube modeling by DSMC, which must correct for the drift when measuring profiles for steady shocks [10]. The general problem has been analyzed for the AERW and many results are known [2,1,16,22] but here we focus on the variance of the shock location as a function of time. We define a shock location, s(t) by fitting a Heaviside function of equal integrated density, that is,

$$\int_{-L/2}^{s(t)} u_{\rm L} \, \mathrm{d}x + \int_{s(t)}^{L/2} u_{\rm R} \, \mathrm{d}x = \int_{-L/2}^{L/2} u(x,t) \, \mathrm{d}x \tag{32}$$

from which we find

$$s(t) = L \frac{\bar{u}(t) - 1/2(u_{\rm L} + u_{\rm R})}{u_{\rm L} - u_{\rm R}},\tag{33}$$

where $\bar{u} = L^{-1} \int_{-L/2}^{L/2} u(x,t) dx$ is the instantaneous average density. The shock location fluctuates with a diffusion similar to that of a simple random walk [16] so averaging over ensembles from the same initial state gives

$$\langle \delta s^2 \rangle \approx 2\mathscr{D}t$$
 (34)

with a shock diffusion coefficient, \mathcal{D} , that depends on Reynolds number, shock strength, etc. Note that this expression for the variance is not accurate at very short times (due to relaxation transient from initial state) or at very long times (due to finite system size). Also note that the variance of the total mass, $\langle L^2 \delta \bar{u}^2 \rangle$, diffuses



Fig. 9. Variance of shock location $\langle \delta s(t)^2 \rangle$ versus time t for a deterministically steady shock ($u_L = 0.1, u_R = 0.9$). Methods used are AERW with $p_{\rightarrow} = 0.7$ (green); SPDE/AERW hybrid with $p_{\rightarrow} = 0.55$ (blue); SPDE/AERW hybrid with $p_{\rightarrow} = 0.7$ (red); SPDE/AERW hybrid with $p_{\rightarrow} = 0.8$ (black); DPDE/AERW hybrid with $p_{\rightarrow} = 0.7$ with 8 cells of additional buffering (magenta).

in the same fashion. This indicates that the diffusion of s is different than other shock profile variables (e.g., center-of-mass location) that fluctuate even if \bar{u} is constant.

Fig. 9 shows typical results for the variance in the shock position from an ensemble of runs versus time. The hybrid algorithm is used for $p_{\rightarrow} = 0.55$, 0.7 and 0.8 with a time step of 0.05 integrated for two million steps; for each of these simulations the dynamic shock refinement criteria was used (see Section 4.3). The statistics were computed from 400 samples for $p_{\rightarrow} = 0.55$, 800 samples for $p_{\rightarrow} = 0.7$, and 1200 samples for $p_{\rightarrow} = 0.8$, which reflects increasing fluctuations in the shock drift at higher Reynolds number, Re_c . For the intermediate case $(p_{\rightarrow} = 0.7)$ we also ran the pure AERW for 400 sample over a shorter interval, demonstrating that SPDE/AERW hybrid accurately captures the behavior of the system. (Only this case was compared and the pure AERW results are for a shorter time because of the large computational expense of running the ensemble of pure particle simulations.) As expected, the shock diffusion depends on the shock strength with the strongest shock $(p_{\rightarrow} = 0.8)$ exhibiting the most drift.

The most interesting feature observed in these simulations was the absence of shock diffusion in the AR hybrid using a deterministic PDE solver. This deficiency persisted even when the refined (i.e., particle) region was widened by eight cells, roughly doubling its size. The absence of shock diffusion may be, in part, due to the definition of shock location yet alternative ways of measuring the position of the shock are expected to also exhibit significantly reduced diffusion. Given that the localization of shock fronts is an important question addressed in gas dynamics simulations, the suppression of shock diffusion is a cautionary warning that the fidelity of multiscale hybrids may depend on the accuracy of the stochastic modeling.

6. Conclusions and further work

We have constructed a hybrid algorithm that couples an excluded random walk with a viscous Burgers' equation that represent the mean field approximation to the dynamics. The algorithm allows the random walk to be used locally to approximate the solution while modeling the system using the mean field equations in the

remainder of the domain. In tests of the method we have demonstrated that it is necessary to include the effect of fluctuations, represented as a stochastics flux, in the mean field equations to ensure that the hybrid preserved key properties of the system. As expected, not representing fluctuations in the continuum regime leads to a decay in the variance of the solution that penetrates into the particle region. Somewhat more surprising is that the failure to include fluctuations was shown to introduce spurious correlations of fluctuations in equilibrium simulations and for rarefactions. Even more troubling is the observation that using a deterministic PDE solver coupled to the random walk model suppresses the drift of shock location seen with the pure random walk model and with the AR hybrid using a stochastic PDE solver.

We plan to extend this basic hybrid framework to the solution of the compressible Navier–Stokes equations in multiple dimension. For that extension we will use a Direct Simulation Monte Carlo algorithm for the microscopic model coupled to a finite difference approximation to the continuum equations, as described in [17]. The Landau–Lifshitz fluctuating hydrodynamic equations will be used to represent microscopic fluctuations at the continuum level. [18,23]. At present, the challenge remains to establish accurate finite difference schemes for solving these stochastic PDEs.

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