

# On the performance of an implicit-explicit Runge–Kutta method in models of cardiac electrical activity

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**Abstract**—Mathematical models of electric activity in cardiac tissue are becoming increasingly powerful tools in the study of cardiac arrhythmias. Considered here are mathematical models based on ordinary differential equations (ODEs) that describe the ionic currents at the myocardial cell level. Generating an efficient numerical solution of these ODEs is a challenging task, and in fact the physiological accuracy of tissue-scale models is often limited by the efficiency of the numerical solution process. In this paper, we examine the efficiency of the numerical solution of 4 cardiac electrophysiological models using implicit-explicit Runge–Kutta (IMEX-RK) splitting methods. We find that variable step-size implementations of IMEX-RK methods (ARK3 and ARK5) that take advantage of Jacobian structure clearly outperform the methods commonly used in practice.

**Keywords:** splitting methods; efficient numerical methods; implicit-explicit Runge–Kutta methods; simulation of electrophysiological models; Rush–Larsen method; ordinary differential equations.

## I. INTRODUCTION

Computer simulation is becoming an important tool in cardiovascular research. Mathematical models of the heart can be used to simulate heart conditions and the effects of certain drugs designed to treat them. Presently, the development of a drug often costs hundreds of millions of dollars [1]. One aim of computer simulation is to reduce this cost, e.g., by reducing the number of physical experiments needed in designing a drug.

Electrophysiological models of the heart describe how electricity flows through the heart, controlling its contraction. In this paper, we consider 4 mathematical models for the flow of ionic currents at the myocardial cell level. The models consist of systems of ordinary differential equations (ODEs). Cardiac electrophysiological

models are often based on the Nobel prize-winning work of Hodgkin and Huxley [2] in the 1950s that modelled neural tissue mathematically as a circuit. Modern cardiac electrophysiological models adapt the work of Hodgkin and Huxley to describe electrical activity in the heart and include data gathered from experiments to form models with increasing physiological accuracy.

A major barrier to obtaining the most useful data from tissue-scale electrophysiological models of the heart is the challenge of performing the simulations efficiently. Often the physiological accuracy of the mathematical model must be sacrificed for a simulation to become feasible; see, e.g., [3], [4], [5], [6]. The ODE systems describing the cellular dynamics in these models are non-linear and stiff (see, e.g., [7]). The consequence of stiffness is that the speed of the solution process is limited by considerations of numerical stability instead of accuracy. Hence the solution process can potentially be made more efficient through the use of appropriate numerical algorithms.

In this paper, we investigate the efficiency of implicit-explicit Runge–Kutta (IMEX-RK) splitting methods (see, e.g., [8]) for the simulation of 4 cardiac electrophysiological models. Cardiac electrophysiological models contain linear and non-linear terms as well as stiff and non-stiff terms, so IMEX methods are a natural choice for their numerical solution. An IMEX-RK method uses both an implicit RK (IRK) method and an explicit RK (ERK) method to approximate the solution to an ODE.

The use of splitting methods for cardiac simulations is quite common, particularly for simulations in which the ODEs for myocardial cell models are coupled with partial differential equations (PDEs) describing the flow of electricity through myocardial tissue. For example, a second-order accurate operator splitting method for the monodomain model of myocardial tissue is studied in [9]. The coupled non-linear PDE model is split into an uncoupled linear parabolic PDE and a system of non-linear ODEs. The PDE is then solved with either the forward Euler method or an alternating direction implicit method, and the ODE system is solved with the Rush–

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Larsen method [10]; see also below. This splitting is also used with first- and second-order IMEX schemes in [11] for both the monodomain and bidomain models. An implicit method is used for the PDE(s), and an explicit method is used to integrate the ODEs. The use of both a second-order Crank–Nicolson method and a first-order semi-implicit method to solve the bidomain model is investigated in [12]. These methods were optimized by splitting the non-linear cell model into two components, one with positive eigenvalues and another with negative eigenvalues. This work is extended in [13], where the cell model is split from the PDEs and solved with the backward Euler method and further optimized by deriving an explicit scheme to solve the ODEs. The work in [12] is extended in [14] to create a first-order method to simulate the flow of electricity in the heart and the surrounding torso. In [15] the ODEs are separated as in [9], and then the approach from [14] is used to solve the PDEs in the heart and torso, resulting in a second-order method. The work of [12] is also extended in [16] to include additional complexities such as rotational anisotropy and blood layer boundary conditions. Other numerical methods used to solve the ODEs in such models include singly diagonally implicit RK (SDIRK) methods, fully implicit RK methods, and multi-step methods based on backward differentiation formulas (BDFs); see below and also, e.g., [17].

In this paper, we consider 4 mathematical models of cardiac electrophysiology: the Luo–Rudy model of guinea pig ventricular tissue [18], the Courtemanche et al. model of human atrial tissue [19], the Winslow et al. model of canine ventricular tissue [20], and the Puglisi–Bers model of rabbit ventricular tissue [21]. We perform a thorough comparison of the performance of two specific IMEX-RK methods, denoted here as ARK3 and ARK5 [22], respectively, to numerical methods commonly used in practice for simulating these 4 models. We also consider problem-specific optimizations to the IMEX-RK methods described. The results presented have the obvious limitation that they are based on single-cell models, and thus further studies are necessary to fully establish their applicability to tissue-scale simulations. However, the present work can provide guidance for development of efficient splitting methods for the simulation of PDE models that use variable step-size ODE solvers for the cellular dynamics.

The rest of this paper is organized as follows. In Section II we give a brief introduction to mathematical models of electrical activity in the heart. In Section III we give a brief overview of the numerical methods used in this paper for the numerical solution of the ODEs arising from these models. In Section IV we discuss the results

of the numerical experiments. We find that a variable step-size implementation of an IMEX-RK method with a customized linear system solver clearly outperforms all the commonly used numerical methods of which we are aware. Finally in Section V we give some conclusions and directions for future research.

## II. ELECTRICAL ACTIVITY OF THE HEART

Because of their intricacy, obtaining physiologically accurate mathematical models is a difficult task. A further challenge to obtaining physiological accuracy is that of performing the simulation efficiently. To move effectively beyond models for one cell, enough cells must be included in the model to realistically approximate the geometry and physiology of the heart. Because the heart has approximately  $10^{10}$  cells [23], any realistic simulation will have enough cells (or clusters of cells) to dramatically magnify any inefficiencies in the numerical method. This has forced some researchers to reduce the physiological accuracy of their models to allow the simulation to be performed within an acceptable amount of time; see e.g., [3], [4], [5], [6]. The models are numerically *stiff*, and so standard (explicit) numerical methods are often unable to provide efficient simulations. If the efficiency of the simulation process can be significantly improved, then greater physiological accuracy and subsequently more useful data can be obtained.

### A. The Luo–Rudy model of guinea pig ventricular tissue

In 1991 Luo and Rudy developed a model of guinea pig ventricular action potentials based on a previous model from Beeler and Reuter [24]. The Luo–Rudy model [18] extended the Beeler–Reuter model to include fast inward sodium and outward potassium currents to make the model more physiologically accurate. The general approach of these models is based on Hodgkin–Huxley type formalism [2]; the Luo–Rudy model itself consists of 8 nonlinear ODEs.

For an individual cardiac cell we have that the transmembrane potential  $V_m$  satisfies [18]:

$$\frac{dV_m}{dt} = -\frac{1}{C_m}(I_{ion} + I_{st}), \quad (1)$$

where  $C_m$  is the membrane capacitance,  $I_{ion}$  is the total transmembrane ionic current, and  $I_{st}$  is the stimulus current. An example of the evolution of  $V_m$  over time for this model is given in Figure 1.

The Luo–Rudy model contains 6 ionic currents that are determined by 6 gating variables [18]. The evolution of each gating variable  $y$  is governed by a nonlinear ODE

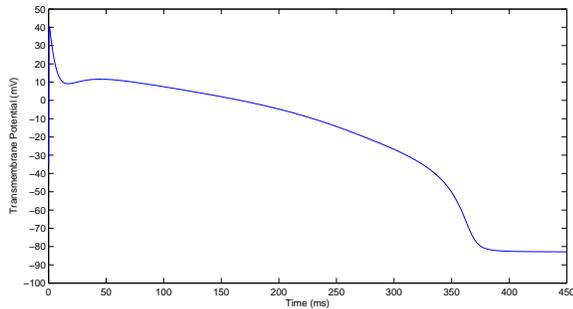


Fig. 1. Transmembrane potential over time in the Luo–Rudy model. This is a numerical solution produced by ARK5 for equation (1) coupled with 7 other ODEs [18] via the  $I_{ion}$  current as produced by ARK5.

involving rate parameters  $\alpha_y$  and  $\beta_y$  in the general form

$$\frac{dy}{dt} = \frac{y_\infty - y}{\tau_y}, \quad (2)$$

where

$$y_\infty = \frac{\alpha_y}{\alpha_y + \beta_y} \quad \text{and} \quad \tau_y = \frac{1}{\alpha_y + \beta_y}.$$

The remaining ODE in the Luo–Rudy model describes calcium concentration in the cell:

$$\frac{d([Ca]_i)}{dt} = -10^{-4}I_{si} + 0.07(10^{-4} - [Ca]_i), \quad (3)$$

where  $[Ca]_i$  is the intracellular calcium concentration and  $I_{si}$  is the slow inward calcium current [18]. The 6 gating equations of the form (2) are coupled with (1) and (3) to form the complete Luo–Rudy model. Full details of the model can be found in [18].

In 1994 Luo and Rudy published an improvement to this model, now known as the Luo–Rudy Phase II model [25], [26]. This model includes the actions of ionic pumps and changes in ionic concentrations. It is a more physiologically accurate yet more complicated model than (1)–(3), consisting of 14 ODEs. We do not consider the Luo–Rudy Phase II model in this study.

### B. The Courtemanche et al. model of atrial tissue

In 1998 Courtemanche, Ramirez, and Nattel developed a model of human atrial action potentials [19]. It was developed in response to findings that show there are important differences in human action potentials when compared to those of other mammals frequently used in models. Courtemanche et al. developed this model with human data supplemented with animal data when needed. The Courtemanche et al. model is an extension of the Luo–Rudy Phase II model. It consists of 21 ODEs. Full details of the model can be found in [19].

### C. The Winslow et al. model of canine ventricular tissue

In 1999 Winslow, Rice, Jafri, Marbán, and O’Rourke developed a model of canine ventricular tissue [20]. This model is based on a guinea pig model that was an extension of the Luo–Rudy Phase II model. The Winslow et al. model was developed using experimental data to modify the guinea pig model so that it would simulate canine ventricular tissue. The Winslow et al. model is particularly detailed when describing the dynamics of  $Ca^{2+}$ , which is an important consideration in heart failure. It consists of 32 ODEs, making it the most complex of the models in this study. Full details of the model can be found in [20].

### D. The Puglisi–Bers model of rabbit ventricular tissue

In 2001 Puglisi and Bers developed a model of rabbit ventricular tissue [21]. Although rabbit ventricular tissue is used frequently in experiment, no mathematical model had been previously developed for it. This model was adapted from the Luo–Rudy model to include data from the literature and from the joint laboratory of Puglisi and Bers. This model was designed to be a learning aid for students as well as a tool for researchers to reproduce experimental data via computer simulation. Thus, physiological accuracy was of paramount importance. The Puglisi–Bers model gives particular detail to calcium handling in order to accurately simulate heart failure. This model contains 17 ODEs. It is also referred to as the *LabHeart* [21] model. Full details of the model can be found in [21].

## III. NUMERICAL METHODS

In this paper we focus on the solution of the *initial-value problem* (IVP) for ODEs, defined as

$$\frac{dy}{dt} = \mathbf{f}(t, \mathbf{y}), \quad \mathbf{y}(0) = \mathbf{y}_0, \quad (4)$$

on the interval  $t \in [0, t_f]$ . We now describe the numerical methods compared here. We also describe aspects of creating an efficient numerical simulation. Except for the fully implicit methods studied, all numerical experiments were run using `odeToJava` [27], a Java-based problem-solving environment for IVPs<sup>1</sup>. Details on how the fully implicit methods were run are given below.

### A. The Forward Euler Method

Arguably the simplest numerical method for approximating the solution of (4) is *Euler’s method*, or *Forward*

<sup>1</sup>Presently, `odeToJava` supports only ERK and linearly implicit IMEX-RK methods.

*Euler* (FE). One step of FE from  $(t_{n-1}, \mathbf{y}_{n-1})$  to  $(t_n, \mathbf{y}_n)$  is given by

$$\begin{aligned}\mathbf{y}_n &= \mathbf{y}_{n-1} + \Delta t_n \mathbf{f}(t_{n-1}, \mathbf{y}_{n-1}), \\ t_n &= t_{n-1} + \Delta t_n,\end{aligned}$$

where  $\mathbf{y}_n \approx \mathbf{y}(t_n)$ . The time step  $\Delta t_n := t_n - t_{n-1}$  need not be constant for each  $n$ . It is well known that FE is first-order accurate [28], which is generally too low for efficiency in practice. Nonetheless, because of its simplicity, it is often used to simulate IVPs (4) associated with models of cardiac activity, and hence it serves as a useful benchmark against which to measure other numerical methods.

### B. Explicit Runge–Kutta Methods

The FE method can be viewed as the simplest of a more general class of numerical methods for solving IVPs known as *explicit Runge–Kutta* (ERK) methods; see, e.g., [7]. ERK methods aim to improve on the FE method by increasing the accuracy of the numerical solution by means of additional  $\mathbf{f}$  evaluations (or *stages*) within a given time step. A general  $s$ -stage RK method has the form

$$\mathbf{y}_n = \mathbf{y}_{n-1} + \Delta t_n \sum_{i=1}^s b_i \mathbf{k}_i,$$

where for  $i = 1, 2, \dots, s$ ,

$$\mathbf{k}_i = \mathbf{f} \left( t_{n-1} + \Delta t_n c_i, \mathbf{y}_{n-1} + \Delta t_n \sum_{j=1}^s a_{ij} \mathbf{k}_j \right),$$

and can be summarized via the *Butcher tableau* [7]:

$$\begin{array}{c|c} \mathbf{c} & \mathbf{A} \\ \hline & \mathbf{b}^T \end{array}$$

A Runge–Kutta method is *explicit* if  $\mathbf{A}$  is strictly lower triangular; otherwise it is *implicit*. With ERK methods, the stages can be computed successively and their contributions combined to produce a high-order approximation at the end of the step. With IRK methods, a (generally) nonlinear system of equations must be solved at every time step for all stages simultaneously. However, because of their superior stability properties, IRK methods are well-suited for stiff problems [7].

Perhaps the most popular high-order ERK method is the classical RK method, which is a four-stage, fourth-

order ERK method, and which we denote by ERK4:

$$\begin{array}{c|ccc} 0 & 0 & 0 & 0 \\ \frac{1}{2} & \frac{1}{2} & 0 & 0 \\ \frac{1}{2} & 0 & \frac{1}{2} & 0 \\ 1 & 0 & 0 & 1 \\ \hline & \frac{1}{6} & \frac{1}{3} & \frac{1}{3} & \frac{1}{6} \end{array}$$

Another popular ERK method is the Dormand–Prince 5(4) (DP) pair, which is of order 5 and is coupled with a method of order 4 for error estimation and step-size control [28]. This method is the basis of the popular Matlab routine `ode45`.

### C. Fully Implicit Methods

We include 3 fully implicit methods in our study. The first is arguably the simplest fully implicit method, the *backward Euler* (BE) method, a one-stage, first-order implicit RK (IRK) method with Butcher tableau

$$\begin{array}{c|c} 1 & 1 \\ \hline & 1 \end{array}$$

The second is the SDIRK method SDIRK4 of [29, p. 107]. This is a 5-stage,  $L$ -stable method of order 4 with an embedded method of order 3.

The third method is the 3-stage Radau IIA method RADAU5 of [29, p. 78]. It is a fully implicit RK method of order 5 and has the property of *stiff decay* [29].

We note that we only consider IRK and not BDF methods in the present study. In the context of splitting methods for mono- or bidomain models, the ODEs and PDEs are only advanced over small intervals in time before updated solution information is exchanged between systems. Results from [17] indicate that BDF methods were not competitive with IRK methods in terms of CPU time in such situations. Moreover, BDF methods require storage of the solution from past steps. This makes them less attractive than RK methods because this extra storage requirement may consequently reduce the overall resolution available for the simulation.

### D. Implicit-Explicit Methods

When the right-hand side of an ODE can be written as the sum of two terms

$$\frac{d\mathbf{y}}{dt} = \mathbf{f}_I(t, \mathbf{y}) + \mathbf{f}_E(t, \mathbf{y}), \quad (5)$$

it is often natural to consider approximating the contributions of  $\mathbf{f}_E(t, \mathbf{y})$  and  $\mathbf{f}_I(t, \mathbf{y})$  using different numerical methods. Such methods are known as *additive* methods. In general, when the right-hand side of an ODE can be written as the sum of  $n$  terms, these methods are called  $n$ -additive methods. When the constituent numerical

methods are RK methods, then they are known as  $n$ -additive RK methods. Furthermore, if  $dy/dt = \mathbf{f}_E(t, \mathbf{y})$  is such that it is best approximated with an explicit method and  $dy/dt = \mathbf{f}_I(t, \mathbf{y})$  is such that it is best approximated with an implicit method, we may use an IMEX method in an attempt to approximate the solution to this ODE efficiently [8]. An example of when an IMEX method would be useful is when  $\mathbf{f}_E(t, \mathbf{y})$  consists of non-stiff and/or nonlinear terms and  $\mathbf{f}_I(t, \mathbf{y})$  consists of stiff and/or linear terms. Again, when the constituent implicit and explicit methods are RK methods, we have an IMEX-RK method.

For example [30], the combination of the FE and BE methods gives the IMEX-RK method:

$$\mathbf{y}_n = \mathbf{y}_{n-1} + \Delta t_n (\mathbf{f}_E(\mathbf{y}_{n-1}) + \mathbf{f}_I(\mathbf{y}_n)).$$

More generally an  $s$ -stage IRK method with coefficients  $\mathbf{A}, \mathbf{c}, \mathbf{b}$  is combined with an  $(s+1)$ -stage ERK method with coefficients  $\hat{\mathbf{A}}, \hat{\mathbf{b}}, \hat{\mathbf{c}}$ . As is conventional, we assume that  $\hat{\mathbf{c}} = (0, \mathbf{c})^T$ , and the IRK method is taken to be an SDIRK method [30], [22]. An SDIRK method has a lower triangular  $\mathbf{A}$  with all diagonal elements  $a_{ii}$  non-zero and equal. This form of  $\mathbf{A}$  creates efficiencies in the solution to the nonlinear equations for the stages at each step, i.e., allowing each stage to be solved for sequentially with the same Newton iteration matrix.

In this study, we consider the IMEX-RK methods ARK3(2)4L[2]SA and ARK5(3)8L[2]SA from [22], which we denote by ARK3 and ARK5, respectively. ARK3 is an IMEX-RK method having 4 stages and order 3 with an embedded method of order 2 for error estimation and automatic step-size control; ARK5 has 8 stages, order 5, and an embedded method of order 4. The Butcher tableaux of ARK3 and ARK5 are listed in Appendix C of [22].

We split the ODE (4) on each time step  $[t_{n-1}, t_n]$  with the (default) dynamic splitting in `odeToJava`, namely  $\mathbf{f}_I(t, \mathbf{y}) := \mathbf{J}(t_{n-1}, \mathbf{y}_{n-1})\mathbf{y}(t)$  and  $\mathbf{f}_E(t, \mathbf{y}) := \mathbf{f}(t, \mathbf{y}) - \mathbf{J}(t_{n-1}, \mathbf{y}_{n-1})\mathbf{y}(t)$ , where  $\mathbf{J} := \partial\mathbf{f}/\partial\mathbf{y}$ . We note that this splitting is such that only the linear term is treated implicitly, and hence there is no need for a Newton iteration when solving the implicit equations. This makes the method similar to *Rosenbrock methods*, see, e.g., [8], which are also linearly implicit methods but for ODEs (4) that are not split.

### E. The Rush–Larsen Method

An alternative to using classical methods such as FE that is popular in the cardiac simulation literature is the *Rush–Larsen* (RL) method. The RL method advances the

solution to the gating equations (2) using

$$y_n = y_\infty + (y_{n-1} - y_\infty)e^{-\frac{\Delta t_n}{\tau_y}},$$

which represents the exact solution of (2) *assuming all variables besides  $y$  are constant*. FE is then used to advance the solution of the remaining equations. Using this method the Luo–Rudy model, for example, is no longer stiff [31]; i.e., the time step-size can be chosen based on accuracy considerations. However, this method is only first-order accurate and thus suffers from the usual drawbacks of low-order methods.

### F. Accuracy and Stability

An IVP is often called *stiff* if the choice of step-size  $\Delta t_n$  of a numerical method is determined by stability requirements rather than by accuracy requirements<sup>2</sup>. Generally, the step-size required for a stiff problem is much smaller than accuracy requirements dictate. In such cases, the numerical solution is typically much more accurate than required by the user.

When a numerical method is able to produce a stable approximation, we are then interested in the accuracy of the approximation. When the exact solution is not known, we may be able to generate a *reference solution* by using a variable step-size solver with low error tolerances until two approximations are produced that agree to a desired number of significant digits. In this study, we generate a reference solution by using a high-order, variable step-size implicit solver and lowering the error tolerances for successive approximations until two approximations are identical for at least 10 significant digits at  $N$  equally spaced output points  $t_i = it_f/N$ ,  $i = 1, 2, \dots, N$ , with  $N = 100$ . We can then measure the error in the approximation,  $\mathbf{y}$ , relative to the reference solution,  $\hat{\mathbf{y}}$ . A popular way to quantify error in the literature on heart simulation is the *Relative Root Mean Squared (RRMS) error* of the transmembrane potential [32]:

$$RRMS := \sqrt{\frac{\sum_{i=1}^N (V_i - \hat{V}_i)^2}{\sum_{i=1}^N \hat{V}_i^2}},$$

where  $V_i$  is the numerical approximation and  $\hat{V}_i$  is the reference solution at time  $t_i$  as described above. Given the many approximations made in creating the model, an RRMS error of 5% is generally considered acceptable.

As a more familiar measure of error, we also quantify error via the *global error*, which we define as

$$e_{\text{global}} := \max_i |V_i - \hat{V}_i|, \quad i = 1, 2, \dots, N.$$

<sup>2</sup>Arguably, there is no universally accepted definition of a stiff problem, but this description of stiffness suffices for our study.

## IV. RESULTS

Using `odeToJava` we performed numerical experiments with the FE, ERK4, RL, DP, BE, SDIRK4, RADAU5, ARK3, and ARK5 methods and compared their performance. Except for  $V_m$ , the initial values of the variables were taken to be those that correspond to the heart in its resting state. For the Luo–Rudy and Winslow et al. models, the initial values of  $V_m$  were chosen to produce the effect of an explicit stimulus current. That is, the initial values used for  $V_m$  are above the thresholds for the cells to fire. For the Courtemanche et al. and Puglisi–Bers models, the initial values for  $V_m$  were taken as their resting values, and an explicit stimulus was applied as given by (1). In particular, for the Courtemanche et al. model,  $I_{st} = -2000$  pA/pF from  $t = 0$  to  $t = 2$ ms and 0 elsewhere; for the Puglisi–Bers model,  $I_{st} = -10$   $\mu$ A/ $\mu$ F from  $t = 1$  to  $t = 5$  ms and 0 elsewhere. The specific initial values used for  $V_m$  are listed in Table I. See [18], [19], [20], [21] for complete listings of the remaining initial values. These values may also be accessed at <http://www.cellml.org/models/> under the directories `luo_rudy_1991_version04`, `courtemanche_ramirez_nattel_1998_version02`, `winslow_rice_jafri_marban_ororke_1999_version01`, and `puglisi_bers_2001_version01`, respectively.

TABLE I  
INITIAL VALUES FOR  $V_m = V_{rest}$ .

Model	$V_{rest}$
Luo–Rudy	-35.0
Courtemanche et al.	-81.2
Winslow et al.	-35.0
Puglisi–Bers	-85.5

The models were solved over time intervals representing one cardiac cycle. Different time intervals were used due to specific physiological properties of the mammalian heart that each model represents. Accordingly, the Luo–Rudy model was solved on the interval [0,450] ms, the Courtemanche et al. model was solved on the interval [0,500] ms, the Winslow et al. model was solved on the interval [0,300] ms, and the Puglisi–Bers model was solved on the interval [0,330] ms.

All the numerical experiments were performed on an Athlon 64 3000+ 1.8 GHz processor with 1 GB RAM. CPU times reported are the minimum of 5 runs. We note that the runs for FE, ERK4, RL, DP, ARK3, and ARK5 were performed within the `odeToJava` framework. The runs for BE were performed using `ode15s`

within Matlab with `MaxOrder` set to 1 and BDF to 'On'. In this case a conversion factor was determined to compare CPU time within Matlab to CPU time within `odeToJava`. This was done by computing the average of the ratios of a number of runs of `ode45` with the DP class within `odeToJava` for different tolerances. Similarly, the SDIRK4 and RADAU5 methods were run using their respective Fortran codes, and a conversion factor for CPU times was calculated by comparing runs of `DOPRI5.f` [33, p. 477] with the DP class in `odeToJava`. All CPU times reported below for BE, SDIRK4, and RADAU5 reflect this conversion.

### A. Customized Linear System Solver

With the splitting employed, a linear system involving  $\mathbf{J}(t, \mathbf{y})$  must be solved at each time step of an IMEX-RK integration. Accordingly, we are interested in the *sparsity pattern* of  $\mathbf{J}(t, \mathbf{y})$  across the entire solution interval of an ODE. A sparsity pattern can be thought of as a map of a matrix describing which entries of a matrix are always zero and which entries can be non-zero. The sparsity patterns of  $\mathbf{J}(t, \mathbf{y})$  for each of the 4 models were generated; see Figure 2 for the sparsity pattern of the Luo–Rudy model. If an element of  $\mathbf{J}(t, \mathbf{y})$  is always zero, it may be possible to omit it during Gaussian Elimination [34], the method used for solving linear systems by our implicit solver. This means that we may further optimize the IMEX-RK results compared to the previous section by customizing a Gaussian Elimination code to take advantage of these sparsity patterns. Results of the ARK3 and ARK5 methods with customized Gaussian Elimination routine are included below. In general not only do we see that such a customization results in a performance improvement for each of the 4 models over a wide range of tolerances, but we also see that the customization results in a variable step-size implementation of either ARK3 or ARK5 being the most efficient numerical method for every model.

### B. Constant Step-Size Tests

In Table II, we report the maximum step-size,  $\Delta t_{\max}$ , to 3 significant figures, for which FE, ERK4, and RL produce an approximation with less than 5% RRMS error. We also report the corresponding CPU times required, the RRMS error, and the global error.

For FE and ERK4,  $\Delta t_{\max}$  is also the step-size that produces a stable solution, indicating that these methods generally view the problems as stiff. Thus the resulting RRMS errors can be well below the desired levels. We also see that FE takes approximately 2–3 times less CPU time than ERK4 on all 4 models studied. Hence, as

TABLE II  
RESULTS FOR THE FE, ERK4, AND RL METHODS.

		$\Delta t_{\max}$	CPU Time	RRMS Error	Global Error
L	FE	1.34E-2	2.24E-1	3.08E-2	2.81E+0
R	ERK4	1.86E-2	6.77E-1	3.87E-2	3.51E+0
	RL	2.50E-1	4.29E-2	4.79E-2	5.39E+0
C	FE	1.94E-2	8.05E-1	2.30E-3	1.94E+0
R	ERK4	2.68E-2	2.30E+0	4.73E-2	6.73E+0
T	RL	3.45E-1	7.89E-2	4.97E-2	3.75E+1
W	FE	1.07E-4	4.04E+1	7.78E-8	9.91E-2
I	ERK4	1.30E-4	1.31E+2	3.40E-2	2.61E-1
N	RL	2.80E-4	2.25E+1	4.86E-2	6.08E+0
P	FE	1.08E-2	4.54E-1	5.20E-3	3.54E-1
B	ERK4	1.48E-2	1.08E+0	1.30E-2	1.33E-1
	RL	4.30E-1	6.50E-2	4.83E-2	8.54E+0

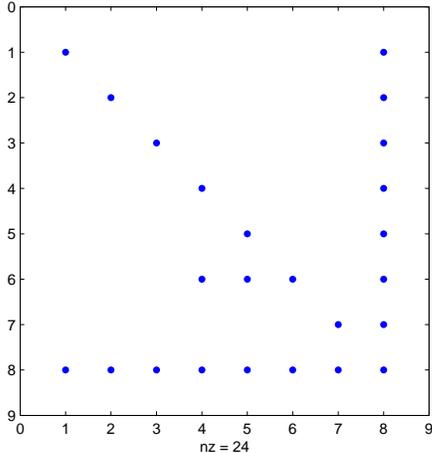


Fig. 2. Sparsity pattern for the Jacobian of the Luo-Rudy model.

is well known, higher order does not lead to greater efficiency when non-stiff methods are applied to stiff problems with moderate accuracy requirements.

We note that care must be exercised when determining  $\Delta t_{\max}$  for the RL method because the RRMS error produced is not a monotonically increasing function of the step-size. In other words, there exist  $\Delta t < \Delta t_{\max}$  for which the RRMS error *exceeds* 5%. We also see that RL always takes less CPU time than FE, ranging from approximately 2 times less on the Winslow et al. model to approximately 5–10 times less on the other 3 models.

### C. Variable Step-Size Tests

Tables III–VI respectively report the results from DP, BE, SDIRK4, RADAU5, ARK3, and ARK5 with vari-

able step-sizes applied to each of the 4 cardiac electrophysiological models. We run the models using standard error estimation and step-size control algorithms (see, e.g., [28]) for a range of absolute and relative tolerances. We set absolute tolerances equal to relative tolerances and define TOL to be their logarithm to base 10; e.g., TOL =  $-3$  implies both absolute and relative tolerances were set to  $10^{-3}$ . Integer values of TOL were run from  $-1$  to  $-6$  for all solvers and all 4 models, but details are reported for only the runs with the best CPU time that met the 5% RRMS error criterion.

For the Luo-Rudy (LR) model, ARK3 and ARK5 produce acceptable solutions in the least amount of CPU time. These methods are able to outperform the other methods in the study even without taking sparsity into account. In this case ARK3 with sparsity is the most efficient method. It is also over 60% faster than RL, its next closest commonly used competitor.

For the Courtemanche et al. (CRT) model, ARK3 produces an acceptable solution in the least amount of CPU time. With sparsity, ARK3 is over 40% faster than RL, its next closest commonly used competitor.

For the Winslow et al. (WIN) model, ARK3 again produces an acceptable solution in the least amount of CPU time. With sparsity, ARK3 is over 25% faster than SDIRK4, its next closest commonly used competitor.

Finally, for the Puglisi-Bers (PB) model, we see that ARK3 and ARK5 produce acceptable solutions in the least amount of CPU time, with or without taking sparsity into account. In this case ARK5 with sparsity produces an acceptable result almost 10 times faster than RL, its next closest commonly used competitor.

TABLE III  
RESULTS FOR THE LUO–RUDY MODEL.

	TOL	CPU Time (s)	Sparse CPU Time	RRMS Error	Global Error	Average $\Delta t$	Max $\Delta t$	Min $\Delta t$
DP	-2	5.20E-1	–	1.85E-2	2.41E+0	9.26E-2	4.99E-1	7.80E-3
BE	-3	1.45E-1	–	1.61E-2	2.92E+0	1.08E+0	5.89E+0	1.85E-3
SDIRK4	-1	4.80E-2	–	1.77E-2	2.05E+0	1.25E+1	6.78E+1	1.00E-3
RADAU5	-1	7.30E-2	–	6.75E-4	1.78E-2	1.25E+1	9.58E+1	1.00E-3
ARK3	-2	4.00E-2	1.80E-2	5.33E-3	1.91E+0	7.25E+0	6.42E+1	1.00E-3
ARK5	-2	3.40E-2	2.40E-2	7.46E-4	2.42E+0	1.15E+1	4.85E+1	6.36E-3

TABLE IV  
RESULTS FOR THE COURTEMANCHE ET AL. MODEL.

	TOL	CPU Time (s)	Sparse CPU Time	RRMS Error	Global Error	Average $\Delta t$	Max $\Delta t$	Min $\Delta t$
DP	-3	6.37E-1	–	5.26E-4	2.71E-1	1.66E-1	4.98E-1	2.84E-3
BE	-2	8.26E-2	–	2.40E-2	1.21E+1	2.34E+0	1.77E+1	2.82E-3
SDIRK4	-2	1.28E-1	–	7.51E-4	1.91E-1	9.43E+0	7.91E+1	1.00E-3
RADAU5	-1	2.40E-1	–	2.61E-3	2.79E+0	1.04E+1	5.00E+1	1.00E-3
ARK3	-2	6.90E-2	4.50E-2	7.87E-3	1.06E+0	7.14E+0	3.32E+1	1.00E-3
ARK5	-2	1.37E-1	6.00E-2	3.60E-2	2.28E+1	1.19E+1	5.43E+1	1.00E-3

TABLE V  
RESULTS FOR THE WINSLOW ET AL. MODEL.

	TOL	CPU Time (s)	Sparse CPU Time	RRMS Error	Global Error	Average $\Delta t$	Max $\Delta t$	Min $\Delta t$
DP	-3	3.00E+1	–	7.92E-3	4.15E-1	4.88E-3	6.08E-1	1.57E-4
BE	-4	9.51E-1	–	3.71E-2	1.25E+1	1.32E-1	5.38E+0	8.58E-5
SDIRK4	-2	2.72E-1	–	5.03E-3	5.81E-1	3.75E+0	4.79E+1	1.00E-3
RADAU5	-3	6.13E-1	–	7.68E-3	9.84E-1	2.65E+0	2.70E+1	1.00E-3
ARK3	-3	2.68E-1	1.97E-1	1.46E-2	1.84E+0	1.69E+0	3.45E+1	1.00E-3
ARK5	-3	4.65E-1	2.92E-1	3.15E-2	3.96E+0	3.06E+0	3.45E+1	1.00E-4

TABLE VI  
RESULTS FOR THE PUGLISI–BERS MODEL.

	TOL	CPU Time (s)	Sparse CPU Time	RRMS Error	Global Error	Average $\Delta t$	Max $\Delta t$	Min $\Delta t$
DP	-2	1.53E+0	–	4.43E-2	1.30E+0	4.09E-2	4.94E-1	4.71E-3
BE	-3	1.74E-1	–	5.69E-3	7.62E+0	6.13E-1	1.01E+1	1.76E-1
SDIRK4	-3	1.36E-1	–	8.48E-3	5.85E+0	4.92E+0	4.45E+1	1.00E-3
RADAU5	-2	1.53E-1	–	4.77E-2	5.53E+1	6.34E+0	7.18E+1	1.00E-3
ARK3	-4	6.90E-2	4.60E-2	2.38E-3	5.04E-1	4.12E+0	5.50E+1	1.00E-3
ARK5	-3	1.12E-2	7.00E-3	1.89E-2	1.63E+1	1.48E+0	7.20E+1	1.00E-4

#### D. Constant Step-Size IMEX

We also investigated the use of constant step-size ARK3 and ARK5 methods. We find that these constant step-size implementations significantly underperform the corresponding variable step-size implementations. They also significantly underperform the RL method on all of the models studied. We omit further details.

#### V. CONCLUSIONS

In this paper we compared the performance of several numerical methods for approximating solutions to ODEs found in 4 popular mathematical models of cardiac electrical activity. In particular, we compared the performance 2 IMEX-RK methods (ARK3 and ARK5) to other commonly used numerical methods for these models, i.e., FE, ERK4, RL, DP, SDIRK4, and RADAU5.

For constant step-sizes, the RL method is the most efficient for all the models studied here. It ranges from being approximately 2 times faster than the FE method for the Winslow et al. model to approximately 5–10 times faster than FE for the other models.

For variable step-sizes, the ARK methods outperformed the Dormand–Prince method for all 4 models. We obtained qualitatively similar results from a comparison of the ARK methods with the Bogacki–Shampine 3(2) method, which is the underlying method behind Matlab’s `ode23` routine; we do not comment on this further.

Overall, a variable step-size implementation of ARK3 or ARK5 with a customized linear system solver was the most efficient numerical method for all 4 mathematical models of cardiac electrical activity considered here. The results in this paper indicate that it is generally advisable to use a numerical method with an inexpensive implicit component and implemented with variable step-sizes and specialized techniques that take advantage of specific problem structure. When variable step-sizes are not possible, the RL method is the most efficient method.

It may be possible to further increase the efficiency advantage of IMEX-RK methods over the methods studied here by modifying the definitions of  $\mathbf{f}_E$  and  $\mathbf{f}_I$  in (5) from the default definitions in `odeToJava` or by constructing new IMEX-RK methods.

In this paper we have only considered ODE models of one cell. Further investigations of models involving large numbers of cells coupled with PDEs in two or three dimensions are necessary to fully establish the potential of the results presented here. In particular, linearly implicit IMEX operator-splitting methods that do not use constant step-sizes may lead to substantial performance gains. We report elsewhere on research efforts in these directions.

#### ACKNOWLEDGMENT

The authors would like to thank Mary MacLachlan for reference Matlab code for the Luo–Rudy, Courtemanche et al., and Winslow et al. models and Sarah Flaim for reference Matlab code for the Puglisi–Bers model. We would also like to thank the reviewers for their comments. R.C.D. thanks Rong Wang for his help with some of the Fortran codes.

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