

In Silico Analysis of the Effects of Fibroblasts Coupling to Atrial Myocytes under Conditions of Atrial Fibrillation Remodeling

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Abstract

Under chronic atrial fibrillation (cAF), cardiac tissue experiences electrophysiological and structural remodeling, including fibrosis. Fibrosis in the atrial tissue has an important impact on myocyte action potential (AP) and its propagation. The objective of this work is to explore fibroblasts characteristics and their impact on myocytes AP. Human atrial myocyte and fibroblast electrophysiology is simulated using mathematical models proposed by Koivumaki et al., including healthy and cAF remodeled myocytes. Cellular simulations were run using three different resting membrane potentials for the fibroblasts (RMPf) and different numbers of coupled fibroblasts to a single myocyte in 0D and with different fibroblast densities in 1D. Results from single cell simulations show that myocyte's AP duration (APD) depends on both RMPf and the number of coupled fibroblasts. In 1D simulation, conduction velocity (CV) depends on RMPf and density along the strand. In conclusion, fibroblasts RMPf significantly alters myocyte AP and its propagation along cardiac tissue. Myocytes affected by cAF remodeling seem to be less sensitive to changes in fibroblasts RMPf.

1. Introduction

Atrial fibrillation (AF) is the most common cardiac arrhythmia. Under chronic atrial fibrillation (cAF) cardiac tissue experiences electrophysiological and structural remodeling. Fibroblasts density increases during AF and has an important role in the generation and maintenance of atrial reentries. Different studies suggest that fibroblasts present in the atrial substrate are electrically coupled to myocytes [1], [2]. Recent experimental studies have shown that fibroblasts express sodium channels allowing an inward current (I_{Na}) and their coupling with myocytes alter the AP.[3–5].

Computational simulations of cellular electrophysiology gives the opportunity to study the role that fibroblasts play in the atrial substrate. It also gives the flexibility to modify cellular electrophysiology and vary different characteristics in order to analyze the

impact of the electrical coupling with normal and electrical remodeled myocytes under cAF.

In this study, fibroblast sodium current will be updated using recent experimental data, RMPf will be varied to analyze its effects on the myocyte AP. This study also analyzes the different effects of fibroblast-myocyte coupling under normal and cAF conditions in structurally and electrophysiologically remodeled 1D atrial tissue.

2. Materials and methods

2.1. Action potential models

In this study we used human models of the atrial myocytes and fibroblasts from Koivumaki et al. [5], [6]. In the fibroblast model the sodium current was updated according to recent experimental results [4]. The electrical remodeling was introduced modifying ion channel conductances for L-type Ca^{2+} current (I_{CaL}), transient outward K^+ current (I_{to}), sustained outward K^+ current (I_{sus}), time independent K^+ current (I_{K1}), ryanodine receptors (RyR) and Na^+/Ca^{2+} exchanger (NCX), sarcoplasmic reticulum Ca^{2+} ATPase (SERCA) pump and parameters of time constants for I_{CaL} , I_{to} and I_{Na} , as described in Koivumaki et al.[7].

2.2. Cellular simulations (0D)

To compute the membrane potential in a cell equation (1) will be used:

$$C_m \frac{dV}{dt} + I_{ion} + \sum_{k=1}^n G_{gap} (V_f - V) = 0 \quad (1)$$

Where C_m is the membrane capacitance, I_{ion} is the ionic current flowing through ionic channels, V is the membrane potential in the myocyte, n is the number of attached fibroblasts to one myocyte and G_{gap} is the coupling conductance between myocyte and fibroblast. G_{gap} was set to 0.5 nS [1]. Additionally, the number of attached fibroblasts to a single myocyte varied from 0 to

9 and also the resting membrane potential of the fibroblast (RMPf) was set to different values (-30, -45, and -65 mV) [8]. APDs were measured at 90% of repolarization (APD₉₀).

2.3. Fiber simulations (1D)

To simulate the electrical propagation of the action potential along a strand, the monodomain approximation diffusion-reaction equation (2) was used.

$$\nabla(D\nabla V) = C_m \frac{dV}{dt} + I_{ion} \quad (2)$$

Where D is the diffusion coefficient, C_m is the membrane capacitance and I_{ion} stands for ionic currents through the membrane. Fibroblasts were randomly distributed along the strand with two different densities (10% and 20%) and three different values for the RMPf (-30, -45, and -65 mV). Diffusion coefficient was calculated considering geometrical dimensions from the cell, gap junction conductance, and intracellular medium conductance which was adjusted to achieve physiological conduction velocities (CV) along a strand of myocytes [9].

Depolarization upstroke was measured in a fibroblast near the centre of the strand and in one myocyte before and after the fibroblast to see the effect on the myocyte

depolarization upstroke as shown in Figure.

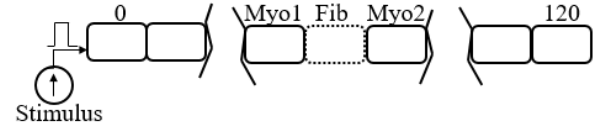


Figure 1. Strand configuration indicating the elements where measurements were taken (Myo1, Fib, and Myo2).

2.4. Simulation protocols and numerical methods

In 0D simulations the number of attached fibroblasts to a single myocyte was varied, as well as the value of RMPf. The applied stimuli had an amplitude of twice the threshold value. For 1D simulations, monodomain approach was solved using FDM with a space discretization of 100 μm for myocytes and 10 μm for fibroblasts and a time step of 10 μs. The first element was stimulated with an amplitude of twice its threshold. Both simulations reached steady state with a BCL of 1s before measurements were taken.

3. Results and discussion

Fibroblast coupling is expected to modulate myocyte AP characteristics. Results from 0D simulations are depicted in Figure 2, which shows how RMPf and the

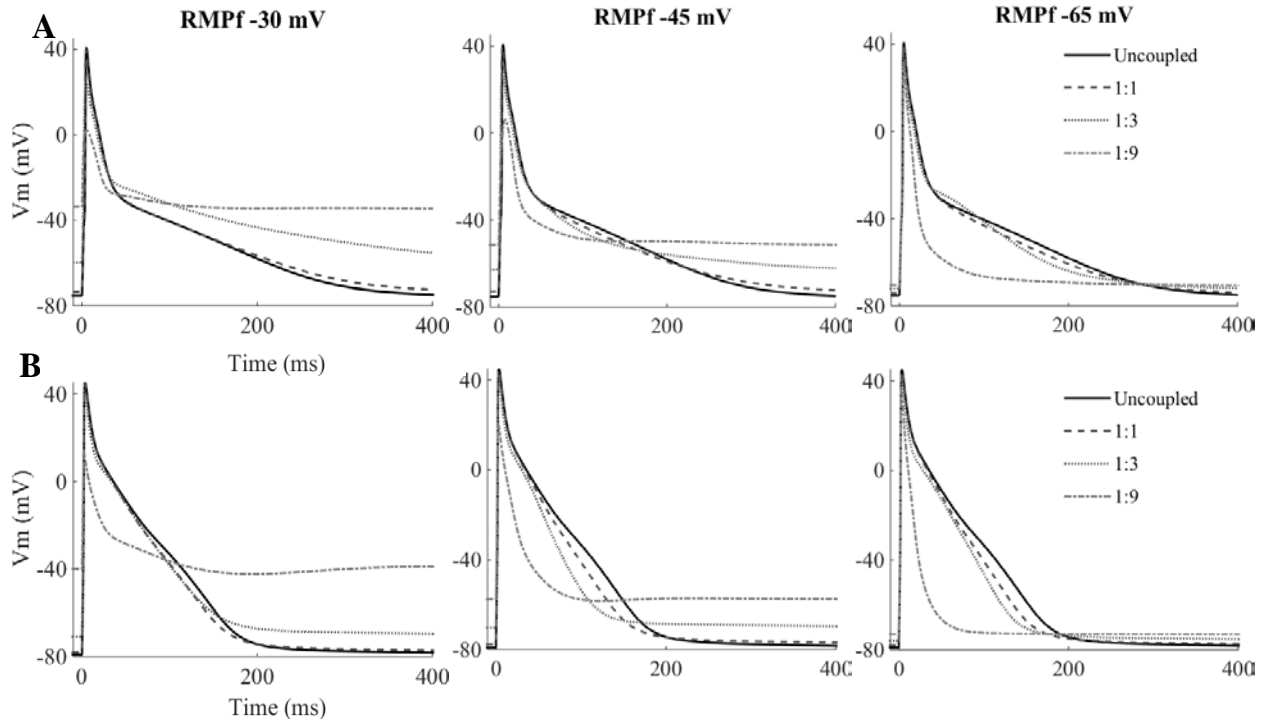


Figure 2. Simulations of isolated myocyte with a different number of attached fibroblasts and different RMPf, from left to right -30, -45, -65 mV, respectively. Panel A represents APs for myocyte under normal conditions (nSR) and panel B APs for myocytes under cAF remodeling.

number of coupled fibroblasts to a single myocyte have an influence on APD₉₀ and resting membrane potential (RMP) of the myocyte in normal nSR conditions (panel A) and under cAF remodeling conditions (panel B).

Uncoupled myocyte APD₉₀ is 233.2 ms and RMP is -74.8 mV under nSR (control values). However, when coupled with 1 or 3 fibroblasts with RMPf of -65 mV (panel A right graph), APD₉₀ is reduced to 91.68% and 78.04% of control and myocyte RMP is reduced to 99.18% and 96.50% of control. When RMPf is -45 mV (panel A middle graph), APD₉₀ is reduced to 94.43% and 70.20% of control value and the myocyte RMP is 97.36% and 83.98% of control. In both cases myocyte APD₉₀ is shortened and myocyte RMP is more depolarized when RMPf is less negative and coupled to 3 fibroblasts. Additionally, when RMPf is -30 mV (panel A left graph) and coupled to 1 or 3 fibroblasts, myocyte APD₉₀ was longer (102% and 137%, respectively) and their RMP is 98.15% and 80.12% of control. When 9 fibroblasts are coupled APD₉₀ is reduced to 30% of control approximately with all RMPf values, also myocyte RMP is significantly depolarized when RMPf is less negative.

Under cAF remodeling conditions (panel B), uncoupled myocyte APD₉₀ is 168 ms and RMP is -79.04 mV (control values) due to electrical remodeling. For 1 or 3 fibroblasts coupled to a single myocyte and RMPf -65 mV (panel B right graph), APD₉₀ is 90% and myocytes RMP 98.78% of control. However when coupled with 1 or 3 fibroblasts with RMPf of -45 mV (panel B middle graph), APD₉₀ is 86.82% and 66.75% of control with myocyte RMP of 98.14% and 88.86% of control. It is important to highlight that when coupled to 1 or 3 fibroblasts and RMPf of -30 mV (panel B left graph), all APD₉₀ are shortened. Furthermore, coupling to 9 fibroblasts results in an APD₉₀ and myocytes RMP reduction of 50% of control. Electrically remodeled

myocytes are less sensitive to the effect of fibroblast coupling and the variation of the RMPf.

As observed in 0D simulations, fibroblasts coupling has an important effect on myocytes AP. In 1D simulations, CV was analyzed to study fibroblasts characteristics in a tissue strand. CV in physiological conditions was 70 cm/s and in cAF CV was reduced to 68.6 cm/s.

Error! Reference source not found. shows the influence of fibroblast coupling on the upstroke of the AP. Values obtained for dV/dt_{max} during the depolarization upstroke of myocytes were 103 mV/s and 100 mV/s in nSR and cAF, respectively (control values) in the absence of fibrosis.

In nSR dV/dt_{max} of myocytes located before the fibroblast (Myo1 in Figure 1) are reduced depending on the RMPf and the fibrosis density in the strand. For 10% density and RMPf of -30 mV (panel A, left graph) dV/dt_{max} is reduced to 57% of control value while for the myocyte after the fibroblasts (Myo2 in Figure 1) dV/dt_{max} is 47% of control. With RMPf of -45 mV (panel C, left graph) dV/dt_{max} is 65% of control for Myo1 while for Myo2 the value is 53% of control. When fibroblast density is increased to 20% dV/dt_{max} with -30 mV (panel A, second graph) in Myo1 is 60% and in Myo2 is 37% of control, and with RMPf of -45 mV (panel B, second graph) the first myocyte presents a dV/dt_{max} of 25% and the second myocyte 34% of control.

CV is also reduced with increasing fibroblast density along the strand and depending on RMPf. Results from simulations in nSR shows that with densities of 10% with all three RMPf, CV is reduced (to 87%, 85%, and 63% of control value without fibrosis, respectively). When fibrosis density increases to 20% CV does not follow the same reduction trend. While with -30 mV RMPf, CV is reduced to 63% of control value, with -45 mV and -65

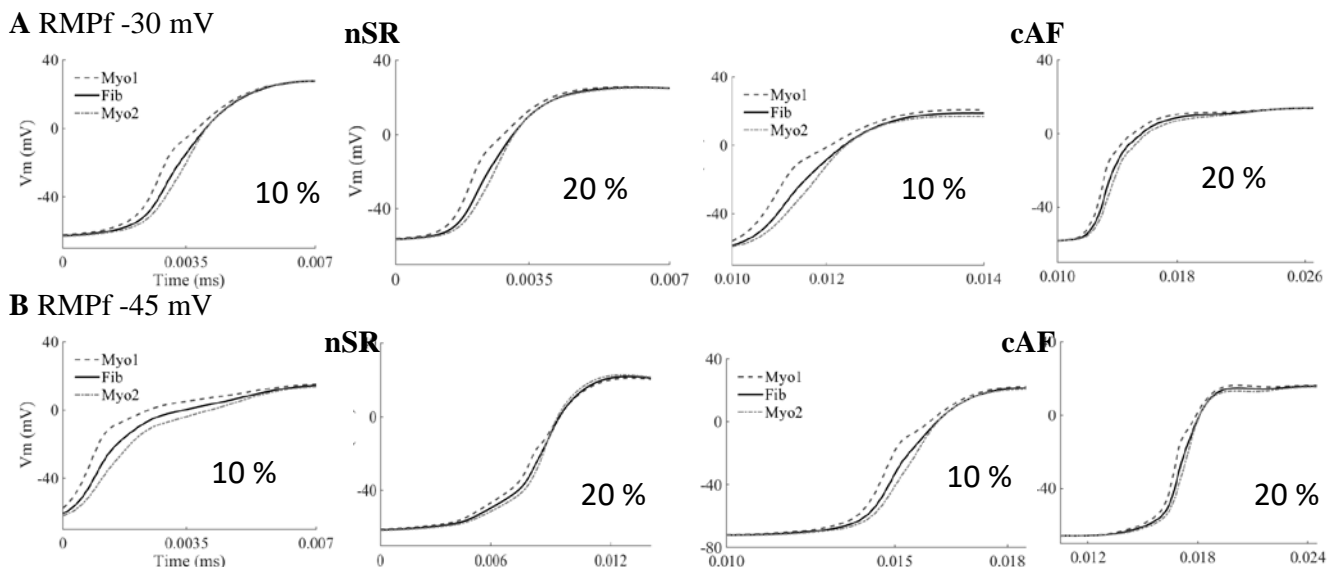


Figure 1. Depolarization upstroke for Myo 1, Fibroblast and Myo2 in 1D simulated strand for different conditions. Panel A) 10% and 20% fibrosis density with a RMPf of -30 mV and nSR. Panel B) 10% and 20% fibrosis density with a RMPf of -30 mV and cAF. Panel C) 10% and 20% density with RMPf of -45 mV and nSR. Panel D) 10% and 20% fibrosis density with a RMPf of -45 mV and cAF.

mV CV is only reduced to 74% and 75% of control value.

When electrical remodeling is introduced to simulate cAF conditions, values of dV/dt_{max} in the upstroke of the depolarization phase are also affected. For density of 10% and RMPf of -30 mV (panel B left graph), dV/dt_{max} for Myo1 is 66% of control and with RMPf of -45 mV (panel D left graph) is 43% of control, values for Myo2 are 51% and 28% of control respectively. When the density of fibroblasts is 20% and RMPf of -30 mV for Myo1 (panel B second graph) dV/dt_{max} is 58% and with a RMPf of -45 mV for Myo1 (panel D second graph) 54% of control, for Myo2 dV/dt_{max} values are 43% and 41% of control, respectively.

CV in cAF for 10% density and with all three RMPf shows the same percentage of reduction as in nSR. When fibroblast density increases, CV decreases as RMPf is depolarized (74%, 71%, and 67% of control, respectively). Additionally, RMP of the myocyte is not significantly affected.

Our results in 0D and 1D show that fibroblasts play an important role in both physiological nSR and in cAF conditions. They modify atrial substrate, which defines tissue properties for the action potential to propagate. Results obtained are in accordance with experimental results that show a decay in CV when density of fibroblast increases [10], [11]. It is interesting to notice that during cAF, CV seems to be less sensitive to present variability with RMPf.

4. Conclusion

In conclusion, fibroblasts effects on the myocyte AP depends on the value of their RMPf and the number of fibroblasts coupled to a myocyte as observed in 0D simulations. In 1D, fibroblast density and RMPf alters the action potential depolarization upstroke which modifies the CV along a tissue strand. Fibroblasts and their RMPf significantly alter myocyte AP and its propagation along cardiac tissue. Myocytes affected by cAF remodeling seem to be less sensitive to changes in fibroblasts RMPf.

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