Spiral- and scroll-wave dynamics in mathematical models for canine and human ventricular tissue with varying Potassium and Calcium currents

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A B S T R A C T

We conduct a systematic, direct-numerical-simulation (DNS) study, in mathematical models for ventricular tissue, of the dependence of spiral- and scroll-wave dynamics on $G_{K_r}$, the maximal conductance of the delayed rectifier Potassium current ($I_{K_r}$) channel, and the parameter $\gamma_{Ca}$, which determines the magnitude and shape of the current $I_{CaL}$ for the L-type calcium-current channel, in both square and anatomically realistic, whole-ventricle simulation domains. We study canine and human models. In the former, we use a canine-ventricular geometry, with fiber-orientation details, obtained from diffusion-tensor-magnetic-resonance-imaging (DTMRI) data; and we employ the physiologically realistic Hund-Rudy-Dynamic (HRD) model for a canine ventricular myocyte. To focus on the dependence of spiral- and scroll-wave dynamics on $G_{K_r}$ and $\gamma_{Ca}$, we restrict ourselves to an HRD-model parameter regime, which does not produce spiral- and scroll-wave instabilities because of other, well-studied causes like a very sharp action-potential-duration-restitution (APDR) curve or early after depolarizations (EADs) at the single-cell level. We find that spiral- or scroll-wave dynamics are affected predominantly by a simultaneous change in $I_{CaL}$ and $I_{K_r}$ from their original values in the model, rather than by a change in any one of these currents; other currents do not have such a large effect on these wave dynamics in this parameter regime of the HRD model. In particular, we examine spiral-wave dynamics for ten different values of $G_{K_r}$ and ten different values of $\gamma_{Ca}$ in our 2D DNSs. For our 3D DNSs in an anatomically realistic domain, we chose 16 parameter sets. In the parameter regime we begin with, the system displays broken spiral or scroll states with S1–S2 initial conditions (see below). We show that, by simultaneously increasing $G_{K_r}$ and reducing $\gamma_{Ca}$, we can get to a parameter regime in which the system displays single, stable rotating spirals or scroll waves. We obtain stability diagrams (or phase diagrams) in the $G_{K_r}$–$\gamma_{Ca}$ plane; and we find that these diagrams are significantly different in our 2D and 3D studies. In the 3D case, the geometry of the domain itself supports the confinement of the scroll waves and makes them more stable compared to their spiral-wave counterparts in our flat, 2D simulation domain. Thus, a combination of functional and geometrical mechanisms produce different dynamics for 3D scroll waves and their 2D spiral-wave counterparts. In particular, the former do not break easily because, in an anatomically realistic ventricular geometry, they are not easily absorbed at boundaries, nor do they break near boundaries. We have also carried out a comparison of our HRD results with their counterparts for the human-ventricular TP06 model; and we have found important differences between wave dynamics in these two models. The region in parameter space, where we obtain broken spiral or scroll waves in the HRD model is the region of stable rotating waves in the TP06 model; the default parameter values produce broken waves in the HRD model, but stable scrolls in the TP06 model. In both these models, to make a transition, (most simply, from broken-wave to stable-scroll states) we must simultaneously increase $I_{K_r}$ and decrease $I_{CaL}$; a modification of only one of these currents is not enough to effect this transition. Furthermore, the converse, i.e., an increase in $I_{CaL}$ along with a decrease in $I_{K_r}$ does not yield any interesting dynamical transitions in the HRD model, for, in this range of currents, this model does not sustain spiral or scroll waves or broken waves.

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1. Introduction

Studies of mathematical models of cardiac myocytes and cardiac tissue play an important role in understanding the complex mechanisms that underlie cardiac arrhythmias, which are a major cause of death. These arrhythmias are believed to be associated with reentrant waves of electrical activation in cardiac tissue; specifically, rotating spiral or scroll waves are associated with ventricular tachycardia (VT) and the breaking of such waves is associated with ventricular fibrillation (VF). Understanding the detailed ionic mechanisms leading to spiral or scroll waves and the response of these wave dynamics to the changes in various ionic mechanisms is still a difficult task, because we must account for the large number of ion channels, ion pumps, and intracellular mechanisms that are involved in producing the action potential (AP). Computational tools are becoming more and more useful in these studies ([1–10]) because they allow us increased control and flexibility in handling each parameter and, hence, the associated ionic current; such control is rarely feasible in experiments. A number of studies have been conducted on the mechanisms of ion-channel kinetics and their dependence on various channel parameters or the AP morphology. But there are a few detailed studies of how the current-channel parameters directly affect scroll waves, at the whole-heart level in three dimensions (3D). Furthermore, experiments on mammalian hearts are challenging because of the difficulty in the visualization of waves of electrical activation below the tissue surface [9–18]; for recent advances in such visualization, we refer the reader to Refs. [19,20].

Canine hearts are often used in studies of the mechanisms of cardiac arrhythmias because their size and physiology are comparable to those of human hearts. Furthermore, detailed mathematical models for canine cardiac tissue are now available; these models incorporate important electrophysiological details, including intracellular calcium dynamics. In our work, we use a detailed mathematical model for canine ventricular myocytes, namely, the Hund-Rudy-Dynamic (HRD) model [21,22]. We use a canine ventricular geometry, with fiber-orientation details, which have been obtained in earlier studies that use diffusion-tensor magnetic-resonance imaging (DTMRI). These DTMRI data have been made available for academic use at the CMISS site (https://www.cmiss.org/). Most of the studies on scroll-wave dynamics have been carried out on models that are not as detailed as the HRD model; e.g., many studies use the Luo–Rudy model for guinea pigs [23]. Thus, our work goes significantly beyond such earlier studies.

In particular, we carry out a systematic, in silico direct-numerical-simulation (DNS) of spiral and scroll waves in the HRD mathematical model for canine ventricular tissue. We explore the dependence of spiral- and scroll-wave dynamics on $G_{K_{r}}$, the maximal conductance of the delayed rectifier Potassium current ($I_{K_{r}}$) channel, and the parameter $\gamma$ which determines the magnitude and shape of the current $I_{Ca,L}$ for the L-type calcium-current channel, in both square and anatomically realistic, whole-ventricle simulation domains. We focus on the dependence of spiral- and scroll-wave dynamics on $G_{K_{r}}$ and $\gamma$; therefore, we limit ourselves to a parameter regime, in which the HRD model does not display spiral- and scroll-wave instabilities arising from other well-explored causes, like a very sharp action-potential-duration-restitution (APDR) [24–26] curve or early after depolarizations (EADs) at the single-cell level. We find that spiral- or scroll-wave dynamics are affected predominantly by a simultaneous change in $I_{Ca,L}$ and $I_{K_{r}}$, rather than by a change in any one of these currents; other currents do not display such a large effect on these wave dynamics in this parameter regime. To carry out a systematic study, we examine spiral-wave dynamics for ten different values of $G_{K_{r}}$ and ten different values of $\gamma$ in our 2D DNSs. For our 3D DNSs in an anatomically realistic domain, we choose 16 parameter sets. In the parameter regime we begin with, the system displays broken spiral or scroll states with the S1–S2 initial conditions (see below). We show that, by simultaneously increasing $G_{K_{r}}$ and reducing $\gamma$, we reach a parameter regime in which the system displays a single, stable rotating spiral or scroll wave. We obtain stability diagrams (or phase diagrams) in the $G_{K_{r}} - \gamma$ plane; and we find that these diagrams are significantly different in our 2D and 3D studies. In the 3D case, the geometry of the domain itself supports the confinement of the scroll waves and makes them more stable compared to their spiral-wave counterparts in our flat, 2D simulation domain. Thus, a combination of functional and geometrical mechanisms produce different dynamics for 3D scroll waves and their 2D spiral-wave counterparts: The former do not break easily because, in an anatomically realistic geometry, they are not easily absorbed at boundaries; nor do they break near boundaries.

We have mentioned above that canine hearts are considered to be similar to human hearts in both shape and electrophysiology. It is important to explore this similarity. We begin such and exploration by comparing our HRD-model results with their counterparts for the human-ventricular TP06 mathematical model [27]. We use a human-ventricular geometry with fiber rotation; to obtain the coordinates in the human-ventricular geometry we use Ref [28]. By carrying out in silico DNSs of spiral and scroll waves in this TP06 model, we find important differences between wave dynamics in these two models. The region in parameter space, where we obtain broken spiral or scroll waves in the HRD model is the region of stable rotating waves in the TP06 model; the default parameter values produce broken waves in the HRD model, but stable scrolls in the TP06 model. However, in both these models, to make a transition (most simply, from broken-wave to stable-scroll states), we must simultaneously increase $I_{K_{r}}$ and decrease $I_{Ca,L}$; a modification of only one of these currents is not enough to effect this transition. Furthermore, the converse, i.e., an increase in $I_{Ca,L}$ along with a decrease in $I_{K_{r}}$ does not yield any interesting dynamical transitions in the HRD model, for, in this range of currents, this model does not sustain unbroken or broken waves.

In the Supplementary Material [29], we describe the models we use for our DNSs, namely, the HRD model, for a canine-ventricular myocyte, and the TP06 model, for a human-ventricular myocyte. We give a description of the two currents which are the subject of this study in each models. We also describe the anatomically realistic geometry and the numerical methods that we use to study scroll dynamics.

The remaining part of this paper is organized as follows. Section 2 describes briefly the DNSs we have conducted, the details of which are given in the Supplementary Material [29]. Section 3 and Section 4 are devoted to our results, which are presented in two parts, the first for the HRD model and the second for the TP06 model. Each part has two subsections, that are devoted, respectively, to our results for 2D tissue, and 3D anatomically realistic domains; we compare our results from HRD and TP06 models. We describe the results of our cellular-level studies in the Supplementary Material [29]. We also present the variation of the two important currents, which we focus on in this study (as we change model parameters), along with the corresponding APDR curves, for both HRD and TP06 models. Section 3 contains a discussion of our results and conclusions.

2. Models and numerical methods

2.1. Canine ventricular (HRD model) simulations

We have used the physiologically detailed HRD mathematical model for canine ventricular tissue. This is a dynamic model that reproduces the experimentally measured action potential (AP) and Calcium-current regulation over a wide range of the pacing frequency. The HRD model incorporates a total of 15 ionic currents:

$$I_{Na}, I_{Na,L}, I_{Ca,L}, I_{Na,Ca}, I_{K}, I_{K_{r}}, I_{Ca}, I_{Ca,T},$$
$$I_{K_{r}}, I_{K_{p}}, I_{Cl}, I_{Cl,T}, I_{Cl}, I_{L}, I_{L,T},$$

This model uses 21 gating variables, namely,

$$H, m, J, d, f_{1}, f_{2}, f_{Ca},$$
$$f_{Ca,T}, p, r, s, t, a, b, y,$$
$$y_{d}, y_{d}, 2, 3, A, M_{1}, f_{2}, e_{2}, r_{1}.$$
and the following 8 ionic concentrations:

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\[ C_{\text{Ca}}, N_{\text{a}}, C_{\text{l}}, K_{\text{r}}, C_{\text{a},\text{in}}, C_{\text{a},\text{out}}, C_{\text{a,M}}, K_{\text{dep}}. \]

The details of the model and complete equations are given in the Supplementary Material [29]. In this model we calculate the transmembrane potential \( V \) as a dynamical function of the above mentioned currents, concentrations, and gating variables. We give tables with (a) a list of the currents in the HRD model and their descriptions and (b) ionic concentrations in the Supplementary Material [29].

We depict in Fig. 1 the shape of the action potential AP for the HRD canine-myocyte model, showing its different phases along with the major currents responsible for each stage. This figure also shows, for a representative set of values for \( G_{\text{Kr}} \) and \( \gamma_{\text{CaO}} \), the APs that we obtain with continuous pacing of the myocyte. We show for comparison, in Fig. 2, similar plots for the TP06 human-myocyte model.

Our 2D simulation domain, for the HRD model, is a square tissue with size 8.32 cm \( \times \) 8.32 cm. For our 3D simulation we use the processed Diffusion-Tensor Magnetic-Resonance Imaging (DTMRI) data for the canine-ventricular anatomy, which is freely available for academic purposes at the CMISS website (https://www.cmiss.org/), the details of which are given in the Supplementary Material [29].

Fig. 3 shows the geometry of the domain in which we investigate scroll-wave dynamics.

We describe the S1–S2 protocol, which we use to produce the initial scroll waves in the Supplementary Material [29].

For the HRD model, we investigate the effect on spiral- and scroll-wave dynamics of the following two currents.

(1) The L-Type Calcium current is given by the equations

\[ I_{\text{CaL}} = d \cdot f_{\text{CA}} \cdot f_{\text{Ca2}} \cdot f \cdot I_{\text{CaL}}. \]

\[ \dot{I}_{\text{CaL}} = P_{\text{Ca}} \cdot \frac{z_{\text{Ca}}}{RT} \cdot \frac{V_{m} - E_{\text{Ca}}}{F^{2}}. \]

\[ \gamma_{\text{CaO}} \cdot [C_{\text{a}}]_{\text{in}} \cdot \exp\left( \frac{\gamma_{\text{CaO}}(V_{m}-15.0)F}{RT} \right) - \gamma_{\text{CaO}} \cdot [C_{\text{a}}]_{\text{in}} \cdot \exp\left( \frac{\gamma_{\text{CaO}}(V_{m}-15.0)F}{RT} \right) \]

where \( d, p, f_{\text{CaO}} \) and \( f_{\text{Ca2}} \) are gating variables; \( P_{\text{Ca}} \) is the membrane permeability to the Ca ion, in units of cm/s, and \( z_{\text{Ca}} \) is the valence of the Ca ion. The parameters \( \gamma_{\text{CaO}} \) and \( \gamma_{\text{CaO}} \) determine the magnitude and shape of the current \( I_{\text{CaL}} \). They are the activity coefficient of the Ca ion. We find that \( \gamma_{\text{CaO}} \) has a stronger influence on the dynamics of the system than does \( \gamma_{\text{CaO}} \). Therefore, we have chosen to study how the variation of \( \gamma_{\text{CaO}} \), over a wide range, affects spiral- and scroll-wave dynamics in the HRD model.

(2) The delayed rectifier Potassium current is

\[ I_{\text{Kr}} = G_{\text{Kr}} \cdot X_{\text{r}} \cdot R_{\text{Kr}} \cdot (V_{m} - E_{\text{K}}). \]
As we have mentioned above, Fig. 2 gives the shape of the AP for this TP06 model; it depicts different phases along with the major currents responsible for each phase; this figure also shows, for representative values of $G_{Kr} = GKr$ and $G_{CaL} = GCaL$, the APs with continuous pacing of the myocyte.

In Fig. 4 we have shown the anatomical geometry that we have created for our simulations by using the DTMRI data for the coordinate mesh of the human heart. This is the lower part of the heart that contains the ventricles and the septum.

The L-Type calcium current in the TP06 model is described by the following equations:

\[
I_{CaL} = G_{CaL} d.f.d.f Ca \cdot \frac{(V - 15)F^2}{RT} \times 0.25Ca \exp \left( \frac{20(V - 15)F}{RT} \right) - Ca \exp \left( \frac{2V - FRT}{RT} \right) - 1 \]

where $d$, $fCa$, and $f$ are gating variables. The parameter $G_{CaL}$ determines the magnitude and shape of the current $I_{CaL}$. As in our work on the HRD model, we study how the variation of $G_{CaL}$ over a wide range, affects spiral- and scroll-wave dynamics in the TP06 model. Note that the variation of $G_{CaL}$ in the TP06 model corresponds to the variation of $\gamma_{CaO}$ in the HRD model.

In the TP06 model, the delayed rectifier Potassium current is

\[
I_{Kr} = G_{Kr} \sqrt{K^+} \times \frac{1}{5.4} \times 153 \times (V - E_{Kr}) \]

For our DNSs of human-ventricular tissue, we use the TP06 [27] model, which is a modified version of the TNNP model for human-ventricular cells [30]; this incorporates a total of 12 ionic currents. The intracellular calcium handling in these models is not as detailed as it is in the HRD model.

The TP06 model uses 19 variables: (a) 1 for the transmembrane potential $V_{m}$; (b) 13 for ion-channel gates, namely, $m, h, l, d, f, f_2, I_{f2}, r, s, x_{1}, x_{2}, x_{3}, x_{4}$, and $R$; and (c) 5 for intracellular, ion-concentration dynamics, namely, $Na_{j}$, $K_{i}$, $Ca_{j}$, $Ca_{Ca}$, and $Ca_{al}$.

As we have mentioned above, Fig. 2 gives the shape of the AP for this TP06 model; it depicts different phases along with the major currents responsible for each phase; this figure also shows, for representative values of $G_{Kr} = GKr$ and $G_{CaL} = GCaL$, the APs with continuous pacing of the myocyte.

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where $d$, $fCa$, and $f$ are gating variables. The parameter $G_{CaL}$ determines the magnitude and shape of the current $I_{CaL}$. As in our work on the HRD model, we study how the variation of $G_{CaL}$ over a wide range, affects spiral- and scroll-wave dynamics in the TP06 model. Note that the variation of $G_{CaL}$ in the TP06 model corresponds to the variation of $\gamma_{CaO}$ in the HRD model.

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In our DNSs, we investigate 20 different cases in 2D tissue: $G_{Kr}$ is taken as $G_{Kr} = GKr \times N_{GKr}$, where $GKr = 0.153$, and $N_{GKr} = 1.3...7$. Along with this, $G_{CaL}$ is varied as $G_{CaL} = GCaL/N_{GCaL}$, where $GCaL = 0.0000398$ and $N_{GCaL} = 1.3...9$. $GKr$ is the original value of $G_{Kr}$, as it appears in the model described by Eq. (5); likewise, $GCaL$ is the original value of $G_{CaL}$, as it appears in the model described by Eq. (7).

For our DNSs in the 3D anatomical geometry, we investigate 20 different cases as follows: $G_{Kr} = GKr \times N_{GKr}$, $N_{GKr} = 1.3...9$; and $G_{CaL} = GCaL/N_{GCaL}$, $N_{GCaL} = 1.3...7$.

3. Results

We present results from our simulations of the HRD model, at the cell level, in the Supplementary Material [29]. The variations of the AP morphologies and APDR curves, for all the parameter values in our DNSs, are presented along with the plots of the currents $I_{Kr}$ and $I_{CaL}$. In this section, we give results from our studies in 2D and then in 3D for the HRD model. Next we present the results of our simulations for the TP06 model.

It has been observed previously that the action potential duration restitution (APDR) is a crucial factor, which determines whether scroll waves or broken scroll waves develop in cardiac tissue [24–26,31,32]. The APDR is the shortening of the action potential duration (APD) as the pacing frequency. A sharp APDR with a slope $< 1$ is a strong indicator of APDR. As in our work on the HRD model, we study the effect of $G_{CaL}$ over a wide range, affects spiral- and scroll-wave dynamics in the TP06 model. Note that the variation of $G_{CaL}$ in the TP06 model corresponds to the variation of $\gamma_{CaO}$ in the HRD model.

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3.1. 2D results

The conduction velocity $c$ of a plane wave, passing from one end to the other end in our 2D simulation domain, is the same for all the cases we study; we find $c = 6.7002$ m/s. By contrast, the wavelength $\lambda$ of the plane wave varies for each case; we define the wavelength to be the distance between the excited front and the 90% recovered back end of the propagating plane wave of the transmembrane potential $V$ (before the initiation of the spiral wave). As we have noted above, we can also
use the formula $\lambda = cu \times APD$, where APD is the action potential duration. We measure $\lambda$ for each of our parameter sets. The spiral wave, which we use as initial condition for our 2D DNSs, is created by using the S1–S2 protocol, which we describe in the Supplementary Material, [29], where we give representative pseudocolor plots of $V$ for plane-wave and spiral-wave initial conditions.

Given the range of parameter values we use, the initial spiral state shows rich and varied spatiotemporal evolution. Here, we give a detailed description of the various kinds of spiral-wave evolutions that we observe in the large parameter space we investigate.

We observe two (main) kinds of spiral arms, which we show in Fig. 5: (1) The first is a spiral with nonuniform arm widths at different regions in the simulation domain; these regions can have inherent instabilities, which cause spiral-arm thinning (but not enough to lead to the breaking up of the spiral-arm). They are stable and preserve their shape. The (average) wavelength of such a spiral wave is remarkably different from that of the plane wave from which it is formed. This kind of spiral arm is formed from plane waves of large width (as we find for low values of $G_{Kr}$, and high values of $\gamma_{Cao}$). (2) A spiral with almost uniform arm width. These spirals, which are stable and retain their uniform arm-width, are formed from small-width plane waves (as we find for high values of $G_{Kr}$, and low values of $\gamma_{Cao}$). Here, the spiral wavelength is comparable to that of the plane wave.

In summary, the spirals that form, in the different parameter regimes in the HRD model, have different sizes and shapes, even though they might evolve in a roughly similar manner to the final state and be characterized by generic terms such as spiral breakup or spiral rotation. In Fig. 6 we give a phase diagram (or stability diagram) in the $G_{Kr}$–$G_{Ca}$ plane. [Recall that $G_{Kr} = G_{RX}N_{GKr}, \text{where } N_{GKr} = 1, 2, ..., 10$, and $\gamma_{Cao} = G_{CAO}/N_{GCao}$, where $N_{GCao} = 1, 2, ..., 10$.] For this diagram we have used the results from the 100 different pairs of parameter values that we have studied in the 2D HRD model. 30. The 3 main types of dynamics we can observe are: broken waves, combined broken and rotating waves, and rotating waves. In each region we see some parts that sustain stable waves, i.e., the waves stay in the medium; in some other parts there are unstable waves, i.e., they move away and disappear from the medium. They are shown with different markers in each colored region (details are given in the figure caption).

In the break-up region we see two different kinds of phenomena: waves breaking up from the spiral core (core break-up); or waves breaking from the outer arms (far-field break-up). The far-field break-up is weak because it does not spread all the way to the core.

In Fig. 7 we show pseudocolor plots of $V$, from different stages of the spiral-wave evolution, going from left to right, for each of the different kinds of wave dynamics. The time-evolution of the spiral waves in the parameter-space we study is of the following seven major types (shown in the seven rows of Fig. 7): (a) No spiral wave is formed ($N_{GKr} = 2, N_{GCao} = 10$); the waves disappear from the medium before evolving into a spiral [magenta square ■ in Fig. 6]. (b) A stable state with broken spirals is obtained ($N_{GKr} = 2, N_{GCao} = 6$) [● in Fig. 6]; the broken spirals interact and regenerate themselves, without disappearing. (c) A stable single rotating spiral is formed ($N_{GKr} = 7, N_{GCao} = 7$) [green bubbles in Fig. 6]. (d) There is unstable break-up in which the broken waves quickly disappear (e.g., for $N_{GKr} = 1, N_{GCao} = 1$) [red stars ✸ in Fig. 6]. (e) A region where a spiral breaks, recombines, rotates, again breaks and so on [green diamond, ◇, in Fig. 6]. (f) There is far-field break-up (observed for $N_{GKr} = 5$ and $N_{GCao} = 10$) [black-faced hexagon ✶ in Fig. 6]. (g) There is an instability in the core region, exhibited by some parameter-combinations in the stable-rotating region (here, $N_{GKr} = 5$ and $N_{GCao} = 10$) [in Fig. 6 the regions marked by cyan colored bubbles]; such a core-break-up does not last, for the spiral core quickly regenerates itself; this has no far-reaching effect on the evolution of the spiral wave.

As we have mentioned earlier, the parameter region that we investigate in the HRD model leads to spiral-wave dynamics that does not obey the restitution hypothesis. According to this restitution hypothesis, if the slope of the APDR curves is $> 1$, the spirals break up; if the slope is $< 1$, the spirals do not break up. In our studies, the APDR slopes are always $< 0.25$, yet we see spiral breakup. However, we see a correlation between the maximal slopes of the APDR restitution curves and the spiral-wave dynamics. The region where we obtain stable rotating spiral wave dynamics corresponds to the lower-right region of the APDR curves [29]; this is where the slopes are smallest, for the APDR curve is flat. The APD gets smaller and smaller in this region. Small values of the APD corresponds to small wavelengths $\lambda$. Such waves, with small widths, are not very prone to break-up in the HRD model.

3.1.1. Dominant frequencies

In this subsection we examine the dominant frequencies of the spiral waves that are formed in our 2D simulations of the HRD model. For this we use the time series of the transmembrane potential $V$ from a few different sites in the simulation domain. The power spectra of these time series yield the dominant frequencies. In Fig. 8 we show the dominant frequencies that we obtain from power spectra for all our parameter values, i.e., for $N_{GKr} = 1, 2, ..., 10$ and $N_{GCao} = 1, 2, ..., 10$. (We give the spectra corresponding to these different wave dynamics in the Supplementary Material [29].) In the region where stable rotating waves exist, we see a prominent frequency in the spectrum. If the waves are unstable and disappear, it is not possible to identify a major frequency; the missing parameter values in this plot correspond to these unstable regions. Note that the frequency of the rotating wave increases as we increase $G_{Kr}$. The variation of $\gamma_{Cao}$ does not affect the dominant frequency substantially, as is evident from Fig. 8, where the curves are almost flat.

3.2. Three-dimensional (3D) results

We now present our results on scroll-wave dynamics from our DNSs of the HRD model on the realistic canine-heart geometry that we have described above.

In Fig. 9 we show different stages of the spatiotemporal evolution of a plane wave passing through our anatomically realistic simulation domain, from one end to the other end (here there are no obstacles). The lower panels show how the wave moves and finally disappears. After a plane wave has passed through this domain, we apply the S1–S2 cross-field stimulus to produce a scroll filament in the middle of the domain [29]. We then vary the parameters $G_{Kr}$ and $\gamma_{Cao}$ simultaneously and examine the effect of this change on development of the scroll wave for 3s in real time.

With the original parameters of the HRD model, we see that the scroll wave immediately breaks up and develops into a spatiotemporally chaotic state. Within the time duration of our DNS, these broken scroll waves continue to spread in the domain, interact, recombine,
and break up without disappearing; so the broken-wave state is statistically steady. The dynamics of broken scrolls for the parameters \( G_{Kr} = GKR \) and \( \tau_{CaO} = GCAO \) is shown in Fig. 10(b). If we decrease \( \tau_{CaO} = GCAO/N_{GCau} \), by using \( N_{GCau} = 3.5,7 \), the spatiotemporal evolution of scroll waves is qualitatively similar, with wave breaks and spatiotemporally chaotic behavior; but, for \( N_{GCau} = 3 \) and \( 7 \), the waves are unstable, they meander, break up, and finally disappear. This behavior is not visible for \( N_{GCau} = 1 \) and \( 5 \). Now we increase \( G_{Kr} \) by factors of \( 3 \), \( 5 \), and \( 7 \). In each case we vary \( \tau_{CaO} \) as above. Thus, we examine scroll-wave dynamics for \( 4 \times 4 \times 4 \) parameter sets.

The different kinds of scroll-wave developments that we observe in our DNSs of the 3D HRD model in the anatomically realistic domain are depicted in Fig. 10: In Fig. 10(a) we show the development of a rotating scroll wave, with meandering, but without break up for a representative parameter set \((N_{GKr} = 7 \) and \( N_{GCau} = 7 \)). In Figs. 10(b) and (c) we show, respectively, how a meandering scroll wave breaks up and spreads through the domain and how a scroll wave breaks while its central region passes through the right ventricle, but such a break-up goes away very soon, in a time \( \approx 100 \) ms, and the wave recombines. We see this anatomical break-up arising solely because of the geometry rather than from functional break-up associated with parameter dependence. This anatomical break-up duration is negligible; it does not have a significant impact on the long-term behavior of scroll-wave dynamics.

The regions of stability of these different scroll-wave behaviors are depicted in a phase diagram in Fig. 11 (see the figure caption for details). Observe that, as we move to the right and upper regions of this phase diagram, scroll waves tend to be stable and rotating; they do not break up to form a chaotic state. This region corresponds to \( N_{GKr} = 7.5 \) and \( N_{GCau} = 3.5,7 \), and also \( N_{GKr} = 3 \) and \( N_{GCau} = 5.7 \). And anatomical breakup of scroll waves plays no significant role in the long-term dynamics of the system. Here also, as in our study of the 2D HRD model, we obtain stable rotating scroll waves in the bottom-right region of the APDR curves [29], where the APDR curve is nearly flat.

### 3.2.1. Dominant frequencies

We now examine, as we did in 2D, the dominant frequencies of the scroll waves that form in our DNSs of the 3D HRD model in the anatomical heart geometry. We use time series of the transmembrane potential \( V \) from a few different sites from the simulation domain. In the Supplementary Material [29] we present two-level isosurface plots of \( V \) (left panel) showing different examples of scroll-wave dynamics and the corresponding power spectra (middle panel) of the time series (right panel) of \( V \) from a representative point in the domain.

In Fig. 12 we plot the dominant frequencies that we obtain from power spectra (see the Supplementary Material [29]), for \( N_{GKr} = 1.2,7 \) and \( N_{GCau} = 1.2,7 \). In the region where a stable rotating scroll wave exists, we see a prominent frequency in the spectrum. If the waves are unstable and disappear, it is not possible to identify a major frequency. Here also, as in the 2D case, the dominant frequency increases as \( G_{Kr} \) increases; and it is not affected significantly by a variation of \( \tau_{CaO} \).

### 4. Human-heart (TP06) model

We repeat the whole procedure described above on a human-heart geometry, with fiber orientation, by using the TP06 human-ventricular model. Our goal is to compare these results with their HRD-model counterparts. We present our results in two subsections, from 2D and 3D simulations. The shapes of the action potential (AP) and the APD-restitution (APDR) curves, for 5 representative cases for the rapid rectifier current \( I_{Kr} \), the Calcium current \( I_{calc} \), and different values of \( G_{CaL} \) and \( G_{Kr} \), for the TP06 model, are given in the Supplementary Material [29].

#### 4.1. 2D results

For our simulations of the 2D TP06 model, we follow the same methods, numerical schemes, and initial spiral conditions we have used in our HRD-model DNSs. The conduction velocity of a plane wave passing from one end to the other end in the 2D domain is the same for all the cases, which is \( 6.960 \) m/s for the TP06 model. We have measured...
Fig. 7. Pseudocolor plots of the transmembrane potential $V$ showing the different kinds of wave dynamics in our DNSs for the 2D-HRD model. The sub-panels, from the left to the right, show the evolution of the system at four different points of time (for the complete spatiotemporal evolution see the Videos S35, S36, S37, S38, S39, S310 and S311 in the Supplementary Material [29]). We observe the following seven types of spiral-wave dynamics (seven panels from the top to the bottom): (a) no spiral is formed ($N_{GKR} = 2$, $N_{GCaL} = 10$); the waves disappear before evolving into a spiral (magenta square ■ in Fig. 6); (b) a stable state with broken spirals ($N_{GKR} = 2$, $N_{GCaL} = 6$), (⧫ in Fig. 6); the broken spirals interact and regenerate themselves, without disappearing; (c) a stable single rotating spiral ($N_{GKR} = 7$, $N_{GCaL} = 7$), (green bubbles in Fig. 6); (d) an unstable break-up in which the broken waves quickly disappear ($N_{GKR} = 1$, $N_{GCaL} = 1$), (red stars ⋆ in Fig. 6); (e) region where a spiral breaks, recombines, rotates, again breaks and so on (green diamond, ⧫, in Fig. 6); (f) far-field break-up (observed for $N_{GKR} = 5$ and $N_{GCaL} = 10$) [black-faced hexagon ◆ in Fig. 6]; (g) a spiral with a local, short-lived instability of its core (here, $N_{GKR} = 5$ and $N_{GCaL} = 9$), in the stable-rotating region [in Fig. 6 the cyan colored bubbles ]. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

As we have described in Section 2, we vary $G_{Kr} = GKR \times N_{GKR}$, and $G_{CaL} = GCaL/N_{GCaL}$ to obtain a total of 20 different cases.

We now describe the types of spiral-wave dynamics that we obtain in the large parameter space we have studied in the 2D TP06 model. As in the canine 2D HRD model (Fig. 5 in Section 3.1), we observe two types of spiral arms. We depict these in Fig. 13.

In Fig. 14 we give the phase diagram (or stability diagram) for the different types of spiral-wave dynamics we obtain in the 2D TP06, for the 20 cases that we have studied. The 3 main colors in this phase diagram denote broken waves, a phase with both broken and rotating spirals, and rotating waves. In each region, we find that some waves are stable (they repeat themselves in time and do not move away from the

Fig. 8. The dominant frequencies that we obtain from power spectra for all our parameter values, i.e., for $N_{GKR} = 1..2..10$ and $N_{GCaL} = 1..2..10$. In the region where stable rotating waves exist, we see a prominent frequency in the spectrum. If the waves are unstable and disappear, it is not possible to identify a major frequency; the missing parameter values in this plot correspond to these unstable regions.

Fig. 9. Two-level isosurface plots of $V$ illustrating a wave passing from one end to the other of our anatomically realistic simulation domain geometry under normal conditions and without any obstacles. The upper panels show the initial stages, time increasing from left to right; and the lower panels show the wave finally disappearing from the domain. For the complete spatiotemporal evolution see the Video S12 in the Supplementary Material [29].

Fig. 10. Two-level isosurface plots of $V$ illustrating the following: (a) A scroll wave in the stable rotating state; it rotates with a slightly meandering core, without breaking up ($N_{GKR} = 7$, $N_{GCaL} = 7$). (b) A meandering scroll wave breaks up and spreads through the domain leading to a statistically stable chaotic state, for the representative parameter set $N_{GKR} = 1$, $N_{GCaL} = 1$. (c) A scroll wave breaking while its central region passes through the right ventricle; this broken part quickly recombines and continues as a rotating stable wave, thus showing that such anatomical structures have only short-term effects on the wave dynamics (here, $N_{GKR} = 7$, $N_{GCaL} = 3$). For the complete spatiotemporal evolution, see Videos S13, S14 and S15 in the Supplementary Material [29].
In this subsection we examine the dominant frequencies of the spiral waves formed in our 2D TP06 simulations. As we did in our study of domain), whereas others are unstable (they move away and disappear from the domain); these waves are shown with different markers in each colored region. As we go from left to right in this phase diagram, the meandering of the wave increases, so it is prone to break up; eventually, for high values of \( G_{Kr} \), the wave does break up. [Fig. 14 is the counterpart of Fig. 6 for the HRD model.]

In Fig. 15 we show the different kinds of spiral-wave dynamics that we observe in our DNSs for the 2D TP06 model. The panels from the left to the right represent the evolution of the system at four different points of time. We observe the following four types of spiral-wave dynamics. (a) A stable, single rotating spiral state (for \( N_{GKr} = 1, N_{GCal} = 3 \); stable spiral states are represented by red bubbles in the phase diagram of Fig. 14. (b) A stable state with broken spirals (here, \( N_{GKr} = 7, N_{GCal} = 9 \); the broken spirals interact with each other, regenerates themselves, and do not leave the domain. This is denoted by a black star (\( \ast \)) in Fig. 14. (c) We see alternately rotating and breaking spirals (here, for \( N_{GKr} = 5 \) and \( N_{GCal} = 9 \); in Fig. 14, a cyan diamond, \( \bigstar \)). (d) In some parts of the stable-rotating-wave region (green regime in Fig. 14), some of the spiral cores show a tendency to break and recombine in a very short interval of time. This instability of the core does not affect the long-term stability and dynamics of the spiral; the spiral core quickly reassembles and continues to rotate (see Fig. 15(d)), and remains as a localized, isolated event at the center; the regions in which we observe this are marked by magenta bubbles in Fig. 14. In addition to the states mentioned above, we find unstable rotating states, for parameters denoted by blue bubbles, and unstable broken spiral waves, for parameters marked as green stars, in the phase diagram Fig. 14. These states last only for a short duration and quickly move away from the domain. [Fig. 15 is the counterpart of Fig. 7 for the HRD model.]
of the time series of the transmembrane potential in the unstable regions (cf. Fig. 8 for the 2D HRD model).

Fig. 16. The dominant frequencies that we obtain from power spectra for all our parameter values (plots versus \( N_{G_{CaL}} \) for all the parameter values; \( N_{G_{Kr}} = 1, 3, 7 \) and \( N_{G_{CaL}} = 1, 3, 9 \)). In the region where stable rotating waves exist, we see a prominent frequency in the spectrum. If the waves are unstable and disappear, it is not possible to identify a major frequency; the missing parameter values in this plot correspond to these unstable regions (cf. Fig. 8 for the 2D HRD model).

the frequency increases as we move to high values of \( G_{Kr} \); by contrast, \( G_{CaL} \) has a very mild effect on the dominant frequency. A few examples of wave dynamics, along with the corresponding power spectra and the time series of \( V \), are shown in the Supplementary Material [29].

4.2. 3D results

We now present our results from our DNSs of the TP06 model in a realistic, human-heart geometry that is reconstructed from DTMRI data (see Fig. 4).

In the normal situation, a plane wave passes from one end to the other end of this geometry (without any obstacle). Fig. 17 shows different stages of such a passing of a plane wave through the human-ventricular geometry. The upper panels show the initial stages; and the lower panels show the final stages when the wave finally moves away from the domain.

We follow the same procedure that we have used above for the HRD canine-ventricular model to initiate a scroll wave. Then, as in the HRD case, we vary the parameters \( G_{Kr} \) and \( G_{CaL} \) simultaneously and investigate the effects of these changes on the scroll-wave development. We record this development for 2.5 s in real time.

With the original parameters of the TP06 model, we see that the scroll wave rotates without breaking; and it is stable for a long time after its initiation. The dynamics of such a scroll, for the parameters \( G_{Kr} = 1 \) and \( G_{CaL} = 1 \), is shown in Fig. 19a.

With a decrease of \( G_{CaL} \) by factors of 3, 5, and 7, the dynamics we observe is the same as above; a stable rotating wave is formed and it continues to rotate, with a small amount of meandering, but it does not break. We then increase \( G_{Kr} \) by factors of 3, 5, 7, and 9, while simultaneously varying \( G_{CaL} \), as above. Thus, we examine \( 5 \times 4 = 20 \) parameter sets. We present the general results of the scroll-wave behaviors in the phase diagram (or stability diagram) of Fig. 18. We observe that, as we move to the right and upper region of this phase diagram, the scroll waves tend to break up. This region corresponds to \( N_{G_{Kr}} = 7, 9 \) and \( N_{G_{CaL}} = 3, 5 \) and 7, and also \( N_{G_{Kr}} = 5, N_{G_{CaL}} = 7 \) and \( N_{G_{Kr}} = 3, N_{G_{CaL}} = 5 \). (This is the counterpart of Fig. 11 for the 3D HRD model.)

Fig. 19 shows the scroll wave’s time development for each type of dynamics that we observe: (a) shows a stable rotating scroll-wave; and (b) depicts a scroll wave breaking up and forming a chaotic state (\( N_{G_{Kr}} = 9, N_{G_{CaL}} = 5 \)). We observe another interesting case in the scroll-wave rotating-meandering region, for \( G_{Kr} = 3 \) and \( G_{CaL} = 7 \): the wave meanders widely throughout the simulation geometry; this...
In our HRD model simulations, the scroll-wave break-up that we observe, without varying the values of \( G_K \) increases and \( G_{CaL} \) decreases. We have observed this qualitative feature in the two distinctly different models, for two different mammalian species, namely the canine-ventricular HRD model and human-ventricular TP06 model [36].

The precise forms of the spiral or scroll wave that are formed in the simulation domain, in the absence of any parameter variation, are, of course, model specific. Hence, changes in these waves and their dynamics, as a function of model parameters, are also model specific. Nevertheless, there is a qualitative similarity in the transitions from one phase to another, the phase diagrams that we have presented for both HRD and TP06 models in both 2D and 3D. Details differ, of course, as we can see by comparing the phase diagrams for these models carefully. Such a detailed comparison between wave dynamics in these different mammalian models has not been attempted hitherto.

In our HRD model simulations, the scroll-wave break-up that we observe, without varying the values of \( G_K \) and \( G_{CaL} \) undergoes a transition to a stable meandering wave, without break up, as we vary the parameters. In the TP06 model simulations, we observe the reverse phenomenon, i.e., a transition occurs, from the stable rotating state, in the initial parameter region, to a broken-scroll or chaotic state (even when we vary the values of \( G_K \) increases and \( G_{CaL} \) decreases. However, this occurrence is not stable, and eventually breaks up (RB). All phases are stable states here, i.e., the waves do not disappear from the medium (cf. Fig. 11 for the 3D HRD model).

is shown in Fig. 19(c); (d) provides an example of the cases \( N_{GKr} = 5N_{GCaL} = 3,5 \), where the scroll-wave rotates for a long time, and finally breaks up after \( \approx 2s \). [This is the counterpart of Fig. 10 for the 3D HRD model.]

By comparing these figures with their 3D-HRD-model counterparts, we can see that scroll-wave dynamics for the TP06 model is quite different from that in the HRD model (for corresponding parameter regions).

4.2.1. Dominant frequencies

In this Subsection we examine the dominant frequencies of the scroll waves formed in the anatomical heart geometry by using the time series of the transmembrane potential from a few different sites of the domain.

Here also, as in the 2D case, the dominant frequency of the scroll wave increases with an increase in \( G_K \); but this frequency is not affected significantly by the variation of \( G_{CaL} \). This is shown in Fig. 20, where we show the dominant frequencies of the scrolls for the whole parameter space that we explore. A few examples of scroll-wave dynamics, along with the corresponding power spectra and the time series of \( V \) for the 3D TP06 model, are given in the Supplementary Material [29].

![Figure 18](image1.png)

**Fig. 18.** The phase diagram for scroll-wave dynamics in 3D TP06 model in a realistic, human-heart geometry. This diagram uses markers in the \( N_{GKr} - N_{GCaL} \) parameter space. In general, a bubble indicates a rotating scroll wave and a star indicates scroll break-up. The red circles indicate stable rotation (R) and the magenta circle indicates a highly meandering rotating scroll (RM). The blue stars indicate a rotating scroll, which breaks up eventually (RB); and the black stars indicates stable break-up (B). All phases are stable states here, i.e., the waves do not disappear from the medium (cf. Fig. 11 for the 3D HRD model).

![Figure 19](image2.png)

**Fig. 19.** Representative figures showing each of the four types of scroll wave dynamics we observe in our system. The panels from left to right show, via two-level isocontour plots of \( V \), the spatiotemporal evolution of the scrolls. (a) A stable rotating state where the scroll wave rotates and slightly meanders, without breaking up (\( N_{GKr} = 5, N_{GCaL} = 1 \)). (b) A stable chaotic state where a scroll wave breaks up and the broken scrolls spread throughout the medium (\( N_{GKr} = 7, N_{GCaL} = 3 \)). (c) A scroll wave meandering throughout the medium (\( N_{GKr} = 3, N_{GCaL} = 7 \)). (d) A scroll wave rotates for a long while and eventually breaks up (\( N_{GKr} = 3, N_{GCaL} = 3 \)); (cf. Fig. 10 for the 3D HRD model). For the complete spatiotemporal evolution see the Videos S23, S24, S25 and S26 in the Supplementary Material [29].

![Figure 20](image3.png)

**Fig. 20.** The dominant frequencies we observe for all the parameter values; \( N_{GKr} = 1, 2, 9 \) and \( N_{GCaL} = 1, 2, 7 \). For stable states, the frequency increases with \( G_K \); the variation with \( G_{CaL} \) is either small or nonexistent.

5. Conclusions

We have conducted extensive, *in silico* studies of the direct effects of two major, ion-channel conductances on spiral- and scroll-wave dynamics in idealized (2D) and anatomically detailed (3D) geometries for the canine- and human-ventricular models (HRD and TP06, respectively). We find that \( I_{Kr} \) and \( I_{CaL} \) are two important currents that determine the characters of spiral- and scroll-wave dynamics, namely, rotation, meandering, or the break-up of these waves in the ventricles. The effects of changes in these currents, on such wave dynamics, are most clearly visible when they are changed together, rather than individually (specifically, when \( G_K \) increases and \( G_{CaL} \) decreases). We have observed this qualitative feature in the two distinctly different models, for two different mammalian species, namely the canine-ventricular HRD model and human-ventricular TP06 model [36].
though we vary parameters over a range that is similar to the one we use in our HRD-model simulations). In both these models, the combination of the ion-channel conductances, for $I_{Kr}$ and $I_{Ca,L}$, plays a crucial role in determining the nature of scroll-wave dynamics. Recall that the parameter region we have explored is devoid of other commonly observed mechanisms that lead to uncontrolled scroll-wave behavior, such as a sharp APD restitution curve, early after depolarizations, and delayed after depolarizations [37–39].

There is no consensus on whether the geometric details of the heart itself affect scroll-wave dynamics or not; and, if they do, to what extent and in which way[40]. We have observed, in our simulations with anatomically realistic geometries and fiber-orientation details, that the long-term effect of the geometry, without abnormal inhomogeneities or other variations, on scroll-wave dynamics is negligible. The role that the geometry itself plays here is to trap the re-entrant waves, preventing them from decaying at the boundaries and thus stabilizing them. The primary determining factors, which affect wave dynamics and transitions from one sort of dynamics to another, are the conductances that govern the values of $G_{Kr}$ and $G_{Ca,L}$.

We conclude that the chaotic dynamics of scroll waves is not only produced by the common causes like a sharp APDR, EADs, and DADs, but also by a combined variation of the rapid-rectifier and calcium currents. $I_{Kr}$ and $I_{Ca,L}$; these play a crucial role in determining the dynamics of spiral and scroll waves in the two mammalian-heart models that we have studied. Our detailed description of the dependence of spiral- and scroll-wave dynamics on changes in these currents should provide insights into an understanding of the effects of drugs that target these current channels.

We mention some limitations of our study. We have used a monodomain model for the cardiac tissue equations in our study; bidomain models are more realistic than monodomain ones; however, a recent study [41] has shown that the latter are adequate when currents are low, as in our study. To impose boundary conditions we have used a phase-field approach [42]; this can also be done with a finite-element model. The dynamics of scroll waves is affected by a many more parameters than the two we study in detail. We have chosen these parameters for the reasons mentioned in this paper. A comprehensive study, including the simultaneous effects of change in more than two parameters, is computationally very expensive.

CRediT authorship contribution statement

K.V. Rajany: Software, Formal analysis, Visualization, Writing – original draft. Alok Ranjan Nayak: Conceptualization, Methodology. Rupamanjari Majumder: Data curation, Investigation, Validation. Rahul Pandit: Resources, Supervision, Validation, Writing – review & Editing, Funding acquisition.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

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