

AN EFFICIENT IMPLICIT FEM SCHEME FOR FRACTIONAL-IN-SPACE REACTION-DIFFUSION EQUATIONS*

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Abstract. Fractional differential equations are becoming increasingly used as a modelling tool for processes associated with anomalous diffusion or spatial heterogeneity. However, the presence of a fractional differential operator causes memory (time fractional) or nonlocality (space fractional) issues that impose a number of computational constraints. In this paper we develop efficient, scalable techniques for solving fractional-in-space reaction diffusion equations using the finite element method on both structured and unstructured grids via robust techniques for computing the fractional power of a matrix times a vector. Our approach is show-cased by solving the fractional Fisher and fractional Allen–Cahn reaction-diffusion equations in two and three spatial dimensions, and analyzing the speed of the traveling wave and size of the interface in terms of the fractional power of the underlying Laplacian operator.

Key words. finite elements, fractional diffusion, numerical solvers

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1. Introduction. Fractional models, in which a standard time or space differential operator is replaced by a corresponding fractional differential operator, have a long history in, for example, physics, finance, and hydrology, with such models often being used to represent so-called anomalous diffusion. In water resources, fractional models have been used to describe chemical and contaminant transport in heterogeneous aquifers [1, 9, 44]. In finance they have been used because of the relationship with certain option pricing mechanisms and heavy-tailed stochastic processes [55]. More recently, fractional models of the Bloch–Torrey equation for describing anomalous diffusion have been considered in the context of magnetic resonance [43]. In cell biology, anomalous diffusion has been measured in fluorescence photobleaching recovery [54] and fractional-in-time models have been developed for simple types of chemical reaction-diffusion equations [69] and to describe microscale diffusion in the cell wall lining of plants [60]. Liu and Burrage [40] have fitted parameters to time fractional gene regulatory models of *Bacillus Subtilis*, a bacterium found in soil.

A number of approaches have been considered for both the time fractional and space fractional differential equations. In the time fractional setting, a common approach is to approximate the fractional-in-time differential operator by a finite difference scheme that has memory going all the way back to the initial condition. Such an approximation can result in explicit [12, 68] or implicit schemes [35]. Fourier series

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and spectral approximations have also been considered [13, 38]. More recently variable order fractional operator problems have been investigated [71]. In the space fractional setting, Liu, Anh, and Turner [39] have considered the numerical approximation of the fractional Fokker–Planck equation. Meerschaert and Tadjeran [46] have used finite difference approximations, and Turner and coauthors have investigated various stability and convergence issues based on finite difference approximations [27, 42, 67]. Liu, Turner, and Cox [41] have also considered a finite volume approach. Li and Xu [37] have considered a spectral approach for the weak solution of the space-time fractional diffusion equation, while Sun and Wu [59] have considered difference schemes for the fractional diffusion wave equation. Many of these approaches have limitations in terms of computational efficiency when two or three spatial dimensions are considered. For example, Roop [50] has considered a finite element approach but computes directly the fractional power of the discrete Laplacian, and this approach does not scale well. Yang et al. [66] have solved the time-space fractional diffusion equation in two spatial dimensions with homogeneous Dirichlet boundary conditions using the matrix transfer technique, and compute the function of a matrix times a vector either by a preconditioned Lanczos (symmetric) or M-Lanczos (nonsymmetric) technique.

The aim of this paper is to obtain a robust, efficient approach that can be equally applicable to fractional-in-space problems in two or three spatial dimensions using a finite element method (FEM) on structured and unstructured grids. The computational heart of this approach is the efficient computation of the fractional power of a matrix times a vector. We will consider three techniques: the contour integral method (CIM), the extended Krylov subspace method (EKSM), and the preassigned poles and interpolation nodes (PAIN) method. We will see that in the first and last cases we can find preconditioners that will allow almost mesh independent convergence.

We will showcase our approach by solving fractional-in-space reaction-diffusion problems of the form

$$(1.1) \quad \frac{\partial u}{\partial t} + K(-\Delta)^\alpha u = g(u), \quad \underline{x} \in \Omega \subset \mathbb{R}^d,$$

using the FEM and a semi-implicit Euler scheme for various $\alpha \in (1/2, 1]$. The system is completed with either a homogeneous Dirichlet boundary condition or a homogeneous Neumann-type boundary condition, where mass is conserved within Ω . A discussion of these boundary conditions can be found in [11]. Standard equations and models that fall into this class include the fractional Allen–Cahn and Fisher equations, as well as, of course, the fractional heat equation when $g = 0$. We have chosen this class of problems as recent work [17] has established how the speed of the traveling wave and the thickness of the interface changes when $\alpha \neq 1$ —see section 6.

The outline of the paper is as follows. In section 2 we review some important issues associated with modeling via fractional differential equations. In section 3 we review the matrix transfer technique that allows us to approximate the fractional Laplacian by the fractional power of a matrix and give a review of relevant computational techniques for computing appropriate matrix functions $f(\mathbf{A})\mathbf{b}$. These include the CIM, EKSM, and PAIN methods. In section 4 we investigate the Neumann formulations of (1.1), which give rise to numerical issues different than Dirichlet formulations as the discrete Laplacian is now singular. Section 5 outlines our approach in terms of iterative methods and preconditioning for the CIM and PAIN methods in terms of mesh independent convergence, while section 6 gives detailed numerical results and analysis of the fractional Fisher and fractional Allen–Cahn models, with particular focus on the relationship between α and the speed of the traveling wave and the size

of the interface. Section 7 gives some conclusions and thoughts for future work.

2. Background to fractional modeling. In order to understand why fractional models are useful and how they might arise, consider the classical advection-diffusion equation which can be derived from two different equations. Let $u(x, t)$ be the particle mass density of some species of particles diffusing and let ν be the constant average velocity of those particles. Fick's law then states that the flux V is the rate at which mass is transported through a unit area and is given by

$$(2.1) \quad V = \nu u - D \nabla u,$$

where D is the diffusion tensor. Conservation of mass requires

$$(2.2) \quad \frac{\partial u}{\partial t} = -\nabla \cdot V,$$

and combining (2.1) and (2.2) gives the classical advection-diffusion equation

$$(2.3) \quad \frac{\partial u}{\partial t} = -\nu u + \nabla \cdot D \nabla u.$$

However, the assumptions that underlie this approach can be questioned for heterogeneous media [64]. The continuum hypothesis uses local averaged values, but averaged quantities fluctuate wildly as the averaging volume becomes smaller. This is also related to the homogenization principle that is fundamental for predicting macroscopic properties from microscopic features [26] that are often assumed to be independent. But in some settings this independence between the microscopic and the macroscopic may not hold and the homogenization principle can fail. It is in this setting that fractional models can offer insights that traditional approaches do not offer, especially for the case of diffusion in heterogeneous environments.

In heterogeneous structures such as those possessing spatial connectivity, movement of particles may be facilitated within a certain scale—so-called superdiffusion. Thus the spatial complexity of the environment imposes geometric constraints on the transport processes on all length scales, which can be interpreted as temporal correlations on all time scales. Nonhomogeneities of the medium may fundamentally alter the laws of Markov diffusion, leading to long range fluxes and non-Gaussian, heavy-tailed profiles [6, 47], and these motions may no longer obey Fick's law [70]. Schumer et al. [56] demonstrated that a fractional Fick's law is a governing equation for solute transport in porous media where temporally correlated velocity fields do not dominate transport processes. A fractional Fick's law naturally implies spatial and temporal nonlocality and can be derived from rigorous approaches using spatial averaging theorems and measurable functions [10].

Meerschaert, Mortensen, and Wheatcraft [45] define a fractional Fick's law as

$$(2.4) \quad V = \nu u - D \nabla^\beta u,$$

where $\beta = 2\alpha - 1$ and $\alpha \in (1/2, 1]$. Combined with (1.2) this gives

$$(2.5) \quad \frac{\partial u}{\partial t} = -\nu u + \nabla \cdot D \nabla^\beta u.$$

In three spatial dimensions,

$$(2.6) \quad \nabla^\beta = \left(\frac{\partial^\beta}{\partial x^\beta}, \frac{\partial^\beta}{\partial y^\beta}, \frac{\partial^\beta}{\partial z^\beta} \right)^T.$$

This is known as the Riemann–Louiville fractional gradient vector. The components of this vector are fractional derivatives which, for $\beta \in (0, 1)$, take the form

$$(2.7) \quad \frac{\partial^\beta}{\partial x^\beta} f(x, y, z) = \frac{1}{\Gamma(1-\beta)} \frac{\partial}{\partial x} \int^x \frac{f(s, y, z)}{(x-s)^\beta} ds,$$

with similar expressions for $\frac{\partial^\beta f}{\partial y^\beta}$ and $\frac{\partial^\beta f}{\partial z^\beta}$ [49].

We can relate fractional derivatives to the underlying behavior of an ensemble of particles undergoing a continuous time random walk. When a fractional derivative replaces the Laplacian operator in the diffusion equation, the resulting solution describes the probability density function of particles undertaking a heavy-tailed random walk, where occasional large jumps dominate the more common smaller jumps. On the other hand, a fractional time derivative leads to subdiffusion in which the waiting time between particle jumps has a long-tailed probability and where the mean square displacement of an ensemble of particles behaves as t^α , $\alpha \in (0, 1]$. The fractional-in-space diffusion equation

$$(2.8) \quad \frac{\partial p}{\partial t} = -K(-\Delta)^\alpha p$$

can be viewed as describing the probability density function of particles undergoing superdiffusion, where $(-\Delta)^\alpha$ is the fractional Laplacian operator. Thus it makes sense that this equation should play the fundamental role in using fractional models.

The relationship between (2.5) and (2.8) is not trivial and so Turner, Ilić, and Perr [60] work with potential theory. They define a potential $\bar{u} = (-\Delta)^{\alpha-1}u$ and note that since $\alpha - 1 < 0$, the expression for the potential can be interpreted as an integral. Thus the fractional gradient ∇^α can be interpreted as $\nabla^\alpha = \nabla(-\Delta)^{\alpha-1}$. In the case that $D = KI$, (2.5) becomes

$$(2.9) \quad \frac{\partial u}{\partial t} = -\nu u - K(-\Delta)^\alpha u.$$

There are some subtle issues associated with the interpretation of the fractional Laplacian and, for example, the Riesz spatial derivatives. These issues manifest when considering boundary conditions, and have been discussed, for example, in [65]. Guan and Ma [21] note that a $(-\Delta)^\alpha$ -harmonic function on a finite domain cannot be determined by its value on the boundary of that domain, but on the whole space minus the domain. On the other hand, Ilić et al. [27] have shown that problems of the form (2.9) with $\nu = 0$ are well defined on finite domains with homogeneous boundary conditions of Dirichlet, Neumann, and Robin type. They show this by noting that $(-\Delta)^\alpha$ has the same interpretation as $-\Delta$ in terms of its spectral decomposition for these homogeneous boundary conditions. These apparent contradictions are due to the differences in the definition of the fractional Laplacian; see [65] for further discussion. In the latter case the fractional Laplacian can be dealt with by using the matrix transfer technique that was introduced in [28, 29]. In essence, the spatial discretization of the fractional Laplacian is obtained by first finding a matrix representation, \mathbf{A} , of the Laplacian (whether it is by finite difference, finite element, or finite volume) and raising it to the same fractional power \mathbf{A}^α .

The case of nonhomogeneous boundary conditions is more complicated, and there is much confusion in the literature. Ilić et al. [28] decompose the operator as $(-\Delta)^\alpha = (-\Delta)^{\alpha-1}(-\Delta)$, where the rightmost operator is used to deal with the nonhomogeneous

boundary conditions for which it is well defined. They then use the matrix transfer technique to solve an ordinary differential equation which requires both the computation of \mathbf{A}^α and $\mathbf{A}^{\alpha-1}$ operating on different vectors. Gunzburger and Lehoucq [22] have considered a nonlocal second order scalar elliptic boundary value problem augmented with nonlocal Dirichlet or Neumann boundary conditions, and under appropriate conditions this reduces to the standard setting. This is an example of a more extensive class of problems with general nonlocal operators. Andreu et al. [3, 4] have considered p -Laplacian diffusion equations with Dirichlet and Neumann boundary conditions. Du and Zhou [16] show through peridynamic nonlocal continuum theory that nonlocal diffusion exhibits multiscale behavior beyond that of standard diffusion.

3. Modeling and computational methods. In this paper we will adopt an FEM approach for solving (1.1), as such problems often arise in heterogeneous settings and FEMs are appropriate for unstructured meshes. Let \mathbf{L} be the standard FEM stiffness matrix with appropriate boundary conditions, and let \mathbf{M} be the corresponding mass matrix. Following the work of Ilić et al. [28], we employ the matrix transfer technique, which states that the error introduced by approximating the fractional Laplacian by a fractional power of the matrix $\mathbf{A} = \mathbf{M}^{-1}\mathbf{L}$ converges at the same rate as the underlying FEM discretization. Both \mathbf{L} and \mathbf{M} are real and symmetric, \mathbf{L} is nonnegative definite, and \mathbf{M} is positive definite. Noting that $\mathbf{M}^{1/2}\mathbf{A}\mathbf{M}^{-1/2} = \mathbf{M}^{-1/2}\mathbf{L}\mathbf{M}^{-1/2}$ is real, symmetric, and similar to \mathbf{A} , it follows that the spectrum $\sigma(\mathbf{A})$ of \mathbf{A} is real. Furthermore the nonnegative definiteness of \mathbf{L} implies $x^T\mathbf{M}^{-1/2}\mathbf{A}\mathbf{M}^{1/2}x \geq 0$ so that \mathbf{A} is also nonnegative definite and $\sigma(\mathbf{A}) \subset [0, \infty)$.

When $\mathbf{A} \in \mathbb{R}^{N \times N}$ is diagonalizable and a function f is defined on the spectrum $\sigma(\mathbf{A})$, the matrix function $f(\mathbf{A})$ can be defined [25, Definition 1.2] as

$$(3.1) \quad f(\mathbf{A}) = \mathbf{Q}f(\mathbf{\Lambda})\mathbf{Q}^{-1},$$

where $\mathbf{\Lambda} = \text{diag}([\lambda_1, \lambda_2, \dots, \lambda_N])$ is a diagonal matrix of the eigenvalues $\lambda_1 \leq \lambda_2, \dots, \leq \lambda_N$ of \mathbf{A} , \mathbf{Q} is a matrix of corresponding eigenvectors, and $[f(\mathbf{\Lambda})]_{j,j} = f(\lambda_j)$. If f is analytic in a connected open subset $\mathcal{H}(f)$ of the complex plane such that $\sigma(\mathbf{A}) \subset \mathcal{H}(f)$, then an equivalent definition [25, Definition 1.11 and Theorem 1.12] is given by

$$(3.2) \quad f(\mathbf{A}) = \frac{1}{2\pi i} \int_{\Gamma} f(z) (z\mathbf{I} - \mathbf{A})^{-1} dz,$$

where $\Gamma \subset \mathcal{H}(f)$ is a closed contour winding once around the spectrum $\sigma(\mathbf{A})$ in the counterclockwise direction.

Discretizing (1.1) semi-implicitly in time with a backward Euler approximation leads, at the m th time step, to linear systems of the form

$$(3.3) \quad (\mathbf{I} + K\Delta t(\mathbf{M}^{-1}\mathbf{L})^\alpha)u_{m+1} = u_m + \Delta t g(u_m).$$

The matrix function we then require is

$$(3.4) \quad f(z) = 1/(1 + K\Delta t z^\alpha),$$

with $\mathcal{H}(f) \subset \mathbb{C} \setminus (-\infty, 0]$. Note that functions of the same form arise when using finite difference approximations to fractional-in-time operators [65, 67], and so the methods we discuss in the next two sections are also applicable in a time-and-space fractional setting. When \mathbf{A} is nonsingular either definition (3.1) or (3.2) can be applied,

but when the smallest eigenvalue $\lambda_1 = 0$, the hypothesis that $\sigma(\mathbf{A})$ is contained in $\mathcal{H}(f)$, and (3.2) cannot be used directly. We discuss this issue further in section 4.

Higham has recently published an extensive monograph [25] on the computation of matrix functions, however the problem $f(\mathbf{M}^{-1}\mathbf{L})$ we consider is so large and dense (even, for example, when $f(x) = x$) that forming it explicitly is infeasible; specialized techniques are required. Fortunately it is not $f(\mathbf{M}^{-1}\mathbf{L})$ we need in (3.3), but rather its action on a vector, and in this section we consider recent methods which allow efficient (and scalable) computation of $f(\mathbf{M}^{-1}\mathbf{L})b$ by taking advantage of the sparsity and structure of \mathbf{M} and \mathbf{L} .

We consider three different but related approaches. The first is a contour integral method (CIM) [24] based upon a quadrature discretization of (3.2). The other two, the extended Krylov subspace method (EKSM) [15, 33] and the preassigned poles and interpolation nodes (PAIN) method [23], are rational Krylov subspace-based techniques. Further comparisons and far more detailed descriptions of these and other rational $f(\mathbf{A})b$ methods can be found in two recent PhD theses, [23] and [58].

In what follows we assume the matrix function f is analytic in $\mathcal{H}(f) \subset \mathbb{C} \setminus (-\infty, 0]$, that the matrix \mathbf{A} has a known decomposition $\mathbf{A} = \mathbf{M}^{-1}\mathbf{L}$ where \mathbf{M} and $\mathbf{L} \in \mathbb{R}^{N \times N}$ are symmetric positive definite, and that the spectrum $\sigma(\mathbf{A})$ of \mathbf{A} is contained in an interval $[\lambda_1, \lambda_N] \subset (0, \infty)$ where λ_1 and λ_N are known (or can at least be computed). In section 4 we extend the discussion to the case where $\lambda_1 = 0$. Furthermore, we suppose that both the action of \mathbf{A} and shifted solves of the form $(\mathbf{A} - \sigma\mathbf{I})x = b$ can be computed efficiently. Further discussion of these linear solves is found in section 5.

3.1. Contour Integral Method. Until recently, Definition 3.2 was rarely exploited for numerical computation, but Hale, Higham, and Trefethen [24] propose efficient quadrature approximations for computing $f(\mathbf{A})b$ when $\sigma(\mathbf{A})$ lies on or near the positive real axis and $\mathcal{H}(f) = \mathbb{C} \setminus (-\infty, 0]$. They show that although a naïve discretization of (3.2) leads to methods requiring a number of quadrature points that increases linearly with the condition number λ_1/λ_N , choosing the contour Γ wisely can lead to methods where the number of quadrature nodes needed to obtain a specified accuracy increases asymptotically as $\log(\lambda_1/\lambda_N)$ [24, Theorem 2.1]. Since the number of nodes equates to the number of linear solves required, this makes the CIM approximation to $f(\mathbf{A})b$ “almost” optimal.

The contour is chosen by a conformal map from an annulus $\mathcal{A}_{1,R}$ to $\mathbb{C} \setminus \{(-\infty, 0] \cup [\lambda_1, \lambda_N]\}$ (Figure 3.1), and the quadrature nodes and weights are given by mapping the trapezoid or midpoint rule applied in the annulus using the same map. This leads to an approximation of the form

$$(3.5) \quad f(\mathbf{A})b \approx r_n(\mathbf{A})b = \sum_{j=1}^n w_j (\xi_j \mathbf{I} - \mathbf{A})^{-1} b,$$

where the ξ_j are complex shifts lying on Γ . Equation (3.5) can be interpreted as a rational approximation to $f(\mathbf{A})b$ given directly in partial fraction form. When $\mathbf{A} = \mathbf{M}^{-1}\mathbf{L}$, this can be rewritten as

$$(3.6) \quad r_n(\mathbf{M}^{-1}\mathbf{L})b = \sum_{j=1}^n w_j (\xi_j \mathbf{M} - \mathbf{L})^{-1} \mathbf{M}b.$$

Note that \mathbf{A} itself need never be formed explicitly. A particular benefit of the CIM is that each of the solves on the right-hand side of (3.6) are independent, making this method easily parallelizable.

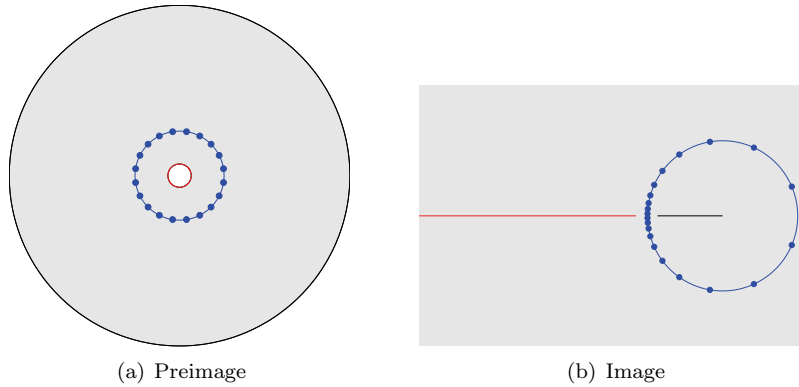


FIG. 3.1. Conformal map from the annulus $\mathcal{A}_{1,R}$ (a) to the slit domain $\mathbb{C} \setminus \{(-\infty, 0] \cup [\lambda_1, \lambda_N]\}$ (b). The dots are the quadrature points in the CIM (i.e., the shifts used in the linear solves) that lie on a circle of radius $\lambda_N \sqrt{1 + \lambda_1/\lambda_N}$ about the point λ_N . For real matrices, only the quadrature points in the upper half-plane are needed.

Furthermore, since \mathbf{A} is real the integrand in (3.2) is real-symmetric and $f(\mathbf{A})$ is twice the real part of the value obtained by integrating over the first half of the contour. This means that we need only half of the quadrature points (i.e., those in the upper half-plane) and therefore compute only half of the shifted solves. To simplify later discussion we will denote by the subscript of r the number of linear solves in the computation, rather than the degree of the underlying rational interpolant, and refer to the n -point CIM approximation in the same manner.

Given $L, M \in \mathbb{R}^{N \times N}$, $\mathbf{b} \in \mathbb{R}^N$, a function \mathbf{f} , and an integer n , the following MATLAB code returns \mathbf{v} , the n -point CIM quadrature approximation to $\mathbf{f}(M \setminus L)\mathbf{b}$. The routines `ellipkqp.m` and `ellipjcc.m` on lines 3 and 5 are from Driscoll's Schwarz–Christoffel toolbox [14] but can be replaced by built-in MATLAB routines as described in [24]. Line 11 uses the MATLAB `\` command to solve the linear systems in (3.6), although more practical methods of doing this when N is large are discussed in section 5. The `parfor` command on line 10 allows each of the shifted linear solves to be solved in parallel when the MATLAB parallel toolbox is available.

```

0 % Contour Integral Method, adapted from [26, method1.m]
1 l = eigs(L,M,2,'BE'); l1 = l(1); lN = l(2); % Spectrum of M\L
2 k = (sqrt(lN/l1)-1)/(sqrt(lN/l1)+1); % A convenient constant
3 [K Kp] = ellipkqp(-log(k)/pi); % Elliptic integrals
4 t = .5i*Kp-K+(n-.5:-1:0)*2*K/n; % Midpoint rule points
5 [sn cn dn] = ellipjcc(t,-log(k)/pi); % Jacobi elliptic functions
6 xi = sqrt(l1*lN)*(1/k+sn)./(1/k-sn); % Quadrature nodes
7 dxidt = cn.*dn./(1/k-sn).^2; % Derivative wrt t
8 wts = f(xi).*dxidt; % Quadrature weights
9 v = zeros(length(b),1); % Initialize output
10 parfor j = 1:n % Parallel solves
11     y = (xi(j)*M-L)\(M*b); % Solve using backslash
12     v = v + wts(j)*y; % Update solution vector
13 end
14 v = -4*K*sqrt(l1*lN)*imag(v)/(k*pi*n); % Scaling

```

3.2. Krylov subspace methods. Krylov subspace methods are well known for their application in solving linear systems (see, for example, [61, 62]). Although rarely considered as an $f(\mathbf{A})b$ problem, i.e., the application of a matrix function to a vector, such linear solves can certainly be viewed as the special case where $f(z) = z^{-1}$. The same subspace projection ideas have been extended to more general functions [61].

From a *polynomial Krylov space* \mathcal{K}_r of order r ,

$$(3.7) \quad \mathcal{K}_r(\mathbf{A}, b) = \text{span}\{b, \mathbf{A}b, \dots, \mathbf{A}^{r-1}b\},$$

the *Rayleigh–Ritz approximation* to $f(\mathbf{A})b$ is given by

$$f_r(\mathbf{A})b = \mathbf{V}_r f(\mathbf{A}_r) \mathbf{V}_r^* b,$$

where \mathbf{A}_r is the *Rayleigh quotient* $\mathbf{A}_r = \mathbf{V}_r^* \mathbf{A} \mathbf{V}_r$, and the columns of \mathbf{V}_r , which form a basis of $\mathcal{K}_r(\mathbf{A}, b)$, are computed by a stabilized Gram–Schmidt process. It is typically assumed that r is not large, so that $f(\mathbf{A}_r)b$ can be computed directly by, say, an eigenvalue decomposition as in Definition 3.1 or by other methods described by Higham [25]. The Krylov spaces (3.7) are nested, i.e., $\mathcal{K}_{r+1}(\mathbf{A}, b) \subset \mathcal{K}_r(\mathbf{A}, b)$, so the basis can be updated efficiently and the approximation refined iteratively.

The Rayleigh–Ritz approach can be interpreted as a polynomial approximation

$$f_r(\mathbf{A})b = p_{r-1}(\mathbf{A})b \approx f(\mathbf{A})b,$$

where the degree $r - 1$ polynomial p_{r-1} interpolates f at the Ritz values $\sigma(\mathbf{A}_r)$ [53]. A natural extension to *rational Krylov methods* based upon rational interpolation was introduced by Ruhe [51, 52]. The motivation here is that the additional cost of needing to solve linear systems in the approximation is more than compensated for by increased rates of convergence, and the overall computational work required can be significantly reduced.

Given a polynomial q_{r-1} of degree $r - 1$ with no poles in $\sigma(\mathbf{A})$, the *rational Krylov space* of order r associated with (\mathbf{A}, b, q_{r-1}) is given by

$$\mathcal{Q}_r(\mathbf{A}, b; q_{r-1}) = \mathcal{K}_r(\mathbf{A}, q_{r-1}(\mathbf{A})^{-1}b).$$

Again it is convenient in computations if the \mathcal{Q}_j are nested, so typically a sequence of *poles* $\{\xi_1, \xi_2, \dots\} \subset \{\mathbb{C} \cup \{\infty\}\} \setminus \sigma(\mathbf{A})$ is defined, and

$$(3.8) \quad q_r(z) = \prod_{j=1}^r (z - \xi_j), \quad r = 1, 2, \dots$$

Once the rational functions q_{r-1} , and hence the subspaces \mathcal{Q}_r , have been chosen, we then obtain the *rational Rayleigh–Ritz approximation*

$$f_r(\mathbf{A})b = \mathbf{V}_r f(\mathbf{A}_r) \mathbf{V}_r^* b,$$

where $\mathbf{A}_r = \mathbf{V}_r^* \mathbf{A} \mathbf{V}_r$, and the orthogonal columns of \mathbf{V}_r form a basis of the now rational Krylov subspace $\mathcal{Q}_r(\mathbf{A}, b)$.

If all $\xi_j = \infty$ in (3.8), then the rational Krylov method reduces to the polynomial Rayleigh–Ritz method described above. Choosing $\xi_{2j} = \infty, \xi_{2j+1} = 0$ gives the *extended Krylov subspace method* (EKSM) [15, 33], discussed below.

3.2.1. Extended Krylov subspace method. The EKSM method, introduced by Druskin and Knizhnerman [15] and recently revived by Knizhnerman and Simoncini [33], is the special case of the rational Krylov methods where the poles $\{\xi_{2j+1}\}$ are all chosen to be at the origin. This simple choice of pole locations means EKSM approximations to $f(\mathbf{A})b$ are generated from the rational subspace

$$\mathcal{K}_{2r}(\mathbf{A}, \mathbf{A}^{-r}b),$$

of which an orthogonal basis $\mathbf{V}_{2r} = [V_2, V_4, \dots, V_{2r}]$ can be constructed by applying Gram–Schmidt orthogonalization to

$$\hat{V}_2 = [b, \mathbf{A}^{-1}b], \quad \hat{V}_{2r} = [\mathbf{A}V_{2r-2}^{(1)}, \mathbf{A}^{-1}V_{2r-2}^{(2)}], \quad r > 1.$$

Here $\mathbf{A} = \mathbf{M}^{-1}\mathbf{L}$, so the \hat{V} are given by

$$\hat{V}_{2r} = [\mathbf{M}^{-1}\mathbf{L}V_{2r-2}^{(1)}, \mathbf{L}^{-1}\mathbf{M}V_{2r-2}^{(2)}],$$

and solves with both \mathbf{M} and \mathbf{L} are required for each dimension of the subspace (except the first). To remain consistent with the CIM method, we therefore refer to the *degree* n EKSM approximation $r_n(\mathbf{A})b$ of $f(\mathbf{A})b$ as that which requires n linear solves to compute, and so $n = 2r - 1$ in the above.

For real-symmetric positive definite matrices \mathbf{A} with spectrum in $[\lambda_1, \lambda_N]$ and functions analytic on $\mathbb{C} \setminus (-\infty, 0]$, it can be shown [8, 33] that convergence of the EKSM satisfies

$$(3.9) \quad \|f(\mathbf{A})b - r_n(\mathbf{A})b\| = O(\exp(-n\sqrt[4]{\lambda_1/\lambda_N})).$$

While this bound suggests that the number of linear solves required to obtain a certain accuracy will increase more rapidly here than in the CIM as the condition number rises (i.e., to the 1/4th power, rather than logarithmically), it is possible to show that, for symmetric matrices, this is an optimal rate for a single pole location [8].

Pseudocode for computing the EKSM was given in the original paper [15], and Simoncini gives MATLAB code in [57] which is easily modified to account for $\mathbf{A} = \mathbf{M}^{-1}\mathbf{L}$ without forming \mathbf{A} explicitly.

3.2.2. Preassigned poles and interpolation nodes. In addition to prescribing the poles $\{\xi_j\}$ that determine the rational Krylov subspace, we can also select the interpolation nodes $\{\alpha_j\}$. The problem then becomes even more closely related to that of best rational approximation, which is itself related to potential theory. While the task of determining a best approximation is known to be a difficult one, there are a number of ways of determining asymptotically optimal points. Levin and Saff [36] give a detailed discussion of the varying approaches, and we recall some of that information here.

Fekete points, which give the supremum of

$$(3.10) \quad \prod_{\substack{j \neq k \\ 1 \leq j, k \leq n}} \frac{|\xi_j - \xi_k| |\alpha_j - \alpha_k|}{|\xi_j - \alpha_k|^2}$$

for $\xi_k \in (-\infty, 0]$, $\alpha_k \in [\lambda_1, \lambda_N]$, are considered the “cleanest” mathematically but are difficult to compute numerically and rarely used in practice. *Leja–Bagby points* are an approximation to Fekete points whereby an initial pole and node $\{\xi_1, \alpha_1\}$ are chosen

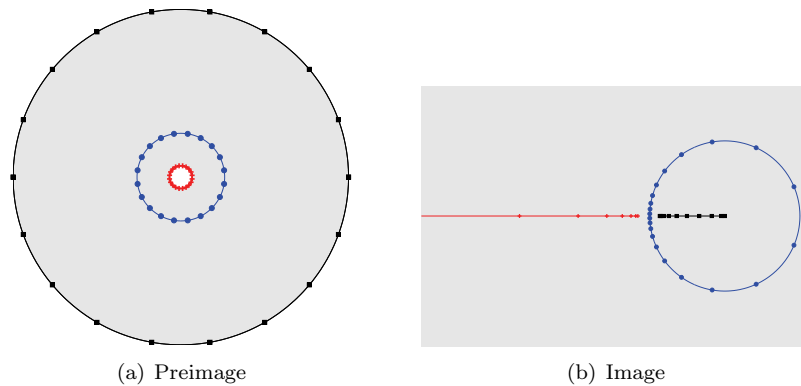


FIG. 3.2. *Fejér–Walsh poles (crosses) and interpolation nodes (squares). Importantly, these are both real.*

arbitrarily and $\{\xi_k, \alpha_k\}, k = 2, \dots, n$ are determined incrementally to minimize the energy

$$(3.11) \quad \prod_{j=1}^{k-1} \frac{|\xi - \xi_j| |\alpha - \alpha_j|}{|\xi - \alpha_j| |\alpha - \xi_j|}$$

for each k . Still these points are cumbersome to compute, and typically the allowed values for $\{\xi_k\}$ and $\{\alpha_k\}$ are restricted to discrete subsets of $(-\infty, 0]$ and $[\lambda_1, \lambda_N]$. A code for solving this discrete problem, `lejadc`, appears in the appendix.

When $\mathcal{H}(f)$ is closed, connected, and disjoint from $[\lambda_1, \lambda_N]$, the *Fejér–Walsh points* are defined by the image of equally spaced points on the boundary of an annulus $\mathcal{A}_{1,R}$ under the unique conformal map from $\mathcal{A}_{1,R}$ to $\mathcal{S} = \mathcal{H}(f) \setminus [\lambda_1, \lambda_N]$. In the problem we consider $\mathcal{S} = \mathbb{C} \setminus \{(-\infty, 0] \cup [\lambda_1, \lambda_N]\}$, and the map is precisely that given for the CIM appearing in Figure 3.1. We repeat the map in Figure 3.2, which now contains the Fejér–Walsh interpolation nodes (squares) and poles (crosses) alongside those from the CIM (circles). Since this map is available, appearing in [24] and [2, section 49], we follow the suggestion of Levin and Saff [36, section 7.c] that these Fejér–Walsh points can be “more efficient” than the other possibilities discussed above. Some brief experiments, at least with discretely chosen Leja–Bagby points, gave similar results.

As with the polynomial Krylov approximation, the PAIN method can be thought of as the evaluation of an interpolant (now rational) in Newton form. As such, certain stability issues arise. In particular we must sensibly order the nodes (and poles) that define the denominators $q_r(z)$, and hence the subspaces \mathcal{Q}_r . We follow the suggestion of Reichel [48] by ordering the Fejér–Walsh points so that (3.11) is satisfied for each $k = 1, \dots, n$, which can be done using the `lejadc` code mentioned above.

The convergence properties of the PAIN method are directly related to those of the underlying rational approximant r_n on the spectrum of \mathbf{A} . Güttel [23] shows that, when choosing the nodes and poles of the rational approximant in any of the ways described above, we have

$$(3.12) \quad \limsup_{n \rightarrow \infty} \|f - r_n\|_{[\lambda_1, \lambda_N]}^{1/n} \leq R^{-1},$$

where R is the outer radius of the annulus $\mathcal{A}_{1,R}$ used to define the Fejér–Walsh points appearing in Figures 3.1(a) and 3.2(a). By the same argument as in [24, Theorem 2.1],

one can show that the degree of the rational function needed to obtain a specified accuracy increases only logarithmically with the condition number λ_N/λ_1 . Güttel also shows that this convergence rate is not more than a factor of two worse than the rational best uniform approximation r_n^* of type $(m-1, m-1)$ to f on $[\lambda_1, \lambda_N]$ [23, Remark 7.7]. For complex matrices, the CIM yields a result similar to (3.12), but with the R being replaced by \sqrt{R} . However, as described in the previous section, when the matrix \mathbf{A} is real we need only use the quadrature points in the upper half-plane, which effectively replaces n with $2n$ in the exponent and recovers (3.12).

Given $L, M \in \mathbb{R}^{N \times N}$, $\mathbf{b} \in \mathbb{R}^N$, a matrix function \mathbf{f} , and an integer n , the MATLAB code below returns \mathbf{v} , the degree n Fejér–Walsh PAIN approximation to $\mathbf{f}(M \setminus L)\mathbf{b}$. Here \mathbf{f} is a matrix version of the function f suitable for evaluating the lower triangular subspace projection $\mathbf{A}\mathbf{r}$, such as that returned by the MATLAB routine `funm`. The inverse action of $f(\mathbf{A})$ can be applied efficiently by uncommenting line 9. The vectors `xi` and `alpha` are given by line 6 in the CIM code, but with line 4 replaced by `t=1i*Kp-K+(n-.5:-1:0)*2*K/n` and `t-K+(n-1:-1:0)*2*K/(n-1)` respectively. The function `lejadc` (which can be found in the appendix) Leja-sorts the nodes and poles as described above. Again we use the MATLAB `\` command to solve the linear systems in line 4, but discuss in section 5 more practical methods for large N .

```

0 % PAIN method, adapted from Guettel's thesis [25]
1 v = zeros(length(b),1); K = eye(n); H = diag(alpha); % Initialize
2 [alpha,xi] = lejadc(alpha,xi); % Leja-sort the nodes.
3 for r = 1:n
4     w = (M*xi(r)-L)\((L-alpha(r)*M)*b); % New basis vector
5     K(r+1,r) = norm(w); % Update K
6     H(r+1,r) = xi(r)*norm(w); % Update H
7     Ar = H(1:r,1:r)/K(1:r,1:r); % Rayleigh quotient
8     fr = f(Ar); % Matrix function
9     % fr = fr\[1 ; zeros(r-1,1)]; % Uncomment for f(A)\b
10    v = v + fr(r,1)*b; % Update solution vector
11    b = w/norm(w); % Renormalize b
12 end

```

3.3. Comparison of CIM, EKSM, and PAIN. We end this section by testing the three $f(\mathbf{A})\mathbf{b}$ methods discussed above on the problem

$$(3.13) \quad u - K(-\Delta u)^\alpha = x(x-1)y(y-1) \quad \text{in } \Omega = (0,1) \times (0,1),$$

with the Dirichlet boundary condition $u(\partial\Omega) = 0$.

We use a sequence of six uniformly refined triangulations of the domain Ω and solve all linear systems using the MATLAB `\` command. We take $K = 0.01$ and show in Figure 3.3 results for $\alpha = 0.75$, which seem representative of the results for all $\alpha \in (0.5, 1)$. The x -axis shows n , the number of linear solves, which are assumed to be the expensive stage of the computation, and the y -axis shows the infinity norm of the error. For the first two levels the reference solution is computed using the MATLAB `mpower` command, and for the more refined grids where this becomes too expensive we choose the 100-point CIM quadrature approximation. Timings are given in Table 3.1, and all computations were performed on an AMD 2.6GHz tricore desktop in MATLAB 7.12.

Figure 3.3 shows that, as a function of the number of linear solves, the convergence of the CIM and the PAIN method are very similar and closely follow the bound (3.12). Although degrading more as the size of the mesh increases, as suggested by (3.9), the

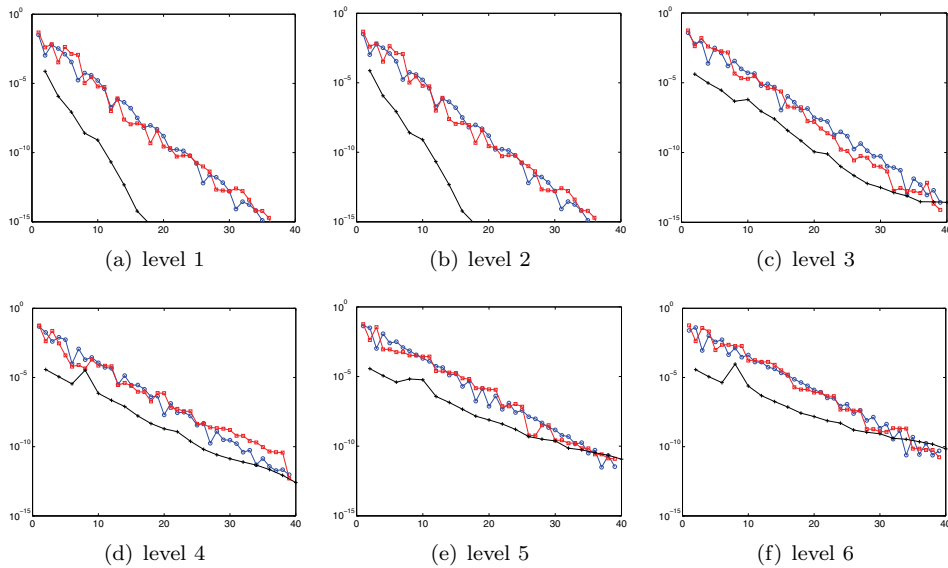


FIG. 3.3. Error in the CIM ($\circ-\circ$), EKSM ($+--+$), and PAIN ($\square-\square$) approximations to the matrix function $f(\mathbf{A})\mathbf{b}$ arising from a FEM approximation to (3.13) as a function of n , the number of linear solves, on a series of increasingly finer meshes.

TABLE 3.1

Timing results for solving (3.13) using the CIM, PAIN, and EKSM methods of degree $n = 40$ (i.e., involving 40 linear solves with backslash). The complex-valued shifts of the CIM make it up to a factor of 2 slower than PAIN, whereas the simplicity of the solves in EKSM makes it around twice as fast.

Level	N	Cond. no.	CIM time	PAIN time	EKSM time
1	145	3,894	0.0262	0.0291	0.0193
2	545	14,488	0.1407	0.3592	0.0563
3	2,113	57,486	0.8678	0.6704	0.2848
4	8,321	230,020	4.7929	2.7610	1.7202
5	33,025	920,038	27.218	15.499	8.5777
6	131,585	3,680,178	165.01	90.953	52.112

curves for the EKSM typically lie below those for the CIM and PAIN methods. In particular, the EKSM gives a surprisingly accurate solution even after just two solves, as it captures the contribution from the unitary eigenvalues arising from the Dirichlet conditions, a phenomenon described in [7]. Recalling that the shifts, and hence all the arithmetic, in the PAIN and EKSM methods are real, it is to be expected that the CIM has the larger computational time when n is large and the linear solves dominate the computational cost. However, it is slower only by a factor of a little less than 2; later in the next section we solve this problem on a 3 core machine, where the timings are comparable. There are no shifts involved in the EKSM, and this makes the solves faster than those in the PAIN method by around a factor of 2.

The CIM is based upon a midpoint rule quadrature discretization, so for each n in Figure 3.3 the shifts used in the linear solves can only be reused each time n is tripled (using the trapezium rule they repeat each time n is doubled). Conversely, the results of the PAIN and EKSM methods are iterative and so need only be applied once. This allows computing the interpolation points and poles corresponding to some

given higher degree rational function, and iterating until some convergence criterion such as [58, Theorem 3.7] is achieved. However, for the application we have in mind where the same matrix function will be computed for many different right-hand sides in a semi-implicit time-stepping scheme, the number of quadrature points (or degree of the rational function) needed to obtain a specified accuracy can be computed offline in advance at little cost.

4. Neumann problems. So far our discussion has been restricted to the Dirichlet problem, where the matrix \mathbf{A} is nonsingular. However, many applications require Neumann-type conditions, such as $\frac{\partial u}{\partial n} = 0$ on $\partial\Omega$, which make the discretized matrix \mathbf{A} singular. In this section we describe extensions to the methods discussed previously, which allow practical computation of $f(\mathbf{A})b$ in such a situation.

We begin by describing a modification of the CIM for singular \mathbf{A} . The contour integral Definition 3.2 of $f(\mathbf{A})$ cannot be applied here, as $(-\infty, 0]$ and $\sigma(\mathbf{A}) = [0, \lambda_N]$ are no longer disjoint, and we cannot find a contour Γ in $\mathcal{H}(f)$ surrounding $\sigma(\mathbf{A})$. However, \mathbf{A} is still diagonalizable, and Definition (3.1) still holds. Theorem 4.1 below suggests how $f(\mathbf{A})b$ might be computed up to the addition of a constant vector by a contour integral which simply ignores the zero eigenvalue.

THEOREM 4.1. *Let $\mathbf{A} = \mathbf{Q}\text{diag}(\lambda_1, \lambda_2, \dots, \lambda_N)\mathbf{Q}^{-1}$ be a diagonalizable matrix. If f is a function analytic in an open connected set $\mathcal{H}(f) \subset \mathbb{C}$ containing $\lambda_2, \lambda_3, \dots, \lambda_N$ and defined at λ_1 , then*

$$f(\mathbf{A}) = \frac{1}{2\pi i} \int_{\Gamma_2} f(z)(z\mathbf{I} - \mathbf{A})^{-1} dz + f(\lambda_1)\mathbf{Q}(1, :)\mathbf{Q}^{-1}(:, 1),$$

where $\Gamma_2 \subset \mathcal{H}(f)$ is a closed contour containing $\lambda_2, \lambda_3, \dots, \lambda_N$, but not λ_1 .

Proof. To simplify notation let $\mathbf{S} = \mathbf{Q}^{-1}$, $s^T = \mathbf{S}(:, 1)$, and $q = \mathbf{Q}(:, 1)$ (the eigenvector corresponding to the eigenvalue λ_1 , which spans the null-space if $\lambda_1 = 0$). Definition (3.1) implies

$$\begin{aligned} f(\mathbf{A}) &= \mathbf{Q}f(\mathbf{\Lambda})\mathbf{S} \\ &= \begin{pmatrix} q & \mathbf{Q} \end{pmatrix} \begin{pmatrix} f(\lambda_1) & \\ & f(\overline{\mathbf{\Lambda}}) \end{pmatrix} \begin{pmatrix} s^T \\ \overline{\mathbf{S}} \end{pmatrix} \\ (4.1) \quad &= f(\lambda_1)qs^T + \mathbf{Q}f(\overline{\mathbf{\Lambda}})\overline{\mathbf{S}}, \end{aligned}$$

where $\underline{\mathbf{Q}} = \mathbf{Q}(:, 2:\text{end})$, $\overline{\mathbf{S}} = \mathbf{S}(2:\text{end}, :)$, and $\overline{\mathbf{\Lambda}} = \mathbf{\Lambda}(2:\text{end}, 2:\text{end})$. We can expand the inverse of $(z\mathbf{I} - \mathbf{A})$ in the same way to find

$$(z\mathbf{I} - \mathbf{A})^{-1} = \frac{1}{z - \lambda_1}qs^T + \underline{\mathbf{Q}}(z\overline{\mathbf{I}} - \overline{\mathbf{\Lambda}})^{-1}\overline{\mathbf{S}},$$

which, multiplying by $f(z)$ and integrating around Γ_2 , gives

$$(4.2) \quad \int_{\Gamma_2} f(z)(z\mathbf{I} - \mathbf{A})^{-1} dz = qs^T \int_{\Gamma_2} \frac{f(z)}{z - \lambda_1} dz + \underline{\mathbf{Q}} \int_{\Gamma_2} f(z)(z\overline{\mathbf{I}} - \overline{\mathbf{\Lambda}})^{-1} dz \overline{\mathbf{S}}.$$

The first integral on the right-hand side of (4.2) vanishes by the Cauchy integral theorem, and since the second is precisely Definition 3.2 of $f(\overline{\mathbf{\Lambda}})$ (scaled by $2\pi i$), we have

$$(4.3) \quad \frac{1}{2\pi i} \int_{\Gamma_2} f(z)(z\mathbf{I} - \mathbf{A})^{-1} dz = \underline{\mathbf{Q}}f(\overline{\mathbf{\Lambda}})\overline{\mathbf{S}}.$$

Combining (4.1) and (4.3) gives the required result. \square

Remark 1. Theorem 4.1 and its proof generalize to the case when \mathbf{A} is not diagonalizable but can be expressed in Jordan canonical form, where λ_1 is semisimple with multiplicity 1. Furthermore, the result can easily be extended to a contour which avoids multiple eigenvalues of \mathbf{A} .

Remark 2. When computing $f(\mathbf{A})b$ the correction $f(\lambda_1)qs^T b$ is a scalar multiple of q . If \mathbf{A} is symmetric, then $s = q$ and the correction can be computed explicitly if q and λ_1 are known.

Remark 3. In the case of Neumann boundary conditions in our FEM discretization, $\lambda_1 = 0$, and the null-space of \mathbf{L} , and hence \mathbf{A} , is a constant vector. Although $\mathbf{A} = \mathbf{M}^{-1}\mathbf{L}$ is not symmetric, both \mathbf{M} and \mathbf{L} are, and so the null-space of \mathbf{A}^T is spanned by $s = \mathbf{M}q/q^T\mathbf{M}q$ (where the scaling ensures $q^T s = 1$). The correction $f(\lambda_1)\mathbf{Q}(1, :)\mathbf{Q}^{-1}(:, 1)b$ when applying Theorem 4.1 to a vector b is therefore given by

$$(4.4) \quad q \frac{f(0)q^T\mathbf{M}b}{q^T\mathbf{M}q} = e \frac{f(0)e^T\mathbf{M}b}{e^T\mathbf{M}e},$$

where e is the constant vector of ones. This correction has a further interpretation in terms of mass conservation. That is, if $f_2(\mathbf{A})b$ is defined by the integral in (4.3), then

$$(4.5) \quad f(\mathbf{A})b = f_2(\mathbf{A})b + \frac{e^T\mathbf{M}(b - f_2(\mathbf{A})b)}{e^T\mathbf{M}e}e$$

ensures that the masses $e^T\mathbf{M}f(\mathbf{A})b$ and $e^T\mathbf{M}b$ are equal (as should be the case when $f(0) = 1$ as in (3.4)). Furthermore, since the vector e is orthogonal to each column of $\mathbf{M}\mathbf{Q} = \mathbf{M}\mathbf{Q} = \overline{\mathbf{S}}^T$, we find $e^T\mathbf{M}f_2(\mathbf{A})b = e^T\mathbf{M}\mathbf{Q}\underline{f}(\underline{\Lambda})\overline{\mathbf{S}}b = 0$, and so (4.5) agrees with (4.4).

The approach of simply ignoring the zero eigenvalue is related to the *spectral splitting* idea introduced by Ilić and Turner for symmetric positive definite matrices [30]. They discuss a method of decomposing $f(\mathbf{A})$ whereby the spectrum $\sigma(\mathbf{A})$ is split into a “regular” part Λ , where f can be well approximated by a low-degree polynomial p_n , and a “singular” part Λ^c , which is less easily approximated. This leads to an approximation of the form

$$(4.6) \quad f(\mathbf{A})b \approx \hat{\mathbf{Q}}f(\mathbf{H})\hat{\mathbf{S}}b + p_n(\mathbf{A})(\mathbf{I} - \hat{\mathbf{Q}}\hat{\mathbf{S}})b,$$

where $\hat{\mathbf{Q}}$ is an orthonormal basis for the invariant subspace corresponding to Λ^c such that $\mathbf{A}\hat{\mathbf{Q}} = \hat{\mathbf{Q}}\mathbf{H}$, and $\hat{\mathbf{S}} = \hat{\mathbf{Q}}^T$. Ilić and Turner suggest that one of the drawbacks of the method is the difficulty in computing the basis $\hat{\mathbf{Q}}$, but they go some way toward addressing this in a subsequent paper [31]. In our case $\mathbf{A} = \mathbf{M}^{-1}\mathbf{L}$ is no longer symmetric, but we can define $\hat{\mathbf{Q}} = (q \text{ zeros}(\mathbb{N}, \mathbb{N} - 1))$ and $\hat{\mathbf{S}} = (\mathbf{M}\hat{\mathbf{Q}})^T$ with $\mathbf{H} = 0$ so that the first term in the sum (4.6) reduces to a familiar $f(0)qs^T$. Some further algebra shows that

$$\begin{aligned} p_n(\mathbf{A})\hat{\mathbf{Q}}\hat{\mathbf{S}}b &= \mathbf{Q}p_n(\Lambda)\mathbf{S}\hat{\mathbf{Q}}\hat{\mathbf{S}}b \\ &= \mathbf{Q}p_n(\Lambda)e_1s^Tb \\ &= p_n(\lambda_1)qs^Tb \end{aligned}$$

and so

$$(4.7) \quad f(\mathbf{A})b \approx p_n(\mathbf{A})b + (f(\lambda_1) - p_n(\lambda_1))qs^Tb.$$

Although in [30] the original motivation was to let p_n be a low-degree polynomial which approximates f well on Λ , there is no need to stick with polynomials. In particular, we can replace $p_n(x)$ in (4.7) with $r_n(x; [\lambda_2, \lambda_N])$, the degree n PAIN approximation assuming that the spectrum of \mathbf{A} is contained in $[\lambda_2, \lambda_N]$. In this way the PAIN method can now also be applied to Neumann-type problems. Similarly, replacing p_n with f_2 from (4.3) (or its CIM approximation), we recover the approach suggested by Theorem 4.1.

For the EKSM, things are not so straightforward. Whereas we could apply the PAIN method to \mathbf{A} even when it is singular (as the only solves needed have negative shifts), the EKSM builds a space containing $(\mathbf{M}^{-1}\mathbf{L})^{-k}b$, which is not well defined if b contains any components in the null-space of \mathbf{L} . However, we propose the following idea, similar to those for the CIM and PAIN methods and related to the well known Sherman–Morrison formula. Let

$$(4.8) \quad \mathbf{B} = \mathbf{A} + \mu q s' = \mathbf{M}^{-1} \left(\mathbf{L} + \mu \frac{\mathbf{M} e e^T \mathbf{M}}{e^T \mathbf{M} e} \right)$$

for some $\mu > 0$; then

$$(4.9) \quad f(\mathbf{A})b = f(\mathbf{B})b + (f(0) - f(\mu))q s^T$$

by (4.1). Now \mathbf{B} is nonsingular and $f(\mathbf{B})b$ can be computed by EKSM. The rank 1 update in (4.8) means that \mathbf{B} is now dense, and a naïve implementation will be impractical. However, noting that $\mathbf{M}\mathbf{B}$ is simply the stiffness matrix \mathbf{L} plus a multiple of the rank 1 matrix $(\mathbf{M}e)(\mathbf{M}e)^T$, this structure can be exploited to solve $\mathbf{B}^{-k}b$ efficiently. In particular, it can be shown that if y is a solution to

$$\mathbf{L}y = \mathbf{M} \left(c - \frac{e^T \mathbf{M} c}{e^T \mathbf{M} e} e \right)$$

(which is well posed as the right-hand side has no component in the kernel of \mathbf{L}), then

$$x = y + \frac{e^T \mathbf{M}}{e^T \mathbf{M} e} \left(\frac{c}{\mu} - y \right) e$$

solves $\mathbf{B}x = c$.

Figure 4.1 and Table 4.1 show results for the same example problem (3.13), but now with Neumann boundary conditions. For CIM and PAIN this is done by applying the relevant boundary conditions to the stiffness matrix \mathbf{L} , with $\mathbf{1} = \mathbf{eigs}(\mathbf{L}, \mathbf{M}, 2, 'SM')$; $\mathbf{1}\mathbf{1} = \mathbf{1}(2)$; $\mathbf{1}\mathbf{N} = \mathbf{eigs}(\mathbf{L}, \mathbf{M}, 1, 'LM')$; in the codes given previously, and adjusting the output \mathbf{v} subject to (4.5) for the CIM or (4.7) for the PAIN method. For EKSM, $f(\mathbf{B})b$ was computed with μ in (4.8) chosen as $\lambda_1/2$ and adjusted subject to (4.9). In solving these Neumann problems we use the MATLAB Parallel Computing Toolbox on a 3 core machine, still with \backslash , and find in Table 4.1 that as well as convergence as a function of the number of solves n , the timings for the CIM and for PAIN are now comparable. This suggests that on multicore or clustered machines the penalty taken by the CIM in using complex arithmetic might be compensated by the better scalability of its independent solves. The EKSM should benefit by a factor of almost two, as the multiplication and solves with \mathbf{A} (corresponding to solves with \mathbf{M} and \mathbf{L} , respectively) can be solved independently, but our attempts to exploit this in the MATLAB code lead to a decrease in speed rather than an increase.

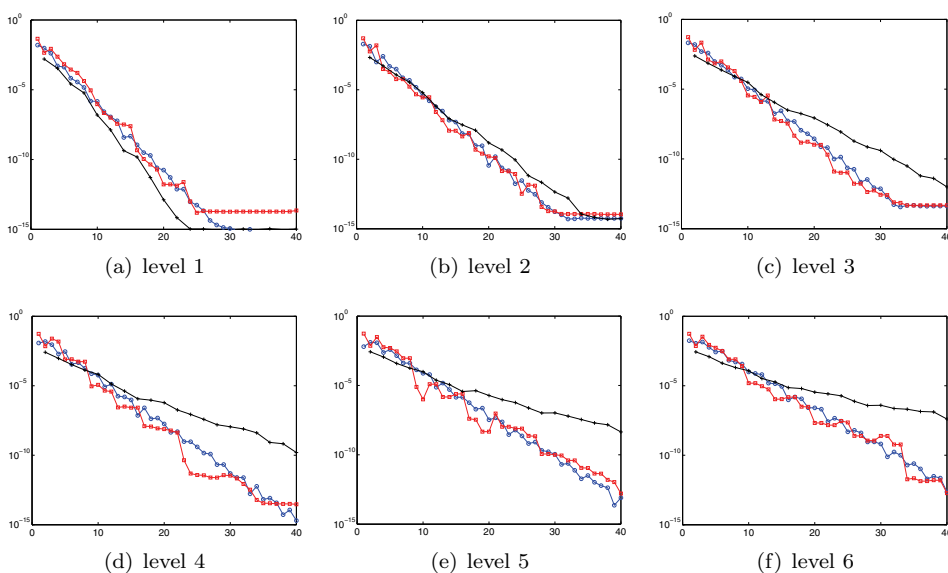


FIG. 4.1. As in Figure 3.3, showing the convergence of the CIM ($\circ-\circ$), EKSM ($+--+$), and PAIN ($\square-\square$) approximations to the $f(\mathbf{A})b$ for the Neumann version of (3.13). Here the less-optimal dependence of the EKSM convergence on the condition number of $\mathbf{M}^{-1}\mathbf{L}$ is more evident as we move to the more refined levels.

TABLE 4.1

Results for solving (3.13), but with homogeneous Neumann boundary conditions. Here we solve on a three core machine using the MATLAB Parallel Computing Toolbox and see that the CIM scales much better than before, and the computational time is now comparable with PAIN. Although the EKSM should theoretically benefit from a factor of 2 speed up in the parallelization of the two independent solves at each iteration, we did not manage to find a way to produce this in practice.

Level	N	Cond. no.	CIM time	PAIN time	EKSM time
1	145	405	0.0445	0.0361	0.0352
2	545	1480	0.1037	0.3985	0.0671
3	2113	5822	0.4077	0.7482	0.3058
4	8321	23301	2.4767	2.3300	1.8584
5	33025	93215	14.779	14.926	9.4105
6	131585	372878	96.589	90.468	54.131

As before, the convergence curves for the CIM and PAIN methods are closely matched. However, the stronger dependence of EKSM on the condition number of \mathbf{A} is more evident in this Neumann case, and furthermore, as there are no longer repeated eigenvalues or gaps in the spectrum close to the origin which it can exploit, the EKSM does not benefit from the instant accuracy we saw in the Dirichlet case. Nonetheless, the simplicity of the linear solves and the trivial parallelization of the two independent solves in the EKSM mean, as we see in Table 4.1, that the same accuracy may be obtained with a comparable computational effort with more iterations.

5. Practical solution methods. In the previous section we provided CPU timings when calculating the fractional Laplacian using the three methods CIM, PAIN, and EKSM. In all cases the majority of this time was taken up solving the required linear systems. Hence, to improve these methods, an efficient solution method for large

sparse systems is required. In the following two sections we will investigate preconditioning choices for both CIM and PAIN. All results presented are for Neumann-type problems, although when applied to Dirichlet-type problems results were seen to be similar. Finally, we recall that there are no shifts in the EKSM, and so standard preconditioners for mass and stiffness matrices can be used.

5.1. Iterative methods. For both the PAIN and CIM methods we are required to solve systems of the form

$$\mathbf{A}_z := (\mathbf{L} - zk^2\mathbf{M})\mathbf{x} = \mathbf{b},$$

where $k \in \mathbb{R}$ and $z \in \mathbb{C}$ with $|z| = 1$. Furthermore, $Re(z), Im(z) > 0$ for the CIM, and z is real and negative for the PAIN method. In both cases these systems are large and very sparse. Using suitably preconditioned Krylov subspace methods may lead to reliable and efficient numerical schemes for such systems.

For \mathbf{A}_z we wish to apply a preconditioner, \mathbf{P}_z , such that

1. \mathbf{P}_z is fast to construct;
2. the action of \mathbf{P}_z^{-1} is fast and requires little storage;
3. the Krylov subspace method acting on $\mathbf{P}_z^{-1}\mathbf{A}_z$ provides mesh independent convergence rates.

5.2. Preconditioning CIM. Due to the complex shifts required in the CIM we are left with an indefinite system. In this case BiCGstab [62] is an appropriate iterative solver to consider. In Table 5.1 we see that applying either an incomplete LU factorization with no fill in, $ILU0(\mathbf{A}_z)$, or an algebraic multigrid, $AMG(\mathbf{A}_z)$, directly to \mathbf{A}_z leads to, at best, mesh dependent convergence and, at worst, total failure. For the incomplete LU preconditioner this degradation with respect to mesh refinement is due to the fact that a high proportion of the shifts result in diffusion-dominated systems, where it is well known that incomplete factorization performs poorly [63]. In the AMG case the action of the inverse of the complex indefinite systems is not approximated well and results in no convergence [19]. Finally, when using $\mathbf{P} = \mathbf{L}_0$, where \mathbf{L}_0 is \mathbf{L} with one node pinned to remove the singularity, motivated by the fact that it is well known that $AMG(\mathbf{L}_0)$ provides an optimal approximation of \mathbf{L}_0^{-1} , iteration counts increase dramatically with mesh refinement [5].

We investigate the use of four possible preconditioners: $\mathbf{P}_1 := \mathbf{L} + k^2z_1\mathbf{M}$, $\mathbf{P}_2 := \mathbf{L} + ik^2z_1\mathbf{M}$, $\mathbf{P}_3 := \mathbf{L} + (1 - 0.5i)z_1k^2\mathbf{M}$, and $\mathbf{P}_4 := \mathbf{L} - k^2(z_1 - 1)\mathbf{M}$, where $z = z_1 + iz_2$. Note, \mathbf{P}_1 was originally considered by Laird and Giles [34], while \mathbf{P}_2 and \mathbf{P}_3 were considered by Erlangga and colleagues [18, 19].

Assuming that w is not an eigenvalue of the generalized eigenvalue problem

$$\mathbf{L}\mathbf{x} = \lambda\mathbf{M}\mathbf{x}, \quad \lambda \in \mathbb{R}^+,$$

any eigenvalue $\sigma := \sigma_1 + i\sigma_2$ of the preconditioned system, $\mathbf{P}_j^{-1}\mathbf{A}_z$, satisfies

$$(5.1) \quad w_2\sigma_1^2 - (z_2 + w_2)\sigma_1 + w_2\sigma_2^2 + (z_1 - w_1)\sigma_2 + z_2 = 0,$$

TABLE 5.1

Average and maximum, in parentheses, number of BiCGStab iterations per quadrature point for the z_i shifted Laplacian solves required in the CIM(20). Here (-) denotes failure to converge. Levels correspond to Table 3.1.

	Level 2	Level 3	Level 4	Level 5
$ILU0(\mathbf{A}_z)$	13 (30)	16(50)	21(80)	29(142)
$AMG(\mathbf{A}_z)$	19 (70)	(-)	(-)	(-)
$AMG(\mathbf{L}_0)$	60 (155)	100 (288)	180 (570)	(-)

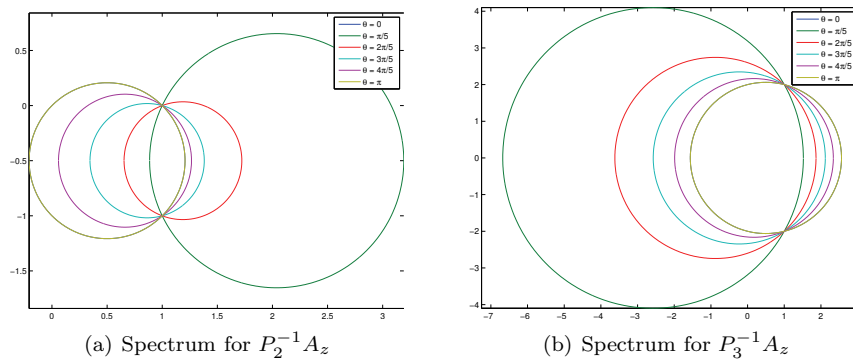


FIG. 5.1. Spectrum of preconditioned system for various positions on the CIM shift circle.

where $\mathbf{P}_j = \mathbf{L} - (w_1 + iw_2)\mathbf{M}$. First, for the preconditioning choices \mathbf{P}_1 and \mathbf{P}_4 , where $w_2 = 0$, we see that the spectrum of the preconditioned system lies on the line in the complex plane satisfying

$$-z_2\sigma_1 + (z_1 - w_1)\sigma_2 + z_2 = 0.$$

Second, for the preconditioners \mathbf{P}_2 and \mathbf{P}_3 , where $w_2 \neq 0$, it can be shown that σ lies on a circle in the complex plain with center C and radius R given by

$$(5.2) \quad C = \left(\frac{w_2 + z_2}{2w_2}, \frac{w_1 - z_1}{2w_2} \right) \quad \text{and} \quad R = \sqrt{\frac{(w_2 - z_2)^2 + (w_1 - z_1)^2}{4w_2^2}}.$$

(See Van Gijzen, Erlangga, and Vuik [20] for details.)

In particular, in the CIM we use a modification of `method1.m` as described in [24], which gives rise to a Γ which is precisely a circle of radius $\lambda_N \sqrt{1 - \lambda_1/\lambda_N}$ about the point λ_N (Figure 3.1). From here on we will call this the CIM shift circle. Writing the shift z in the form

$$z(\theta) = (\lambda_N + \lambda_N \sqrt{1 - \lambda_1/\lambda_N} \cos(\theta)) + i(\lambda_N \sqrt{1 - \lambda_1/\lambda_N} \sin(\theta)),$$

we see that the preconditioned system remains bounded independently of the magnitude of the shift. Furthermore, when considering \mathbf{P}_2 we see that as $\theta \rightarrow \pi$ the CIM shift circle approaches the origin. Figure 5.1 shows the circles that the preconditioned systems lie on for several positions on the CIM shift circles. In practice, $\theta \rightarrow \pi$ as $\lambda_N \rightarrow \infty$, or the number of CIM quadrature points is increased. The consequence can be seen in Table 5.2, where the maximum iteration counts for \mathbf{P}_2 increase as the mesh is refined, where this refinement has led to an increase in λ_N .

The first four rows of Table 5.2 show iteration counts when using exact inverses, and the final four rows show iteration counts when we apply just one AMG V-cycle to the preconditioners. From this table we see that \mathbf{P}_2 is not only mesh dependent when used exactly but also suffers when used inexactly via AMG. Furthermore, it can clearly be seen that \mathbf{P}_1 , \mathbf{P}_3 , and \mathbf{P}_4 perform well when used exactly and, more importantly, inexactly, where only a few more iterations are required compared to the exact solves. In Table 5.3 we show the effects of improving the CIM accuracy by increasing the number of quadrature points in the numerical contour integration on the preconditioners. On average, \mathbf{P}_4 requires the least number of iterations per

TABLE 5.2

Average and maximum, in parentheses, number of BiCGStab iterations for various preconditioners for the z_i shifted Laplacian solve required in the 10-point CIM quadrature discretization, CIM(10). Levels correspond to Table 3.1.

	Level 2	Level 3	Level 4	Level 5
\mathbf{P}_1	17 (23)	18 (24)	17 (23)	18 (24)
\mathbf{P}_2	28 (48)	31 (50)	33 (56)	38 (66)
\mathbf{P}_3	15 (18)	15 (17)	15 (16)	14 (16)
\mathbf{P}_4	12 (16)	12 (14)	12 (14)	11 (13)
AMG(\mathbf{P}_1)	18 (22)	18 (22)	18 (22)	18 (23)
AMG(\mathbf{P}_2)	35 (70)	40 (99)	96 (567)	61 (133)
AMG(\mathbf{P}_3)	16 (19)	16 (19)	15 (18)	15 (17)
AMG(\mathbf{P}_4)	13 (16)	13 (16)	12 (16)	12 (15)

TABLE 5.3

Average and maximum, in parentheses, number of BiCGStab iterations for various preconditioners for the z_i shifted Laplacian solve required in CIM with various numbers of quadrature points.

		Level 2	Level 3	Level 4	Level 5
CIM(20)	AMG(\mathbf{P}_1)	17(21)	18 (22)	19 (24)	19 (24)
	AMG(\mathbf{P}_3)	19(24)	19 (23)	19(22)	18 (22)
	AMG(\mathbf{P}_4)	13 (23)	13 (22)	13 (21)	12 (20)
CIM(40)	AMG(\mathbf{P}_1)	17 (21)	18 (23)	19 (24)	19 (23)
	AMG(\mathbf{P}_3)	23 (31)	24 (30)	23 (30)	23 (30)
	AMG(\mathbf{P}_4)	13 (36)	13 (40)	13 (39)	13 (38)

TABLE 5.4

Average and maximum, in parentheses, number of PCG iterations for AMG(\mathbf{A}_i), ILU0(\mathbf{A}_i), and HYBRID(\mathbf{A}_i) preconditioners for the z_i shifted Laplacian solve required when using PAIN(n), $n = 10, 20, 40$.

		Level 2	Level 3	Level 4	Level 5
AMG(\mathbf{A}_i)	PAIN(10)	8 (23)	9 (26)	9 (25)	10 (35)
	PAIN(20)	–	–	–	–
	PAIN(40)	–	–	–	–
ILU0(\mathbf{A}_i)	PAIN(10)	10 (21)	12 (29)	15 (40)	19 (57)
	PAIN(20)	8 (22)	9 (31)	12 (46)	19 (66)
	PAIN(40)	9 (22)	12 (33)	15 (48)	19 (71)
HYBRID(\mathbf{A}_i)	PAIN(10)	6(10)	7(8)	7(8)	7 (8)
	PAIN(20)	6 (9)	6(8)	7 (8)	7 (8)
	PAIN(40)	6 (8)	6 (8)	7(8)	7 (8)

fractional calculation, although the maximum number of iterations is larger than that of \mathbf{P}_1 . In all cases the maximum iteration count occurs for the largest shift and in this case we could replace the preconditioner in \mathbf{P}_4 with that of \mathbf{P}_1 , although no significant improvement will be obtained.

5.3. Preconditioning the PAIN method. The solves required within the PAIN method are of the form

$$\mathbf{A}_z := (\mathbf{L} - zk^2\mathbf{M})\mathbf{x} = \mathbf{b},$$

where z are real negative. Hence, the system is symmetric positive definite and preconditioned conjugate gradients (PCG) is the iterative method of choice. In Table 5.4 we see that both algebraic AMG and ILU0 lead to unsatisfactory methods. In the case of ILU0(\mathbf{A}_i) we see strong mesh dependence on the iteration counts, while in the AMG case the method fails to converge. In the case of AMG(\mathbf{A}_i) this failure is due to

the large shifts required within the PAIN method, as either PAIN iterations increase or mesh size decreases. While ILU0 is not a reliable method for small shifts where it exhibits mesh dependent convergence [63], if we combine these two preconditioners into a hybrid scheme using AMG for shifts z less than the spectral radius of \mathbf{LM}^{-1} and ILU0 otherwise, we may obtain a reliable mesh and iteration number independent solver. This can be seen in the bottom block of Table 5.4.

6. Numerical results. We have given a thorough discussion and analysis on scalable numerical methods for computing a matrix function with a fractional power, and in this section we apply these to simulate some classes of fractional reaction-diffusion equations. We consider two types of reaction terms,

$$g_1(u) := u(1 - u) \quad \text{and} \quad g_2(u) := u(u - 0.5)(1 - u),$$

for the equation

$$\begin{aligned} \frac{\partial u}{\partial t} + \nu(-\Delta)^\alpha u &= g_i(u) \quad \text{in } \Omega, \\ \frac{\partial u}{\partial n} &= 0 \quad \text{on } \partial\Omega, \\ u(x, 0) &= u_0(x) \quad \forall x \text{ in } \Omega, \end{aligned}$$

where $\nu > 0$. The function $g_1(u)$ represents the Fisher reaction equation leading to logistic growth, with $u = 0$ being an unstable equilibrium point. The second function $g_2(u)$ represents an Allen–Cahn equation with a quartic double well potential, which results in two stable modes with motion driven by curvature.

Within each backward Euler time step $[t_n, t_{n+1}]$ we will treat the nonlinear term, $g_i(u^{n+1})$, using the following simple fixed point iteration: Given u^n , define $u^{n+1,0} := u^n$, and for $k = 1, 2, \dots, N$ find $u^{n+1,k+1}$ such that

$$\frac{u^{n+1,k} - u^n}{\Delta t} + \nu(-\Delta)^\alpha u^{n+1,k} = g_i(u^{n+1,k-1}),$$

where N is to be chosen. Clearly, $N = 1$ leads to the fully explicit treatment of the nonlinear term, and for sufficiently large N the method is fully implicit.

Throughout the results section we terminate the above iteration when

$$\frac{\|u^{n+1,k} - u^{n+1,k-1}\|}{\|u^n\|} < 10^{-4}.$$

In practice this results in three to four iterations at every time step.

In all the results that follow we apply CIM(20). From the previous section we have seen that the use of preconditioner AMG(\mathbf{P}_4), using just one V-cycle, within BiCGStab leads to an effective numerical solution. For both the following two- and three-dimensional results we apply this preconditioner. Table 6.1 shows average CPU timings for three typical time steps applied to a two-dimensional fractional Allen–Cahn problem with $\alpha = 0.85$. The table clearly shows excellent scaling of the model with respect to mesh refinement. Furthermore, these results are indicative of the timings for all subsequent problems.

TABLE 6.1
Average CPU timings per iteration.

DOFs	145	545	2113	8321	33025	131585
$\Delta t = 0.05$	0.43	0.63	1.27	7.54	28.9	126.5
$\Delta t = 0.1$	0.48	0.67	1.27	7.74	31.25	145.0
$\Delta t = 0.2$	0.51	0.78	1.42	8.64	33.25	158.0

Finally, time steps are typically in the region $0.001 < \Delta t < 0.1$. In the case of spinodal decomposition, initially the solution undergoes a rapid transformation into bulk regions. The motion is then seen to slow dramatically. Hence, we use an initial small time step $\Delta t = 10^{-5}$ and increase this to the value $\Delta t = 0.1$ at a rate proportional to $t^{1/4}$ (see [32]).

6.1. Fisher equation. When $\alpha = 1$, the Fisher reaction produces a traveling wave with a smooth interface between the $u = 1$ stable state and the $u = 0$ unstable state. The thickness of this interface and the speed of the wave are determined by ν . For the Allen–Cahn reaction the interface thickness between the two stable states $u = 0$ and $u = 1$ and the speed of the curvature driven flow are determined by ν . For $0 < \alpha < 1$, Engler [17] has shown, in the case $\Omega = \mathbb{R}$, that for Fisher-type reactions with $g > 0$ on $(0, 1)$, the speed of spread of the wave front is unbounded. On the other hand, for Allen–Cahn type reactions, with $g'(0) > 0$ and $g \leq 0$ in a neighborhood of 0, the motion of any interfaces remains finite. In the following we will produce numerical results for both types of models. These results are consistent with the bounds given by Engler [17].

Here we define $\Omega_1 = (0, 20) \times (0, 0.1)$, $u_0(x, y) = e^{-5x}$, and $\nu = 10^{-3}$. This domain is chosen so as to accommodate the resulting unbounded speed of spread of the traveling front. The meshes used within the following two-dimensional example have between 40,000 and 140,000 degrees of freedom. Figure 6.1 shows several evenly spaced time slices for the evolution of the model for $\alpha = 1$ and $\alpha = 0.98$ through the line $y = 0$. In Figure 6.1(a), the steady speed of the standard Fisher equation with $\alpha = 1$ is visible. Figure 6.1(b) shows the effects of fractional diffusion, in this case, $\alpha = 0.98$. The exponential spread of the interface, derived by Engler [17], is apparent.

6.2. Allen–Cahn equation—interfacial properties. In this first example we are interested in the effects on the interface when fractional diffusion is used for Allen–Cahn reactions. To enable us to focus on this issue we consider the long thin domain Ω_1 and use the initial condition $u_0(x, y) = \max(\text{sign}(10 - y), 0)$. This leads to an interface centered along the line $x = 10$, the thickness of which is a function of ν . Since it is only within a region of this interface that the solution changes rapidly, we use a highly nonuniform mesh, with the smallest elements around the region $5 - 1/\nu < x < 5 + 1/\nu$. Figures 6.2(a), 6.2(b), and 6.2(c) show the effects of α on the profile of the steady state solution for fixed $\nu = 0.01$. As expected, in the case $\alpha = 1$, we obtain a well-defined interface with the profile attaining the extreme values of $u = 1$ and $u = 0$ around $x = 8$ and $x = 12$, respectively. For the cases where $\alpha < 1$, we see that the solution changes significantly faster near the center of the interface with respect to the $\alpha = 1$ case. Away from the center the solutions become less steep, and as α is reduced the whole interface becomes thicker.

Figures 6.3(a) and 6.3(b) show three choices for $\nu = 0.1, 0.01, 0.001$ with standard $\alpha = 1$ diffusion compared to the choice $\alpha = 0.5, \nu = 0.01$. As ν decreases, Figure 6.3(a) shows how the interface becomes thinner and steeper at the center. From Figure 6.3(b), which zooms in on the interface region, it is clear that the tail

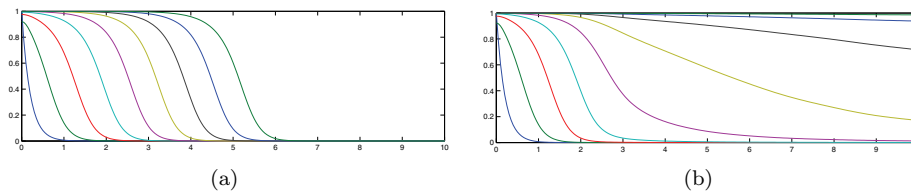


FIG. 6.1. Fisher reaction for standard diffusion, $\alpha = 1$ (left), and fractional diffusion, $\alpha = 0.98$ (right).

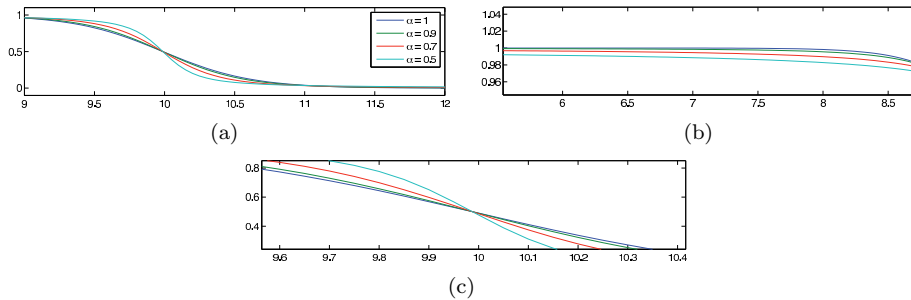


FIG. 6.2. Allen-Cahn reactions for various powers of diffusion, $\nu = 0.01$.

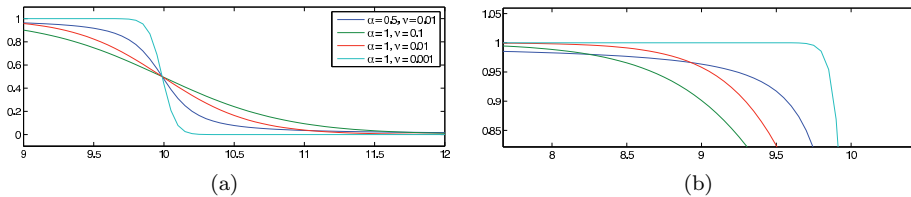


FIG. 6.3. Allen-Cahn for various ν with $\alpha = 1$, compared to $\alpha = 0.5$ with $\nu = 0.01$.

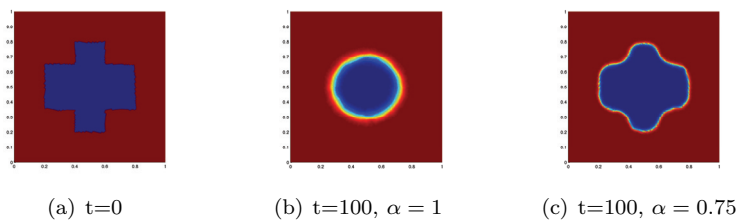


FIG. 6.4. Allen-Cahn diffusion starting from an initial cross (a) and at time $t = 100$ with diffusion power $\alpha = 1$ and $\alpha = 0.75$, respectively ((b) and (c)).

of the interface behaves very differently when fractional diffusion is considered. All solutions obtained using standard diffusion pass the fractional wave at some stage, due to the long tail in the fractional solution.

6.3. Allen-Cahn equation—Motion from an initial cross. In this model we define $\Omega = (0, 1) \times (0, 1)$, $u_0(x, y)$ such that the $u = 0$ phase is in the shape of a cross (Figure 6.4(a)), and $\nu = 10^{-3}$. With standard diffusion, i.e., $\alpha = 1$, we see

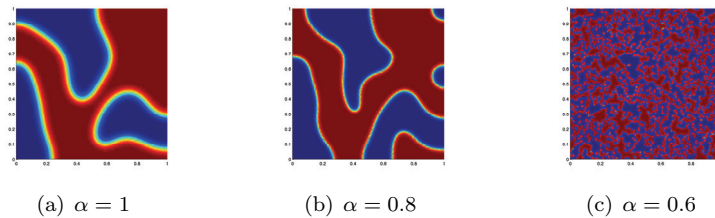


FIG. 6.5. *Spinodal decomposition starting from a random mixture. Plots from left to right show the state at time $t = 100$ for $\alpha = 1.0, 0.8, 0.6$.*

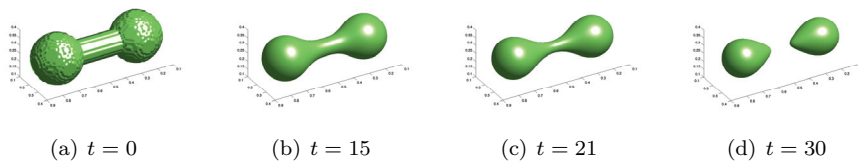


FIG. 6.6. *Level lines $u = 0.5$ for $\alpha = 1$ at times 0.4, 15, 21, and 30.*

in Figure 6.4(b) that the curvature drives the $u = 0$ phase toward a circle (constant curvature) and shrinks. The motion for fractional diffusion, with $\alpha = 0.75$, is similar, although the rate is slower and the interface is thinner (Figure 6.4(c)).

6.4. Allen–Cahn equation—Spinodal decomposition. In this final two-dimensional example we investigate the effects of fractional diffusion when spinodal decomposition is considered. The initial state is well mixed, with $u_0(x, y)$ drawn from a random normal distribution about 0.5. For $\alpha = 1$ the early stages of phase transition produce a rapid movement to bulk regions of both phases, and then motion slows resulting in the state given in Figure 6.5(a) at time $t = 100$. Reducing the fractional power leads to thinner interfaces that allow for smaller bulk regions and a much more heterogeneous phase structure. Furthermore, motion to large bulk regions is dramatically slowed for fractional models; see Figures 6.5(b) and 6.5(c).

6.5. Allen–Cahn equation in three-dimensional space. We now compare standard diffusion with fractional diffusion in three spatial dimensions. Due to the significantly large number of degrees of freedom required to obtain accurate approximations in three dimensions, the methods considered in this paper are essential when considering fractional models. The following two examples are of fractional Allen–Cahn models with diffusion coefficient $\nu = 0.02$. The approximation is on a structured cuboidal mesh with 262,701 degrees of freedom.

For the first example the initial condition is given by $u(x, y) = 0$ if $(x, y) \in D$ and 1 otherwise, where D is the dumbbell occupying the region given in Figure 6.6(a) and $\Omega = (0, 1) \times (0, 0.5) \times (0, 0.5)$. We take the time slices in Figures 6.6 and 6.7 and show the level line $u = 0.5$ for standard diffusion and fractional diffusion with $\alpha = 0.8$ and $\alpha = 1$, respectively. From these we see that the motion is slowed when a fractional power is considered. We also observe that, as was the case in two dimensions, the interfacial region $0 < u < 1$ in the fractional case is sharper.

In the second example we look at spinodal decomposition. Figures 6.8 and 6.9 show characteristics similar to those seen in the two-dimensional case. In particular, for $\alpha = 0.8$, the bulk regions persist longer when compared to $\alpha = 1.0$, where at time

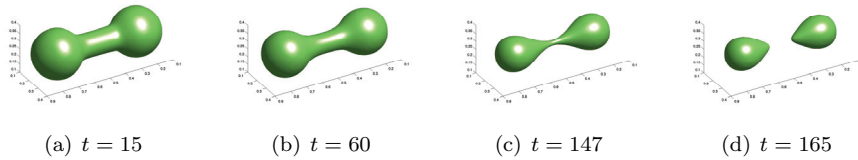


FIG. 6.7. Level lines $u = 0.5$ for $\alpha = 0.8$ at times 15, 60, 147, and 165.

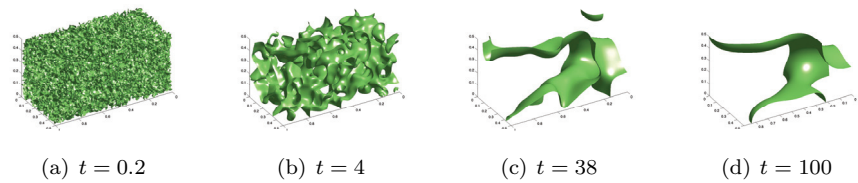


FIG. 6.8. Level lines $u = 0.5$ for $\alpha = 1$ at times 0.4, 4, 38, and 100.

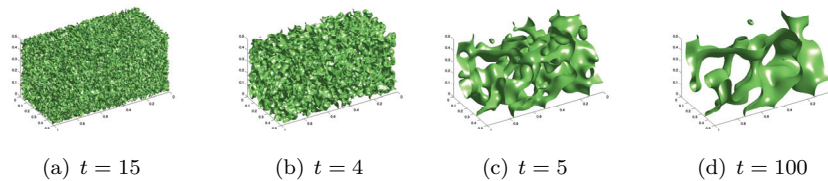


FIG. 6.9. Level lines $u = 0.5$ for $\alpha = 0.8$ at times 0.4, 4, 38, and 100.

$t = 100$ the $\alpha = 1$ case has only a couple of bulk regions, while the fractional model has numerous bulk regions.

7. Conclusion. In this paper we have developed robust, efficient, and scalable techniques for solving fractional-in-space reaction-diffusion equations using finite elements on unstructured grids. We have considered three approaches: the contour integral method (CIM), the poles and interpolation nodes (PAIN) method, and the extended Krylov subspace method (EKSM). CIM is trivially parallelizable through a set of shifted linear solves, and effective preconditioners can be found that lead to mesh independent convergence. The effects of inexact solves are well understood, but a negative aspect is that complex arithmetic is required. In the case of the PAIN method all computations are real and there are no issues that arise from using effective iteration schemes, but the method is not so easily parallelizable and the effect of inexact solves is less well understood. Finally, in the case of EKSM, no shifted solves are required, it is trivially parallelizable on two cores only, the convergence appears strongly mesh independent, and the effect of inexact solves is unknown at this stage. Simulation results on the fractional-in-space Fisher and Allen–Cahn equations show that such problems can have very different dynamics to standard diffusion models and as such represent a powerful modeling approach for understanding the many aspects of heterogeneity.

This paper has used the framework of fractional-in-space reaction-diffusion equations as an aid to understanding the effects of spatial heterogeneity, but clearly homogeneity can be much more complicated than the ideas expressed here. Issues that we have not covered, but which would be needed to address in order to use nonlocal models as a mechanism for understanding heterogeneity in depth, include considering the more general framework of nonlocal operators, anisotropy, spatially dependent diffusion tensors, spatially dependent fractional powers, and more complex boundary conditions. In many of these cases it would be more natural to consider numerical approximations to spatial Riesz derivatives in the various coordinate components, as this approach can cope with these more subtle generalizations. However, the ideas expressed in this paper would still have applicability in this setting. These aspects are beyond the scope of this paper but we note that all of these are active research areas of many groups around the world, including ours, and will be the focus of future work.

Appendix.

```

0 %LEJADC Leja-Bagby points from two discrete sets. Adapted from [25]
1 function [a, b] = lejadc(A, B)
2 sA = ones(size(A)); sB = sA; % Initialize sum vectors
3 a = sA; a(1) = max(A);      % Initialize a and b
4 b = sB; b(1) = min(B);      % Initialize b
5 for j = 1:length(A)-1      % Loop over the discrete sets
6   sA = sA .* ((A-a(j)) ./ (A-b(j))); % Update sum A
7   sB = sB .* ((B-a(j)) ./ (B-b(j))); % Update Sum B
8   [~,indA] = max(abs(sA)); [~,indB] = min(abs(sB)); % Find min & max
9   a(j+1) = A(indA);      b(j+1) = B(indB); % Update outputs
10 end

```

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