A Protocol for Dual Calcium-Voltage Optical Mapping in Murine Sinoatrial Preparation With Optogenetic Pacing

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Among the animal models for studying the molecular basis of atrial and sinoatrial node (SAN) biology and disease, the mouse is a widely used species due to its feasibility for genetic modifications in genes encoding ion channels or calcium handling and signaling proteins in the heart. It is therefore highly valuable to develop robust methodologies for studying SAN and atrial electrophysiological function in this species. Here, we describe a protocol for performing dual calcium-voltage optical mapping on mouse sinoatrial preparation (SAP), in combination with an optogenetic approach, for studying SAP membrane potential, intracellular Ca\textsuperscript{2+} transients, and pacemaker activity. The protocol includes the details for preparing the intact SAP, robust tissue dual-dye loading, light-programmed pacing, and high-resolution optical mapping. Our protocol provides an example of use of the combination of optogenetic and optical mapping techniques for investigating SAP membrane potential and intracellular Ca\textsuperscript{2+} transients and pacemaker activity with high temporal and spatial resolution in specific cardiac tissues. Thus, our protocol provides a useful tool for studying SAP physiology and pathophysiology in mice.

Keywords: sinoatrial preparation, optogenetic pacing, optical mapping, murine heart, Ca\textsuperscript{2+} transient

INTRODUCTION

The heartbeat begins in the sinoatrial node (SAN) arising in a subset of nodal cells that display a spontaneous diastolic depolarization (Bleeker et al., 1980). Unique to such automaticity of the SAN, pacemaker cells possess “membrane clock” contributions by several membrane currents including the funny current, $I_f$, carried by hyperpolarization-activated cyclic nucleotide-gated (HCN) channels. $I_f$ is initiated specifically at the initial phase of the diastolic depolarization, following the deactivation of outward delayed rectifier K$^+$ current, and triggering the activation of inward currents. Inward currents include Na$^+$-dependent background current; the T- and L-type Ca$^{2+}$ currents, $I_{CaT}$ and $I_{CaL}$; and possibly, sustained inward current, $I_{st}$ (Lei et al., 2018). However, more recently, intracellular signaling involving the sarcoplasmic reticulum (SR) Ca$^{2+}$ stores and the cellular cAMP levels and consequent phosphorylation of their signaling proteins have also been implicated in a “Ca$^{2+}$ clock” contributing to diastolic depolarization (Lei et al., 2018). Whereby, spontaneous RyR2-mediated Ca$^{2+}$ release may enhance electrogenic effects of the Na$^+$-Ca$^{2+}$ exchanger.
in late diastolic depolarization (Lei et al., 2018). These SR calcium release events may be triggered by calcium-induced calcium release following the activation of Ca\textsuperscript{v,1.3} calcium channels (Torrente et al., 2016). The relative importance of both the “membrane clock” and “Ca\textsuperscript{2+} clock” in pacemaker function remains controversial.

On the other hand, sinus node dysfunction (SND) associated with abnormal impulse formation or propagation in the SAN affects ≈1 in 600 cardiac patients aged >65 years and is responsible for ≈50% of the 1 million permanent pacemaker implants per year worldwide (de Marneffe et al., 1993). Although SND occurs most commonly in elderly patients in the absence of clinically apparent accompanying cardiac disease (de Marneffe et al., 1993), its pathogenesis is unclear. The development of novel pharmacological therapies to cure these patients relies on the thorough understanding of both normal physiology and pathophysiology of the SAN and atria.

Among the animal models for studying the basic molecular mechanisms of the SAN and atrial biology and disease, mice are widely used due to their feasibility for modifications in the expression of different genes that encode ion channels or calcium handling proteins. It is therefore highly valuable to develop robust methodologies and techniques for studying SAN and atrial electrophysiological properties.

Recent advances in the technique of optogenetics provide an unprecedented opportunity for defining specific cell-type function in a highly complex multicellular system. The technique has enabled control over the activity of one specific cell type while leaving others unaltered, with high temporal resolution and cellular precision. Since the first demonstrations of utility in mammalian neurons in 2005 (Boyden et al., 2005), optogenetics has spurred immense research activity in neuroscience and has extended to the cardiac field over the past few years (Entcheva, 2013; Nussinovitch and Gepstein, 2015; Boyle et al., 2018). We recently deployed a cell-type specific optogenetic approach to study the properties of Pnmt\textsuperscript{+} cell-derived cardiomyocytes in the murine heart (Wang et al., 2017). Additionally, optical mapping using fluorescent dyes provides exciting opportunities for high spatio-temporal study of cellular electrophysiological events, including Ca\textsuperscript{2+} dynamics of the heart (Wang et al., 2017; Wen et al., 2018). The combination of these two technologies therefore allows development of unprecedented platforms for studying physiological events (e.g., voltage and Ca\textsuperscript{2+} signals) in a selective manner at high spatio-temporal resolution (O‘Shea et al., 2019b).

In this manuscript, we describe a detailed protocol for the use of optogenetic and optical mapping techniques for investigating membrane potential and intracellular Ca\textsuperscript{2+} transients and pacemaker activity with high temporal and spatial resolution in sinoatrial preparations (SAPs). The protocol can be a powerful tool for studying SAP physiology and pathophysiology using the mouse as a model system.

**METHODS AND MATERIALS**

**Animals**

Pnmt-cre mice used in this study are as described previously (Wang et al., 2017). Animals used were <6 months old. Pnmt-cre/ChR2 mice exhibit cell-type specific expression of ChR2 by crossing B6.Cg-Gt (ROSA)26Sor$_{tm1(CAG-COP4H134R/tdTomato)Hze}^{tm27.1}$ (Jackson Labs) with a Cre transgenic strain under the control of a Pnmt promoter (Wang et al., 2017). Channelrhodopsin 2 (ChR2) was specifically introduced into murine cells expressing the Phencylenedihydroxyamine n-methyltransferase (Pnmt) gene, which encodes for the enzyme responsible for conversion of noradrenaline to adrenaline. The murine model led to the identification of a distinctive class of Pnmt-expressing neuroendocrine cells and their descendants (i.e., Pnmt\textsuperscript{+} cell-derived cardiomyocytes) within the heart (Wang et al., 2017).

All procedures have been approved by Institutional Animals Ethics Committees at Southwest Medical University, Luzhou, China or Department of Pharmacology at University of Oxford, UK and the national guidelines under which the institution operates. All mice used in this study were maintained in a pathogen-free facility at Southwest Medical University or University of Oxford. Mice were given ad libitum access to food and water. The authors confirm that they have taken all steps to minimize the animals’ pain and suffering.

**Materials and Equipment**

Table 1 shows the details of materials and buffers.

| Table 1 | Details of materials and buffers. |
|---|---|
| Chemical and catalogue reference | Supplier |
| NaCl (SLBS2340V) | Sigma-Aldrich, St Louis, MO, USA |
| NaHCO\textsubscript{3} (SLBX3605) | Sigma-Aldrich, St Louis, MO, USA |
| NaHPO\textsubscript{4} (BCBW9042) | Sigma-Aldrich, St Louis, MO, USA |
| KCl (SLBSS0009) | Sigma-Aldrich, St Louis, MO, USA |
| MgCl\textsubscript{2} (SIGB8641V) | Sigma-Aldrich, St Louis, MO, USA |
| CaCl\textsubscript{2} (SLBK1794V) | Sigma-Aldrich, St Louis, MO, USA |
| Glucose (SLBTA4811V) | Sigma-Aldrich, St Louis, MO, USA |
| HEPES (W1122DC10) | Sigma-Aldrich, St Louis, MO, USA |
| 2,3-butanedione monoxime (BDM) (29297) | Sigma-Aldrich, St Louis, MO, USA |
| Blebbistatin (SLBV5564) | Sigma-Aldrich, St Louis, MO, USA |
| Voltage-sensitive dye RH237 (1971387) | Thermo Fisher Scientific, Waltham, MA, USA |
| Calcium indicator Rhod-2 AM (DSMO) (RNB77442) | Invitrogen, Carlsbad, CA, USA |
| Heparin Sodium (H10121200) | Sigma-Aldrich, St Louis, MO, USA |
| Avertin (2,2,2-tribromoethanol) | Sigma-Aldrich, St Louis, MO, USA |
| Pluronic F127 (1899021) | Sigma-Aldrich, St Louis, MO, USA |

**Figure 1** shows the optical imaging system for light stimulation of ChR2 light-sensitive channels and optical recording.

The tissues were paced through the activation of ChR2 light-sensitive channels. This was achieved by the delivery of 470 nm blue light pulses (13–14 ms pulse width) generated by OptoFlash (Cairn Research, Faversham, UK). Pulses were triggered by a 1,401 digitizer and Spike 2 software (Cambridge Electronic Design). Approximate blue light intensity was measured with a 818-ST2 Wand Detector connected to a 843-R power meter (both Newport Corporation, CA, USA), and we estimate an average irradiance in our experiments of 0.1–0.3 mW/mm\textsuperscript{2} based on an approximate distance of 1–2 cm between Sylgard and liquid light guide (Oriel instruments Model No. 77525).
Buffer Compositions

Physiological Salt Solution
Freshly prepared physiological salt solution (PSS) containing in mM: NaCl 119, NaHCO₃ 25, NaH₂PO₄ 1.0, KCl 4.7, MgCl₂ 1.2, CaCl₂ 1.35, and glucose 10; equilibrated with 95% O₂/5% CO₂ at 37°C, pH = 7.35.

Dye Loading Solution
Calcium indicator Rhod-2 AM (Invitrogen, Carlsbad, CA, USA) was dissolved in dimethyl sulfoxide (DMSO, Sigma-Aldrich, St. Louis, MO, USA) to prepare a stock concentration of 10 mg/ml (8.9 mM). The voltage indicator stock was prepared by dissolving Di-4-ANBDQPQ (University of Connecticut Health Center) in ethanol to obtain a stock concentration of 29 mM. To avoid repeated freezing and thawing, both stocks were stored in 50 μl aliquots at −20°C. For working concentrations of Rhod-2 AM dye loading solution, 15 μl Rhod-2 AM stock solution was mixed with 15 μl Pluronic F-127 (20% solution in DMSO, Invitrogen, Carlsbad, CA, USA) and then dissolved in 50 μl PSS at −20°C. For working concentrations of Rhod-2 AM dye loading solution, 15 μl Rhod-2 AM stock solution was mixed with 15 μl Pluronic F-127 (20% solution in DMSO, Invitrogen, Carlsbad, CA, USA) and then dissolved in 50 μl PSS at −20°C. For working concentrations of Rhod-2 AM dye loading solution, 15 μl Pluronic F-127 (20% solution in DMSO, Invitrogen, Carlsbad, CA, USA) and then dissolved in 50 μl PSS at −20°C.

Blebbistatin Physiological Salt Solution
Excitation-contraction uncoupler blebbistatin (Invitrogen, Carlsbad, CA, USA) was dissolved in in DMSO (Sigma-Aldrich, St. Louis, MO, USA) to make up 10 mM stock solution. To avoid repeated freezing and thawing, aliquots of 15 μl were stored at −20°C. A 10 μl of 10 mM stock blebbistatin solution was added to 15 ml PSS to make blebbistatin PSS solution (10 μM final concentration).

Preparation and Equipment

1. Dissection set-up includes stereomicroscope, dissection chamber filled with Sylgard gel, oxygenation device, LED light source, and surgical instruments.

2. The optical mapping system was set up as detailed below:

A custom-designed system equipped with an EMCCD camera (Evolve 128, Photometrics, Tucson, AZ, USA) was used (see Figure 1). Two green (to excite Ca²⁺ sensitive dye Rhod-2 AM; a peak wavelength of 530 nm) and two red (to excite the voltage sensitive dye Di-4-ANBDQPQ; a peak wavelength of 627 nm) LED lights (Cairn Research, Faversham, UK) were placed around the imaging chamber so as to uniformly illuminate the SAP. A multi-stream light switcher (Cairn Research, Faversham, UK) was set up to automatically switch between the red and green LED lights in accordance with the frame rate of the camera. Signal recording using this setup is in such a way that one frame data recorded will have only the red light turned on (thereby recording voltage signal alone) and the next frame data recorded will have only the green light turned on (thereby recording CaT signal alone), and the process is repeated until the desired number of frames are recorded.
PROTOCOL

Harvest the Mouse Heart and Prepare the Sinoatrial Preparation (30 min)
The steps for preparation of the SAP are outlined in Figure 2 and as described previously (Liu et al., 2007; Hao et al., 2011; Sharpe et al., 2016). The details are as follows:

1. Non-recovery terminal general anesthesia was induced by injection of an overdose of 1.2% Avertin solution (0.5–0.8 ml i.p., 2,2,2-tribromoethanol, Sigma-Aldrich, Poole, Dorset, UK), followed by intraperitoneal injection with heparin (200 units).
2. Following sacrifice via cervical dislocation, the heart was rapidly excised and placed in warmed (37°C) and oxygenated (95% O2 + 5% CO2 gas) PSS solution in a dissection chamber.
3. The heart was gently pressed with the tips of fingers to push the blood out of heart. Fresh oxygenated PSS was then added for further dissection.
4. SAP containing two atria and the SAN region was dissected after clearing off the lung, thymus, ventricles, and connective tissues (Figure 2B).
5. The right atrium was opened from atrioventricular (AV) junction to expose the intercaval region containing the SAN, following which left atrium was opened to expose the left atrial endocardium. As the pacemaker cells are widely distributed throughout the entire region located between the superior (SVC) and inferior vena cava (IVC) and between the crista terminalis and intra-atrial septum (Glukhov et al., 2010), the SAP contains the SAN and the two atria intact.
6. The SAP was pinned onto a small rectangular piece of Sylgard gel by using fine insect pins. All dissection steps were performed in a dissection chamber with oxygenated PSS solution under a stereomicroscope.

Dual Dye Loading of the Sinoatrial Preparation (25 min)
7. A 50 ml falcon tube was filled with 15 ml of the Rhod-2 AM dye loading solution.
8. The Sylgard gel (with SAP attached) was then dropped into the falcon tube.
9. The SAP-containing falcon tube was placed in a 36°C water bath. The solution was oxygenated constantly by gently bubbling with 95% O2/5% CO2 (Note: heavy bubbling can lead to the formation of foam due to the presence of Pluronic F-127 in the solution. This can lead to the suboptimal dye loading). The SAP was incubated in the dye loading solution for 15 min.
10. Dual dye loading of the SAP was performed by incubating the SAP in Rhod-2 AM dye loading solution as detailed above (points 7–9), following which the Sylgard gel with SAP attached was transferred into a new 50 ml falcon tube containing 10 μM Di-4-ANBDQPO dye loading solution. Incubation with Di-4-ANBDQPO was performed for 10 min at 36°C and constantly bubbled with 95% O2/5% CO2.

Optical Mapping (30 min)
11. The imaging chamber was filled with 15 ml of blebbistatin PSS solution and was placed directly under the camera. The recording chamber was heated up to 36°C by a heating plate.
12. The Sylgard (with SAP attached) was removed from the falcon tube and pinned down in the imaging chamber. The sample was placed between the two field-stimulation electrodes. (Note: Removing the SAP from the Sylgard and pinning down directly in the chamber cause further damage to the SAP. Therefore, pinning down the Sylgard with the intact
tissue is recommended). The blebbistatin PSS solution was bubbled in the imaging chamber with 95% O\textsubscript{2}/5% CO\textsubscript{2} gas.

13. Imaging was performed after cessation of motion by blebbistatin. Imaging used the EMCCD camera controlled by the acquisition software Metamorph (Molecular Devices). During image recordings, the excitation lights were turned on, and bubbling was temporarily stopped (to avoid distortion of the signals due to bubbles and motion of the solution). Since Pnmt\textsuperscript{+} cell-derived cardiomyocytes (PdCMs) express channelrhodopsin-2 (ChR2), they can be excited by exposure to 470 nm blue light. The stimulation protocol was then run, and baseline calcium transient or action potential recordings were taken. Recordings were taken again after stimulating the left atrium using blue light (pacing intervals: 150–160 ms, pulse duration: 13–14 ms, intensity: 1,992 mA). The frame rate of these recordings was 0.52 kHz.

14. For dual dye imaging, a multistream light switcher (Cairn Research, Faversham, Kent, UK) was set up to automatically switch between the red (627 nm) and green (530 nm) LED lights according to the frame rate of the camera. Signal recording using this setup was performed, so that one frame of the data was recorded under red light illumination (thereby recording $V_m$ signal alone), and the next frame was recorded under green light illumination (thereby recording CaT signal alone) and so forth until the desired number of frames was recorded. This resulted in an interlaced signal containing alternate frames of CaT and $V_m$ signals. The interlaced signal was separated into CaT and voltage signal using ImageJ and saved as separate files. Due to the signals being interposed, the frame rate recording was 0.26 kHz.

Analysis

15. The saved image files were loaded, processed, and analyzed using the optical mapping analysis software ElectroMap (developed by Dr. Pavlovic’s group, https://github.com/CXO531/ElectroMap; O’Shea et al., 2019a).

16. Images were pre-processed by applying a 4 × 4 pixel Gaussian spatial filter (standard deviation = 1.5), a Top-Hat baseline correction (kernel size = 100 ms), and a third order Savitzky-Golay temporal filter. A window size of 40 ms before transient peak to 100 ms after transient peak was used. Action potential duration (APD) and CaT duration (CaD) were measured at the desired repolarization/decay percentage at each pixel across the tissue, as measured from time of maximum upstroke velocity (maximum dF/dt).

17. For assessment of local conduction velocities, activation maps were generated using time of depolarization midpoint. Conduction velocity across the SAP was then quantified using a multi-vector polynomial method (Bayly et al., 1998) with a local window size of 5 × 5 pixels.

18. To further assess conduction across the entire SAP, we used activation curve analysis, where the percentage of tissue activated was plotted as a function of time (O’Shea et al., 2019a).

Representative Results

A typical example of an SAP Ca\textsuperscript{2+} transient activation map reconstructed from spontaneous sinus rhythm Ca\textsuperscript{2+} transients (by Rhod-2 AM) recorded with the optical mapping system is shown in Figure 3A for a SAP. The early activation point is located within the intercaval region near to where the SAN is located. However, as shown in Figure 3B, the calcium activation pattern and atrial conduction changed after stimulating with blue light (interval: 150 ms; duration: 13 ms; intensity: 1,992 mA; pulses per train: 50). The leading pacemaker site was shifted to the location of left atrium, suggesting that PdCMs were activated earlier than the SAN pacemaker cells. This is consistent with our previous report that PdCM cells predominantly localized to the left atrium and left ventricle, and the heart rate was controlled by stimulating PdCMs selectively with blue light (Wang et al., 2017). We hypothesize that PdCM cells overexpressing ChR2 may be a viable site for spatially-targeted optical pacing when sinus node pacemaker cells are dysfunctional. Further
investigation is required. Figure 4 shows representative $V_m$ and CaT maps and voltage and calcium traces from different regions of the SAP, illustrating the robustness of this preparation for murine research. We monitored the “rundown” of the $V_m$ and CaT signals for up to 4 h – the average signal “rundown” is less than 25%. Our experiment usually finished within 3 h from the beginning of Langendorff perfusion of the heart. We again observe augmentation of the activation pattern induced by optical pacing, shown by both the voltage (Figure 4, upper panel) and calcium (Figure 4, lower panel) activation maps. Activation maps were produced by mapping the midpoint time of the AP/CaT rise.

We found that optogenetic pacing of a SAP can shift the activation point to the left, creating a new leading pacemaker site or “neonode.” Activation patterns in the SAP were plotted using ElectroMap (O’Shea et al., 2019a), with the midpoint of the upstroke of the calcium transient used as a pixel-by-pixel readout of tissue activation. The earliest-activating pixel—discarding outliers—was designated the activation point for that tissue. The earliest activation point before pacing was always in the nodal region. Optogenetic pacing of PdCMs can move this activation point to the left (Figure 5), establishing a new leading pacemaker site toward the left atrium. This change in the location of the leading pacemaker is reversible upon cessation of pacing (Figures 5A,E). Left shift of the leading pacemaker occurred in four of seven SAPs (Figure 5D). One leftward deviation appears more upward relative to the SVC-IVC axis—this is due to contortion of the tissue. Although no leading pacemaker shift occurred in three SAPs, all seven SAPs were successfully paced by blue light: with a decrease in mean cycle length from 264.3 ± 42.7 ms before optogenetic pacing to 159.2 ± 1.4 ms during ($p < 0.02$). Mean cycle length after pacing was 281.9 ± 48.9 ms—not significantly different from before ($p = 0.63$). Therefore, we can always successfully and reversibly pace PdCMs using selective ChR2 overexpression. The variability in pacemaker shift observed (Figure 5D) is consistent with the variable distribution of murine PdCMs across both the SAN region and the left atrium (Ni et al., 2017; Wang et al., 2017). In summary, we report the possibility of using optogenetic pacing of PdCMs to establish novel left-shifted leading pacemaker sites in the murine heart.

To study the effects of optogenetic pacing on overall conduction, we utilized activation curve analysis where the percentage of tissue activated is plotted as a function of time (O’Shea et al., 2019a). Optogenetic pacing of PdCMs augments conduction in the SAP—producing a left shift of the activation curve (Figure 6A). This is most noticeable in the first few milliseconds of tissue activation (Figure 6A, inset). During optogenetic pacing, the rate of tissue activation is initially very rapid (Figure 6B, inset), otherwise, there are no major alterations to tissue activation rate (Figure 6B). Optogenetic pacing reduces the time to 5% tissue activation from 1.59 ± 0.24 to 0.88 ± 0.17 ms ($p = 0.03$),

**FIGURE 4** Simultaneous voltage (top panels) and Ca$^{2+}$ transient (bottom panels) activation maps reconstructed from spontaneous sinus rhythmic (left panels) and light paced (right panels) SAP. Maps were produced by mapping the midpoint time of the voltage and CaT rise with examples of voltage and Ca$^{2+}$ transient traces from indicated regions of the preparation (white and red cross).
the time to 10% tissue activation from 2.40 ± 0.25 to 1.46 ± 0.26 ms (p = 0.02), and the time to 20% tissue activation from 3.91 ± 0.52 to 2.27 ± 0.33 ms (p = 0.02) (Figure 6C), but at all points ≥30% tissue activation, there is no significant difference in activation time (Figures 6A, B).

We also wanted to explore whether the normal conduction pathways in the tissue remained intact during optogenetic pacing. Using ElectroMap, we computed local conduction velocities using the multi-vector method (Bayly et al., 1998) with a 5 × 5-pixel grid size. The magnitude of conduction velocity vectors (Figure 6D) did not change significantly with optogenetic pacing from 26.9 ± 3.3 before to 28.6 ± 2.5 cm/s during pacing (p = 0.34), and there were no major changes in the directionality of conduction vectors either (Figure 6E). In summary, we report the possibility of establishing a novel left-shifted leading pacemaker site, with only minimal, controlled disturbance to normal tissue activation rate, and little or no disturbance to the pathways along which conduction normally propagates.

FIGURE 5 | Mapping of activation in SAPs under sinus rhythm and under optogenetic pacing. (A) (i) Representative SAP—camera view. (ii) Activation map for this SAP before optogenetic pacing. (iii) Activation map during pacing (note formation of “neonode”, white asterisk). (iv) Activation map after pacing. Activation maps are calculated from Ca²⁺ transients. Fifty pacing spikes given, at 6.67 Hz. Contours drawn at 1/frame rate of the EMCCD. Scale bar, right. (B) 3-D graphical rendering of activation maps (i) before pacing, (ii) during pacing. SVC-IVC axis in red. Activation time on z-axis. (C) Representative SAP—microscope view. Pacemaker shift overlaid. SVC-IVC axis shown. Distances in mm. (D) SVC-IVC axis with relative pacemaker shifts plotted (n = 7). (E) Mean change in “shiftedness.” LA, left atrium; RA, right atrium; SAN, sinoatrial node; SVC, superior vena cava; IVC, inferior vena cava; SEP, septum; NeoN, neonode.
We found small alterations to calcium cycling in our tissues, but for the most part, there was no significant change in the characteristics of calcium transient rise time, calcium transient duration to 50% of the decay (CaD50), and calcium transient duration to 75% of the decay (CaD75) with optogenetic pacing. We measured the rise time, calcium transient duration to 50% of the decay (CaD50), and calcium transient duration to 75% of the decay (CaD75) as illustrated in Figure 7A. Rise time is a measure of upstroke duration, calculated between 10 and 90% of time to peak, which minimizes analytical uncertainty (Choi and Salama, 2000; Jaimes et al., 2016). We used a mixed-effects model, with Tukey’s post hoc tests where required, to examine whether there were any significant effects of optogenetic pacing. For rise time, there was a significant overall effect of optogenetic pacing \( (p < 0.0001) \). Post hoc testing revealed that there was no significant effect of pacing on rise time in the right atrium or in the nodal regions—but in the left atrium, rise time fell from 17.21 ± 0.94 ms before pacing to 15.24 ± 1.14 ms during pacing \( (p < 0.002) \) and rose again after pacing to 18.60 ± 1.00 ms \( (p < 0.002) \). In both regions, there was no significant difference between the before and after conditions (left atrium: \( p = 0.39 \); nodal region: \( p = 0.61 \))—so the reduction in rise time with optogenetic pacing is reversible upon cessation of pacing (Figure 7B). There was no overall significant effect of optogenetic pacing on either CaD50 \( (p = 0.15) \) or CaD75 \( (p = 0.34; \text{Figures 7C,D}) \). In summary, rise time is shorter in the left atrium and in the SAN region during optogenetic pacing, and this is a reversible change. Otherwise, calcium handling in the murine SAP appears to be only marginally affected.

DISCUSSION

The success in obtaining high quality \( V_m \) and Ca\(^{2+} \) mapping results in SAPs requires several important technical considerations, including high consumption of oxygen and high sensitivity to temperature and pH of the nodal tissue. Furthermore, the dense arrangement of nodal cells combined with the significant presence of connective tissue makes it hard for a Ca\(^{2+} \) dye to reach to the nodal and atrial cells. The critical steps include as follows:
**Tissue Preparation**

Intact SAPs were micro-dissected as described previously (Ju et al., 2015). The SAN region was identified from anatomical landmarks, including the superior vena cava, the crista terminalis, and the interatrial septum. The tissue needs to be oxygenated constantly. At this point, temperature is not critical.

**Dye Loading**

The $\text{Ca}^{2+}$-sensitive dye used to load the SAP tissue was Rhod-2 AM, a dye in the acetoxymethyl ester form. Esterification of the fluorescent molecule (Rhod-2 AM) results in an uncharged molecule that can permeate cell membranes and chelate cytosolic $\text{Ca}^{2+}$. A 100-fold increase in the molecule's fluorescence intensity results from $\text{Ca}^{2+}$ chelation (Choi and Salama, 2000). The voltage sensitive dye used was Di-4-ANBDQPQ, due to its favorable spectral properties for use with Rhod-2 AM. In our experiments, dye loading was carried out by incubating the tissue in a solution consisting of Pluronic F-127 dissolved in DMSO. Pluronic F-127 was incorporated into the dye loading solution to better disperse the dye into the physiological solution. Dye loading was observed to be generally better in younger mice than old, which may be related to connective tissue making up a smaller volume of the node.

The SAP’s repetitive contractions may have disrupted dye loading and/or aided the leaking of the dye out from the tissue, making it a challenge to optimize dye loading conditions. A fluorescent dye that diffuses across the cell membrane well can also exit from the cytoplasm at a high rate. Reagents like probenecid help prevent this to an extent. Combining these with optimized temperatures, timings and oxygen supply are necessary for effective loading of the tissue with dye.

**Optical Imaging**

During imaging, motion artifacts caused disruption in the measurement of optical signals. Hence, to reduce contraction of the tissue, blebbistatin was used. Blebbistatin is an inhibitor of the adenosine triphosphatases associated with class II isoforms of myosin (Allingham et al., 2005). This compound has been identified as an effective excitation-contraction uncoupler, and previous investigators have used it at several different concentrations (Fedorov et al., 2007). The use of blebbistatin at 10 μM appeared to effectively abolish any motion artifact due to contraction of the SAP without harming SAP.

**Pnmt$^+$ Cell-Derived Cardiomyocytes (PdCMs) and Biopacemaking Strategies**

The advent of cardiac optogenetics has facilitated the identification of a new cardiomyocyte subpopulation in the murine heart, phenylethanolamine-N-methyl transferase Pnmt$^+$-derived cardiomyocytes (PdCMs; Wang et al., 2017). Through selective
manipulation of these cells using optogenetics, our study demonstrated how optogenetics can be used to target specific cardiomyocyte subpopulations, overcoming many of the issues associated with traditional electrophysiological techniques. Selectively stimulating PdCMs with rhythmic blue (470 nm) light pulses was sufficient to maintain sinus rhythm in murine atria, supporting previous observations from whole heart studies (Wang et al., 2017). It has also been found previously that the left atrium alone – dissected out – can be paced by light (Wang et al., 2017), but a capacity to optogenetically control the SAP has not been tested before.

The observation that a “neonode” is typically set up toward the left is consistent with previous findings that PdCMs occupy a left-sided distribution in murine heart (Wang et al., 2017). We therefore assessed the effect of pacing on left atrial activation. Optogenetic pacing of PdCMs leads to alterations in the conduction pattern across murine atria. We found (Figure 5) that optogenetic pacing of Pnmt-cre/Ai27D SAPs can often set up new sites of earliest activation (“neonodes”) toward the left of the tissue. Our proof-of-principle studies clearly demonstrate feasibility of optogenetic control of heart rate via the specific cell population. Indeed, in four of seven SAPs, we observed “neonode” formation, with the leading pacemaker shifting toward the left atrium. Lack of leftward shift in three of seven preparations may be due to the heterogeneity in cell numbers and clustering of the Pnmt cells in the SAPs, thus altering sink-source properties of the myocardium. While the physiological consequences of the studies presented here (specifically using Pnmt cells) are not immediately clear, application of such approaches is of interest to the community and may lead to important findings on the consequences of the spatial shift in the leading pacemaker.

As recently reviewed by Boyle et al. (2018), initial work in this area assessed possibility to override the intrinsic pacing frequency of a beating heart using optical stimuli. Bruegmann et al. (2010) explored this question in a preclinical model by generating ChR2 transgenic mice. They showed in an open-chest configuration that pulses of light from a focused light source could be used to maintain ventricular activation at rates faster than intrinsic sinus rhythm. Light-induced wavefronts were initiated from different parts of the heart (e.g., atrial or ventricular sites). Remarkably, the authors found that 1:1 capture could be maintained when the illuminated area was small (0.05 mm², ~50 myocytes), suggesting that optogenetic pacing via a small population of cells is possible and robust. Several recent studies also provided evidence for the development of optogenetic defibrillating device for termination of tachyarrrhythmias (Nyns et al., 2016; Bruegmann et al., 2017). Bruegmann et al. (2017) reported optogenetic termination of atrial tachyarrrhythmia in intact hearts from transgenic as well as wild-type mice ex and in vivo. Thus, they suggested that their report could lay the foundation for the development of implantable devices for pain-free termination of AF. Nyns et al. (2016) demonstrated that forced expression of a light-gated depolarizing ion channel (ReaChR) in the adult rat heart allows contact- and shock-free termination of VTs through brief local illumination of the ventricular surface, i.e., without relying on conventional drugs, tissue ablation, or electroshocks. Both mono- and polymorphic VTs could be terminated in an effective and repetitive manner by a light-induced electrical current driven by natural electrochemical gradients, providing proof-of-concept for biological arrhythmia termination. Our study also provides proof-of-concept evidence supporting the development of light-based pacing devices targeted to a particular cellular population.

Limitations of the Protocol

There are a number of possible limitations in this protocol including: (1) possibility of injury during the dissection of the preparation; (2) effectiveness of dye loading may vary between preparations, thereby affecting the results; and (3) partial uncoupling (reduction in conductance) due to the use of blebbistatin, which could increase the regional differences in Ca²⁺ dynamics in SAPs.

Conclusions and Future Directions

Here, we have described a detailed protocol for dual optical mapping of murine SAPs with optogenetic pacing. Combination of high-resolution optical imaging with cell-specific optogenetic stimulation allows unique spatial control and study of electrophysiology of the SAP, not possible with traditional direct or field electrode techniques. In the present protocol, we have demonstrated this principle with PdCMs; however, the protocol can be valuable for any mouse models that cell specifically express opsins such as ChR2 within the sinoatrial region of the heart. Therefore, such an approach represents a powerful tool for enhanced electrophysiological understanding of supraventricular physiology and pathophysiology.

Future improvements may focus on exploring dual dye imaging and improved image processing software. Higher resolution and novel optical imaging modalities for 3D optical mapping are also important future directions of optical mapping.

ETHICS STATEMENT

Pnmt-Cre mice used in this study are as described in our previous paper (Wang et al., 2017). Animals used were <6 months old, 20–25 g. All procedures including animal subjects have been approved by Institutional Animals Ethics Committees at Southwest Medical University, Luzhou, China or Department of Pharmacology at University of Oxford and the national guidelines under which the institution operates. All mice used in this study were maintained in a pathogen-free facility at Southwest Medical University or University of Oxford. Mice were given ad libitum access to food and water. The authors confirm that they have taken all steps to minimize the animals’ pain and suffering. Our work complies with the journal policy and regulations.

AUTHOR CONTRIBUTIONS

RD, RM, AR, and SH performed the experiments and produced the data. CO’S, AR, and DP analysed the results.
AG-R, KD, XT, and KK assisted with data processing and manuscript preparation. ML and XO drafted the manuscript with the assistance of RM, DP, and AR. ML revised the manuscript with the assistance of CO’S, DP, RM, AR, and AG-R.

**FUNDING**

This study was supported by Medical Research Council (G10002647, G1002082, ML), British Heart Foundation (PG14/80/31106, PG/16/67/32340, PG/12/21/29473, PG/11/59/29004 ML; PG/17/55/33087, RG/17/15/33106, FS/19/16/34169, FS/19/12/34204 DP), BHF Centre of Research Excellence, Oxford (ML) grants, EPSRC (L016346 DP), National Natural Science Foundation of China (Nos. 81700308 and 31871181).

**REFERENCES**

Allingham, J. S., Smith, R., and Rayment, I. (2005). The structural basis of blebbistatin inhibition and specificity for myosin II. *Nat. Struct. Mol. Biol.* 12, 378–379. doi: 10.1038/nsmb908

Bayly, P. V., KenKnight, B. H., Rogers, J. M., Hildlsey, R. E., Ideker, R. E., and Smith, W. M. (1998). Estimation of conduction velocity vector fields from epicardial mapping data. *IEEE Trans. Biomed. Eng.* 45, 563–571. doi: 10.1109/10.668746

Bleeker, W. K., Mackay, A. J., Masson Pevet, M., Bouman, L. N., and Becker, A. E. (1980). Functional and morphological organization of the rabbit sinus node. *Circ. Res.* 46, 11–22. doi: 10.1161/01.RES.46.1.11

Boyd, E. S., Zhang, F., Bamberg, E., Nagel, G., and Deisseroth, K. (2005). Millisecond-timescale, genetically targeted optical control of neural activity. *Nat. Neurosci.* 8, 1263–1268. doi: 10.1038/nn1525

Boyle, P. M., Karathanos, T. V., and Trayanova, N. A. (2018). Cardiac optogenetics. *JACC Clin. Electrophysiol.* 4, 155–167. doi: 10.1016/j.jacep.2017.12.006

Bruegmann, T., Beiert, T., Vogt, C. C., Schrickel, J. W., and Sasse, P. (2017). Optogenetic termination of atrial fibrillation in mice. *Cardiovasc. Res.* 114, 713–723. doi: 10.1093/cvr/cvx250

Bruegmann, T., Malan, D., Hesse, M., Beiert, T., Fuegemann, C. J., Fleischmann, B. K., et al. (2010). Optogenetic control of heart muscle in vitro and in vivo. *Nat. Methods* 7, 897–900. doi: 10.1038/nmeth.1512

Choi, B. R., and Salama, G. (2000). Simultaneous maps of optical action potentials and calcium transients in Guinea-pig hearts: mechanisms underlying concordant alternans. *J. Physiol.* 529, 171–188. doi: 10.1111/j.1469-7793.2000.00171.x

de Marneffe, M., Greigoire, J. M., Waterschoot, P., and Kestemont, M. P. (1993). The sinus node function: normal and pathological. *Eur. Heart J.* 14, 649–654. doi: 10.1093/eurheartj/14.5.649

Entcheva, E. (2013). Cardiac optogenetics. *Am. J. Physiol. Heart Circ. Physiol.* 304, H1179–H1191. doi: 10.1152/ajplegacy.00432.2012

Fedorov, V. V., Lozinsky, I. T., Sosunov, E. A., Anyukhovsky, E. P., Rosen, M. R., Balke, C. W., et al. (2007). Application of blebbistatin as an excitation-contraction uncoupler for electrophysiologic study of rat and rabbit hearts. *Heart rhythm* 4, 619–626. doi: 10.1016/j.hrthm.2006.12.047

Glukhov, A. V., Fedorov, V. V., Anderson, M. E., Mohler, P. J., and Efimov, I. R. (2010). Functional anatomy of the murine sinus node: high-resolution optical mapping of ankyrin-B heterozygous mice. *Circ. Arrhythm. Electrophysiol.* 3, 897–900. doi: 10.1161/CIRCEP.110.960807

Jaimes, R. 3rd, Walton, R. D., Pasdois, P., Bernus, O., Efimov, I. R., and Kay, M. W. (2016). A technical review of optical mapping of intracellular calcium within myocardial tissue. *Am. J. Physiol. Heart Circ. Physiol.* 310, H1388–H1401. doi: 10.1152/ajplegacy.00865.2015

Ju, Y. K., Lee, B. H., Trajanovska, S., Hao, G., Allen, D. G., Lei, M., et al. (2015). The involvement of TRPc3 channels in sinoatrial arrhythmias. *Front. Physiol.* 6, 686. doi: 10.3389/fphy.2015.00686

Lei, M., Wu, L., Terrar, D. A., and Huang, C. L. H. (2018). Modernized classification of cardiac antiarrhythmic drugs. *Circulation* 138, 1879–1896. doi: 10.1161/CIRCULATIONAHA.118.035455

Liu, J., Dobrzynski, H., Yanni, J., Boyett, M. R., and Lei, M. (2007). Organisation of the mouse sinoatrial node: structure and expression of HCN channels. *Cardiovasc. Res.* 73, 729–738. doi: 10.1016/j.cardiores.2006.11.016

Ni, H., Wang, Y., Crawford, W., Zhang, S., Cheng, L., Zhang, H., et al. (2017). Three-dimensional image reconstruction of distribution of Pnm+ cell-derived cells in murine heart. *Sci. Data* 4, 170134.

Nussinovitch, U., and Gepstein, L. (2015). Optogenetics for in vivo cardiac pacing and resynchronization therapies. *Nat. Biotechnol.* 33, 750–754. doi: 10.1038/nbt.3268

Nyns, E. C. A., Kip, A., Bart, C. I., Plomp, J. J., Zeppenfeld, K., Schali, M. J., et al. (2016). Optogenetic termination of ventricular arrhythmias in the whole heart: towards biological cardiac rhythm management. *Eur. Heart J.* 38, 2132–2136. doi: 10.1093/eurheartj/ehw574

O’Shea, C., Holmes, A. P., Winter, J., Correa, J., Ou, X., Dong, R., et al. (2019). Cardiac Optogenetics and Optical Mapping – Overcoming Spectral Congestion in All-Optical Cardiac Electrophysiology. *Front. Physiol.* 10, 1–14.

O’Shea, C., Holmes, A. P., Yu, T. Y., Winter, J., Wells, S. P., Correa, J., et al. (2019a). ElectroMap: high-throughput open-source software for analysis and mapping of cardiac electrophysiology. *Sci. Rep.* 9, 1389. doi: 10.1038/s41598-018-38263-2

Sharpe, E. J., St Clair, J. R., and Proenza, C. (2016). Methods for the isolation, culture, and functional characterization of sinoatrial node myocytes from adult mice. *J. Vis. Exp.* 116:54555. doi: 10.3791/54555

Torrente, A. G., Barrere, C., Nargeot, J., Mangoni, M. E., Mesirca, P., Rizzetto, R., et al. (2016). L-type Cav1.3 channels regulate ryanodine receptor-dependent Ca2+ release during sino-atrial node pacemaker activity. *Cardiovasc. Res.* 109, 451–461. doi: 10.1093/cvr/cvw006

Wang, Y., Lin, W. K., Crawford, W., Ni, H., Bolton, E. L., Khan, H., et al. (2017). Optogenetic control of heart rhythm by selective stimulation of cardiomycocytes derived from Pnm+ cells in murine heart. *Sci. Rep.* 7, 40687. doi: 10.1038/s41598-017-18302-0

Wen, Q., Gandhi, K., Capel, R. A., Hao, G., O’Shea, C., Neagu, G., et al. (2018). Transverse cardiac slicing and optical imaging for analysis of transmural gradients in membrane potential and Ca2+ transients in murine heart. *J. Physiol.* 596, 3931–3965. doi: 10.1113/JP276239

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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ACKNOWLEDGMENTS

We thank Dr. Leslie M. Loew at University of Connecticut for providing Di-4-ANBDQPO.