

# **e-Scientific Reconstruction of Cardiac Systems @ Manchester**

**– combined approaches of  
simulation and electrophysiology**

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# Reconstruction of the cardiac pacemaker and conduction system (sinoatrial node and atrioventricular node)

Funding bodies: BHF & BBSRC

## Teams

Mark Boyett  
(Manchester)

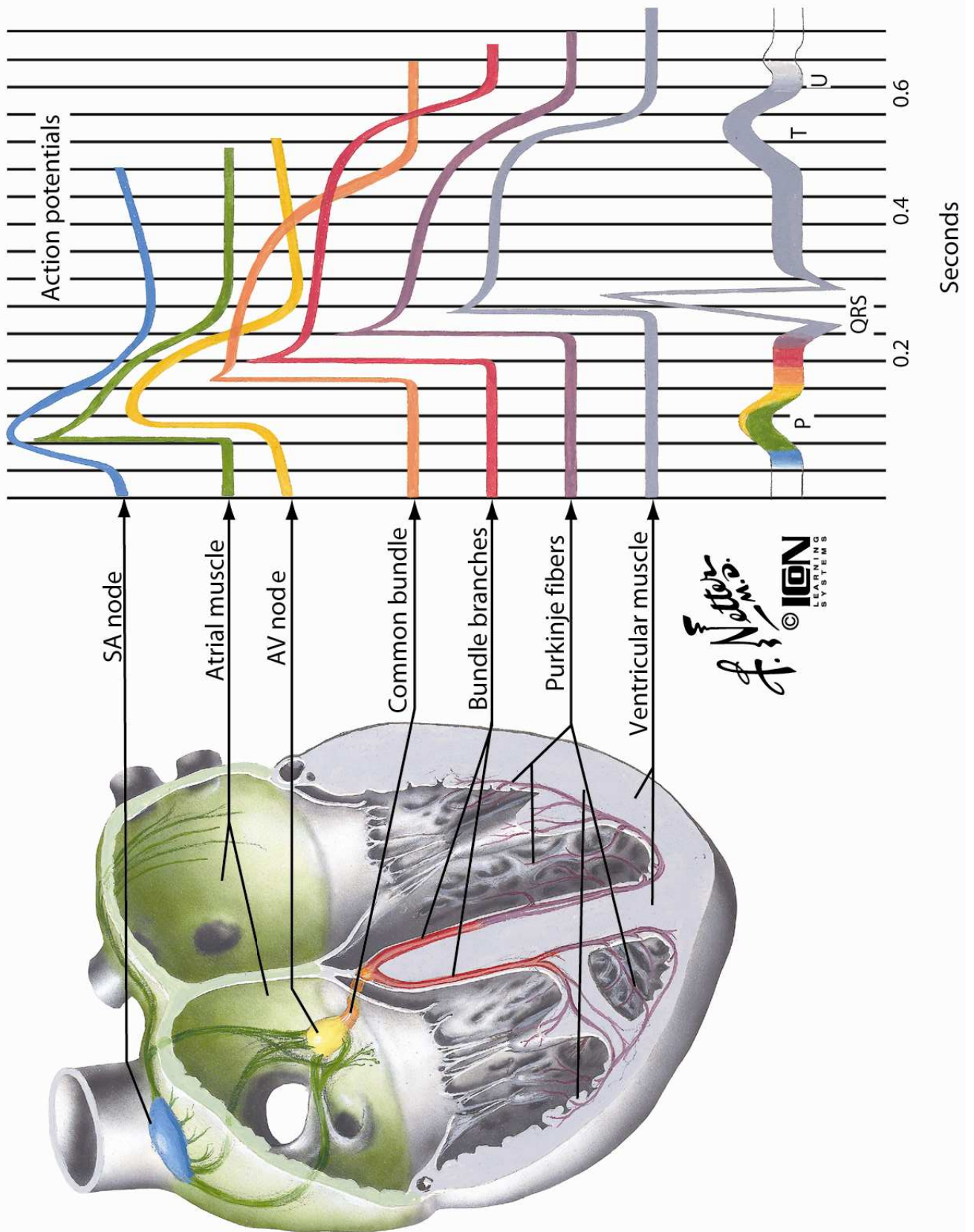
Henggui Zhang  
(Manchester)

Arun Holden  
(Leeds)

Jules Hancox  
(Bristol)

John Tucker  
(Swansea)

Min Chen  
(Swansea)



# Model of right atrium of rabbit

Jue Li, Jurgen Schneider,  
Mitsuru Yamamoto and  
Halina Dobrzynski

# Magnetic resonance imaging - rabbit

1 mm

Superior vena cava)

Sinoatrial node

Crista terminalis

Right atrium

Left atrium

Enclosed part of atrioventricular node

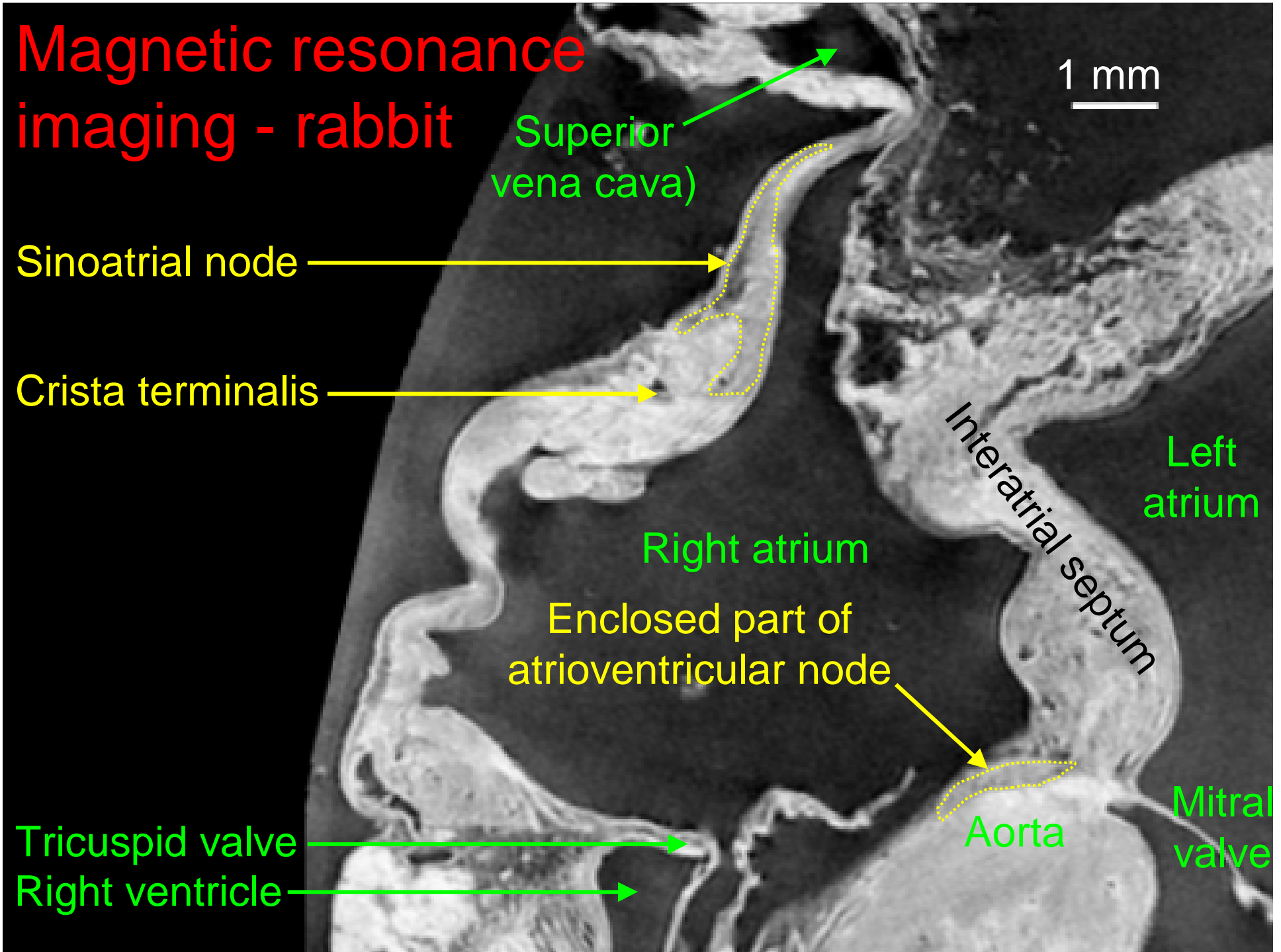
Interatrial septum

Tricuspid valve

Aorta

Mitral valve

Right ventricle



Superior vena cava →

Right atrial appendage →

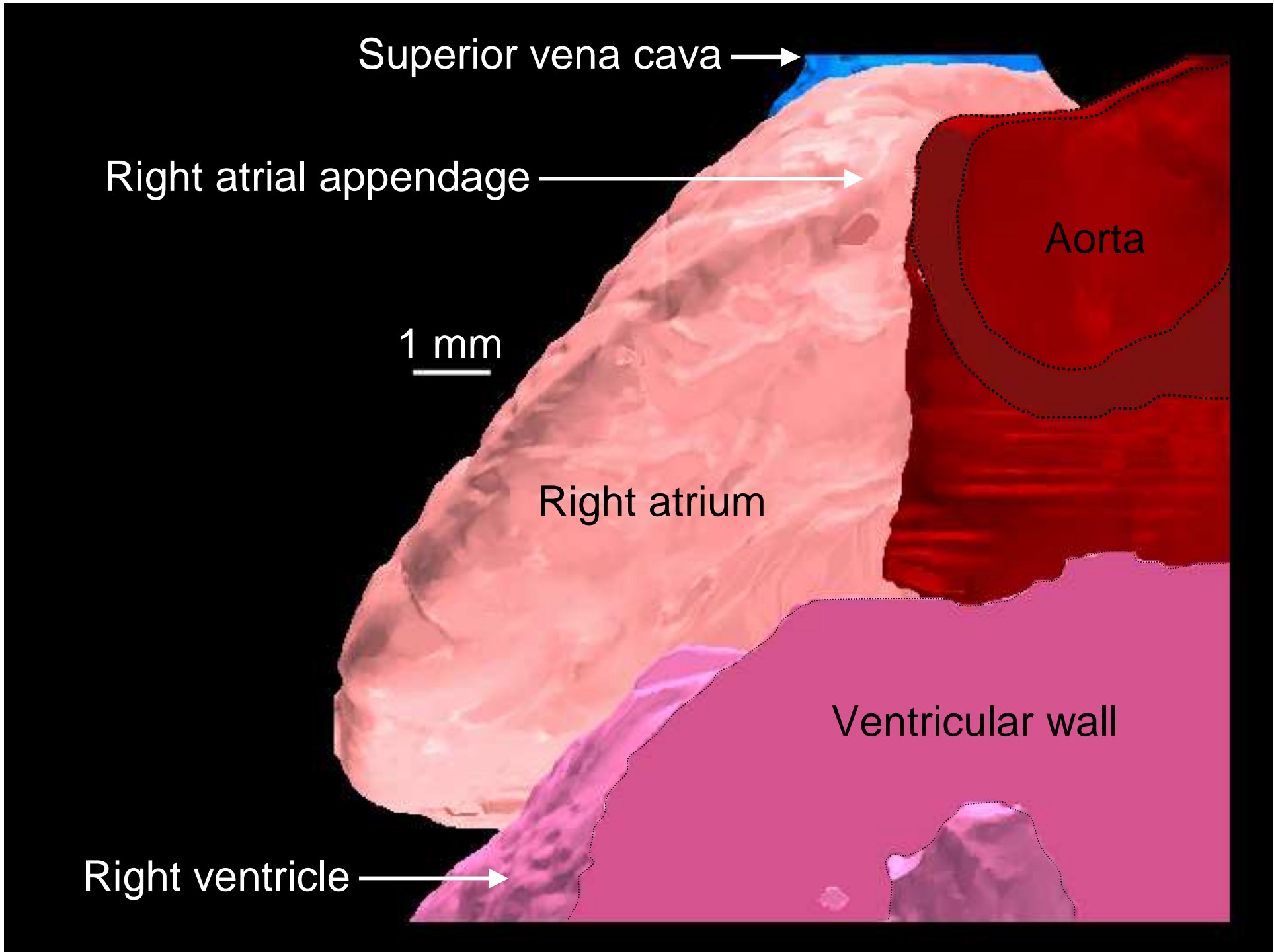
1 mm

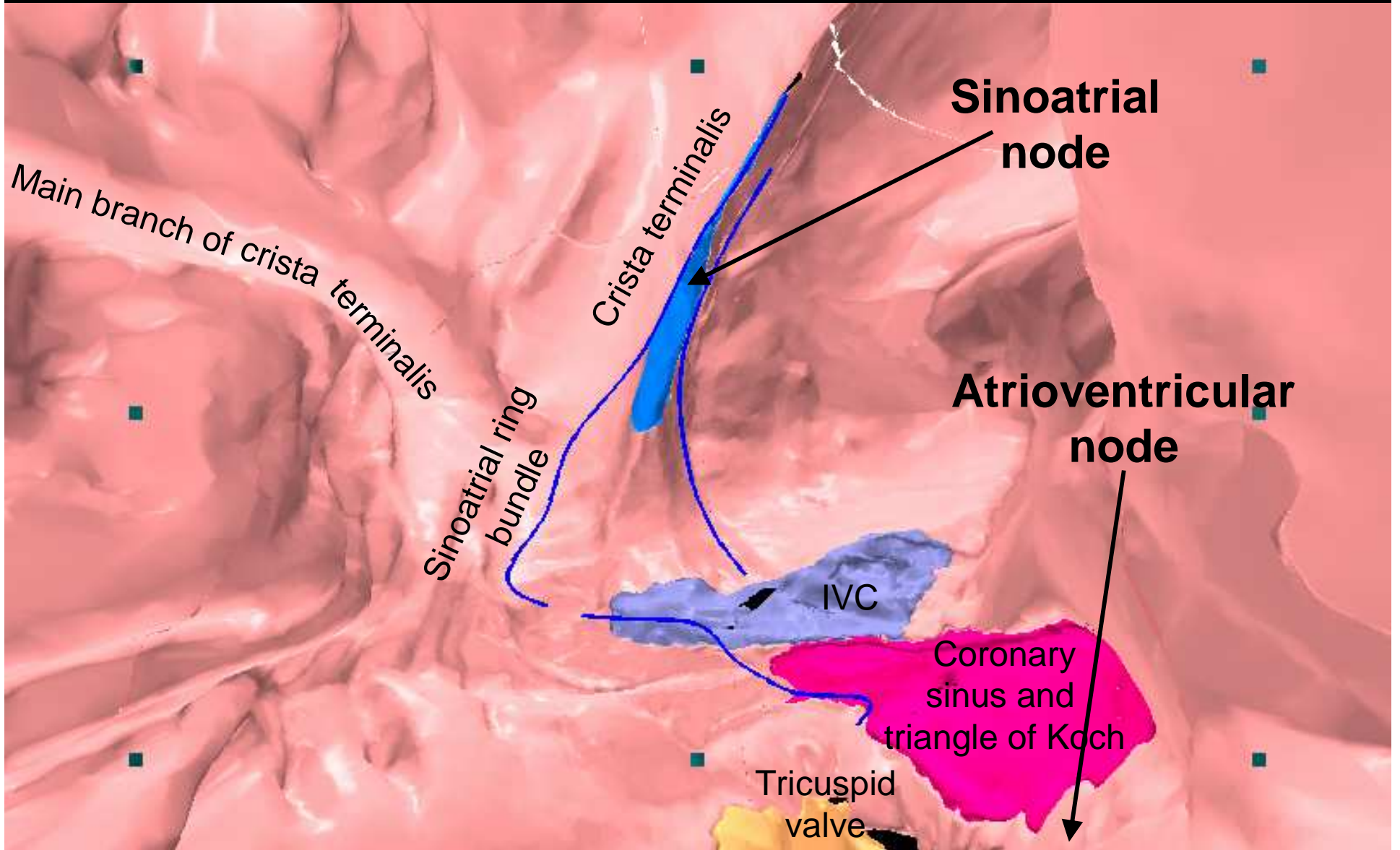
Right atrium

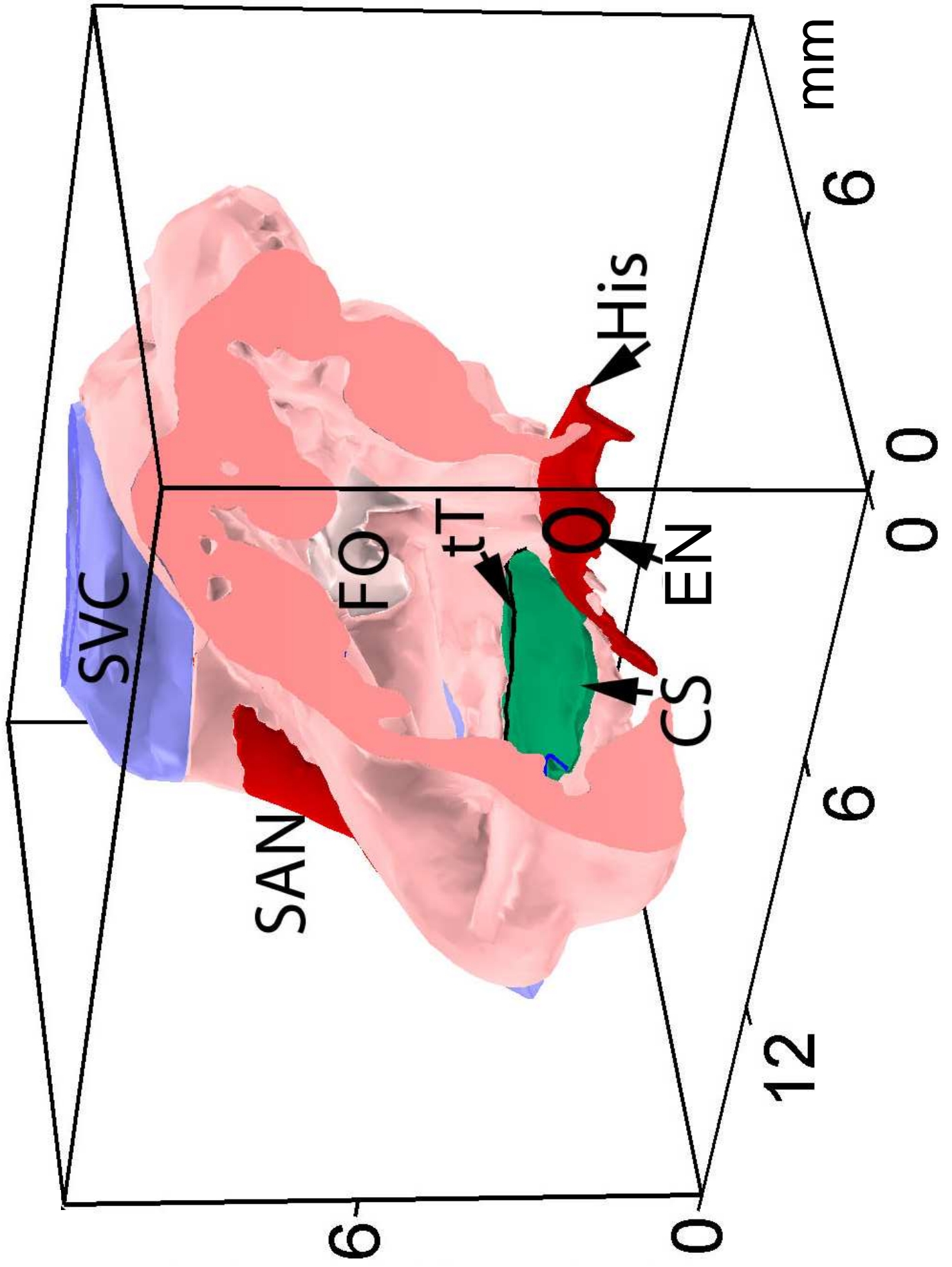
Aorta

Ventricular wall

Right ventricle →



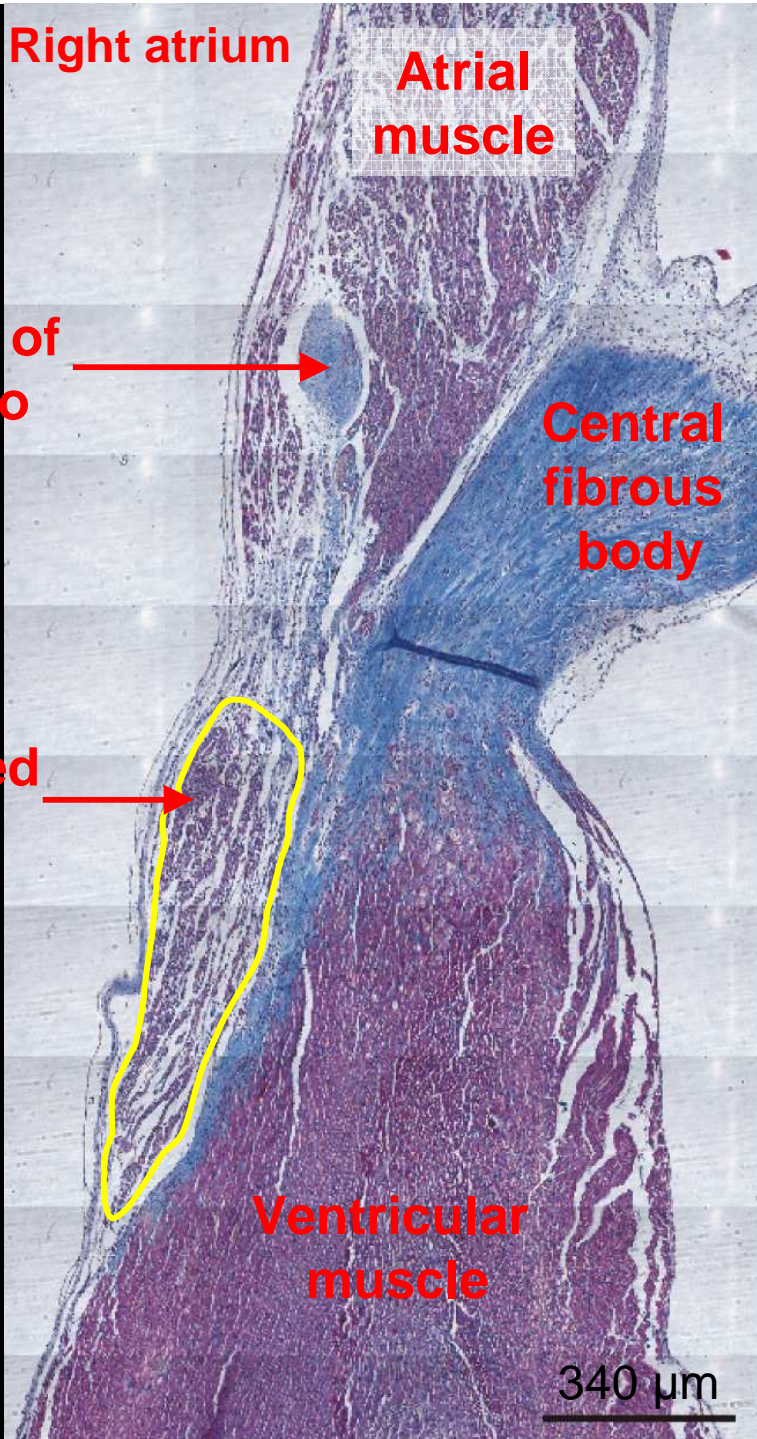




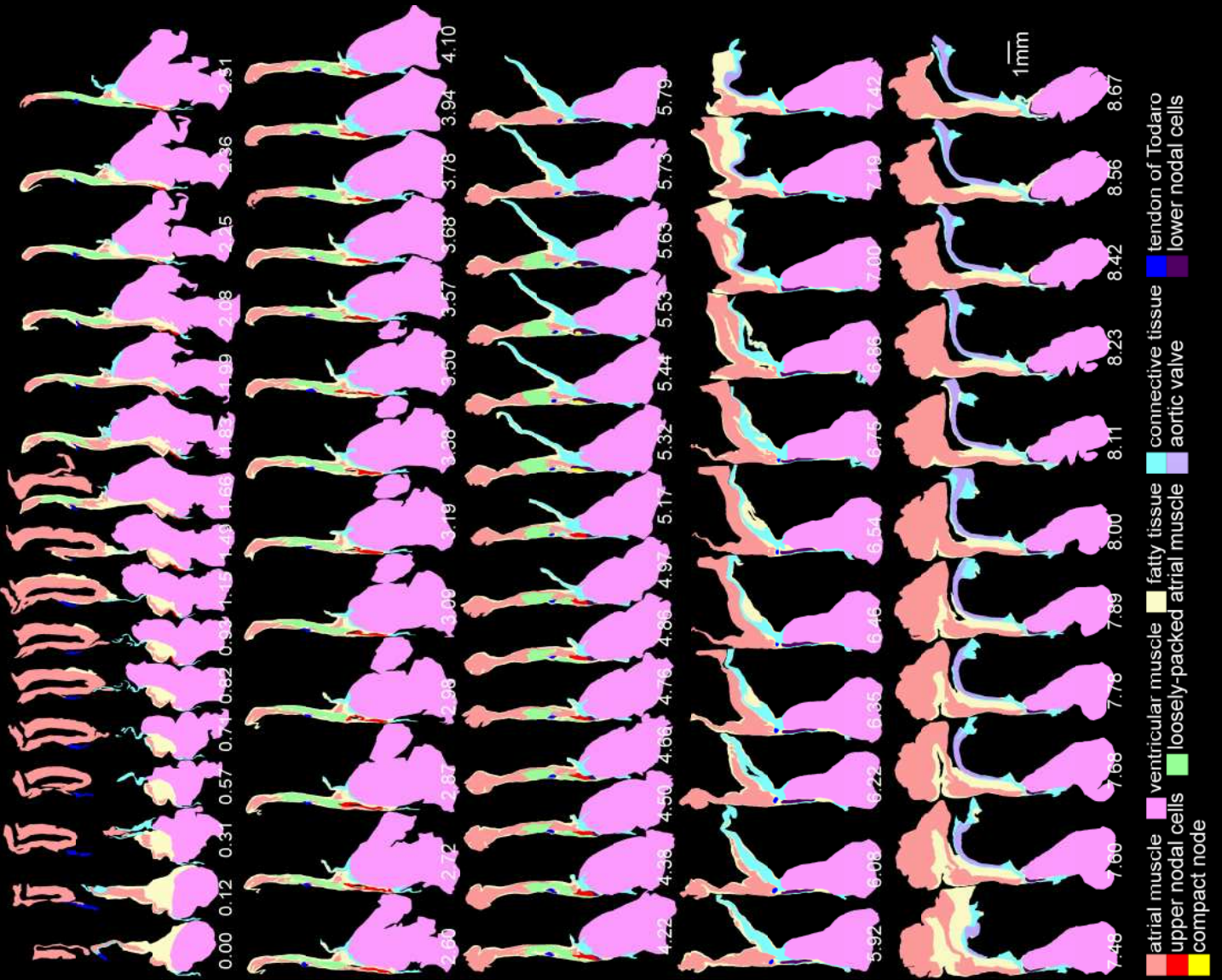


# Model of atrioventricular node of rabbit

Jue Li, Halina Dobrzynski,  
Ian Greener, Vladimir Nikoloski and  
Igor Efimov

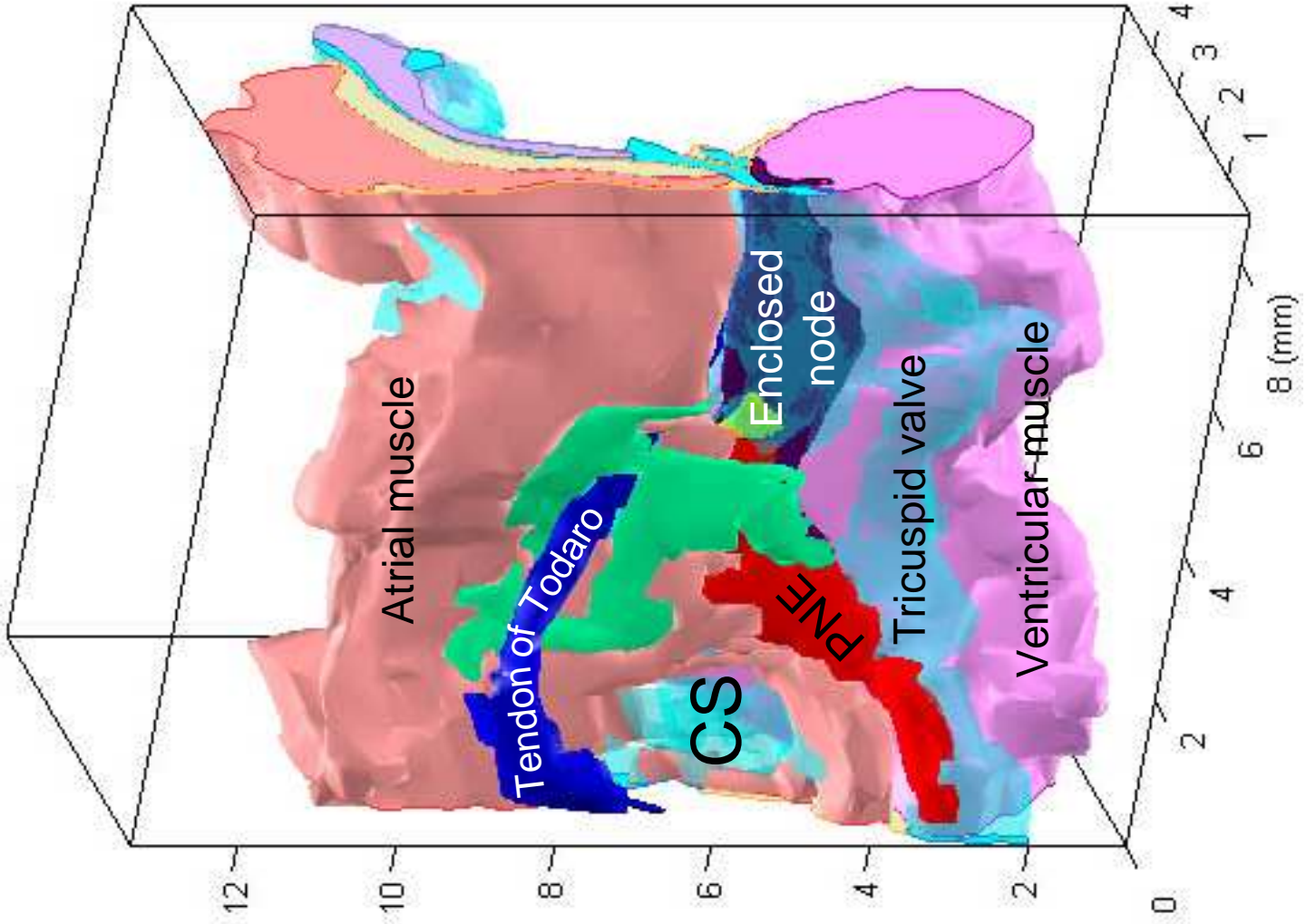


# AV node



- atrial muscle
- ventricular muscle
- fatty tissue
- connective tissue
- tendon of Todaro
- upper nodal cells
- loosely-packed atrial muscle
- aortic valve
- lower nodal cells
- compact node

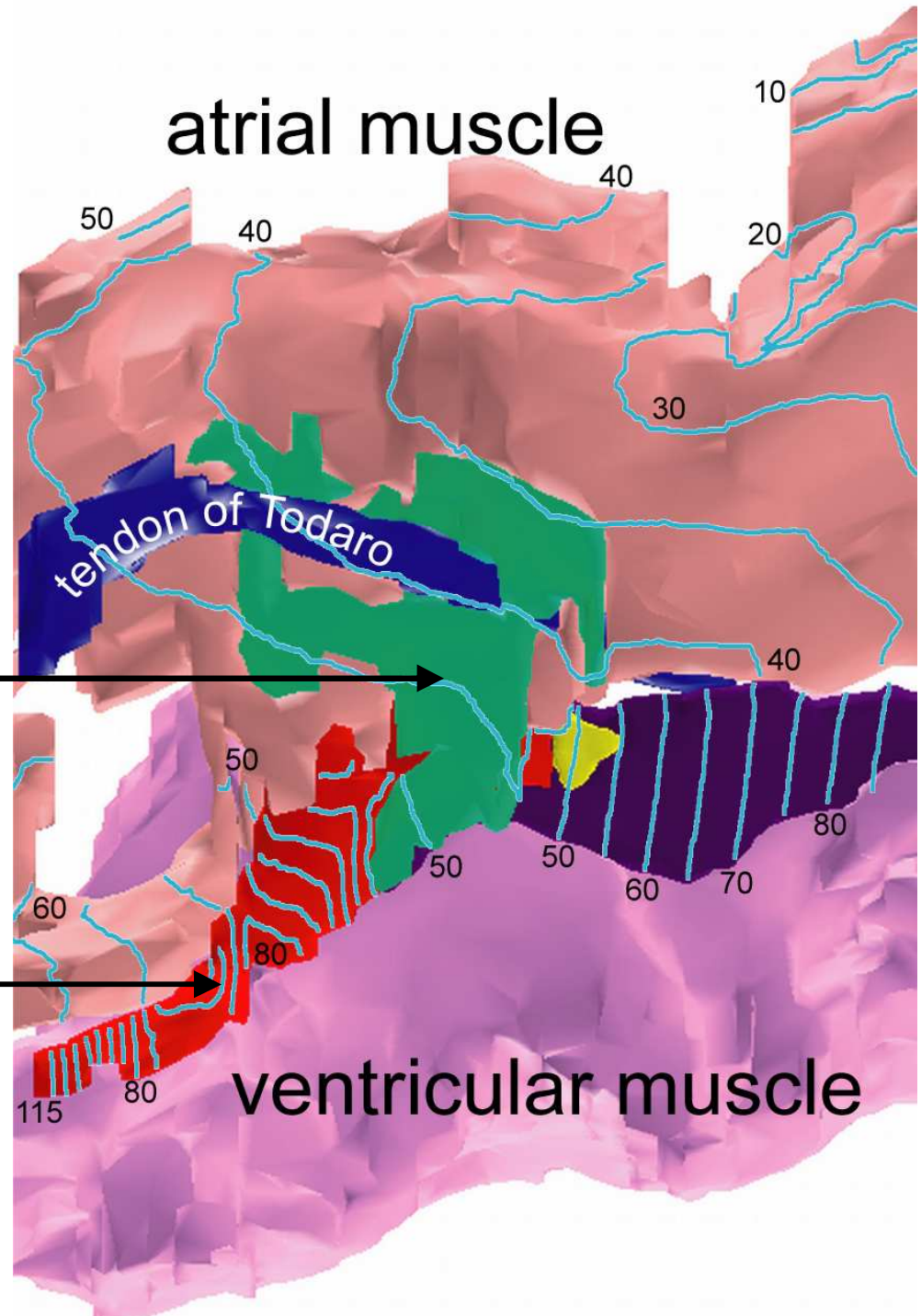
# AV node



# AV node activation sequence (cellular automaton model)

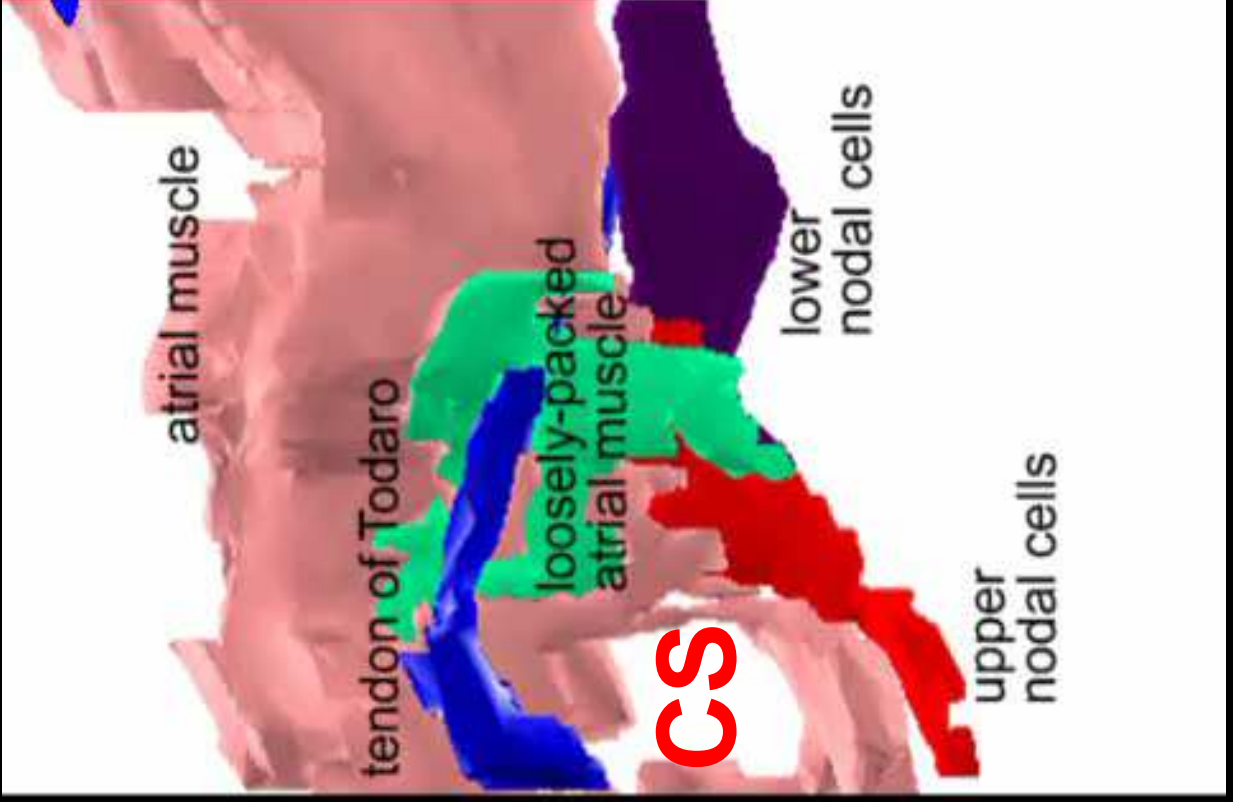
Fast pathway

Slow pathway

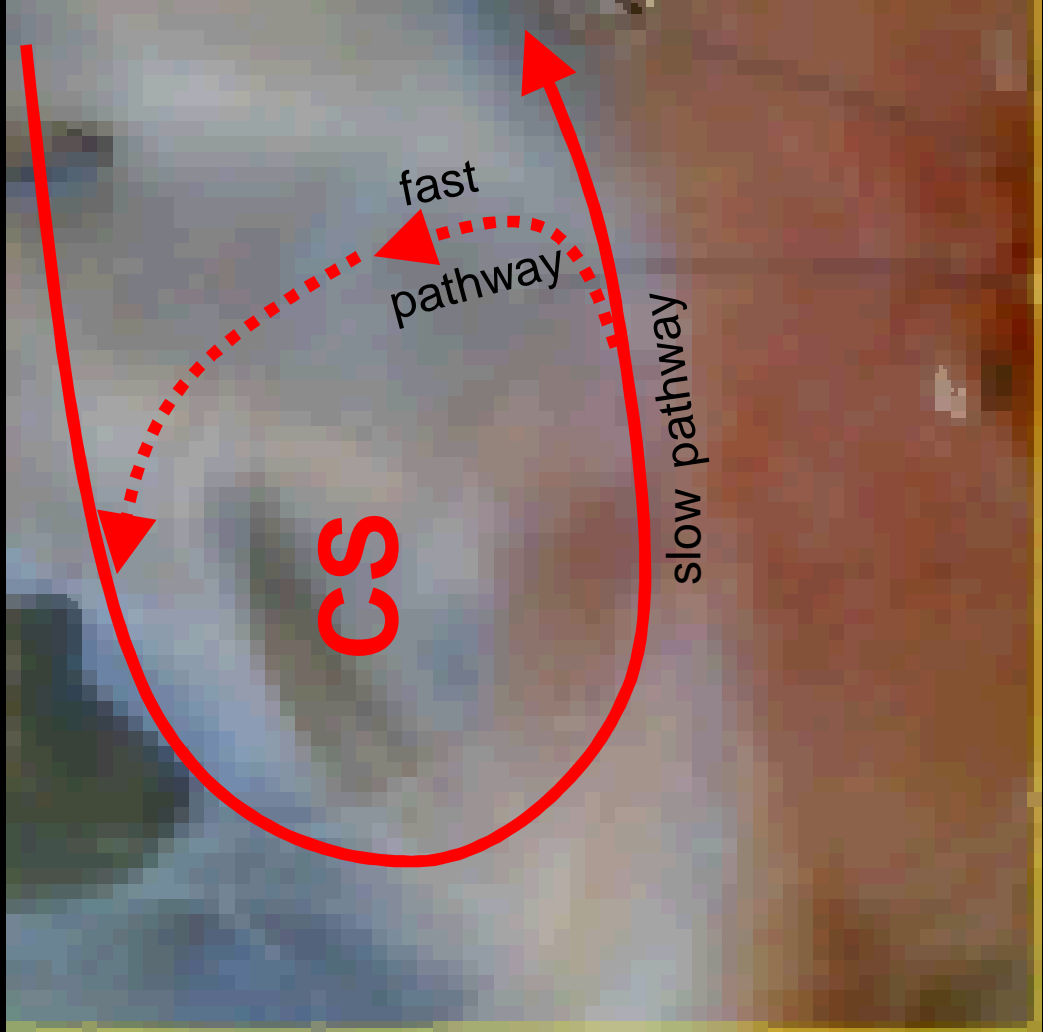


# AV nodal reentry

Model (cellular automaton model)

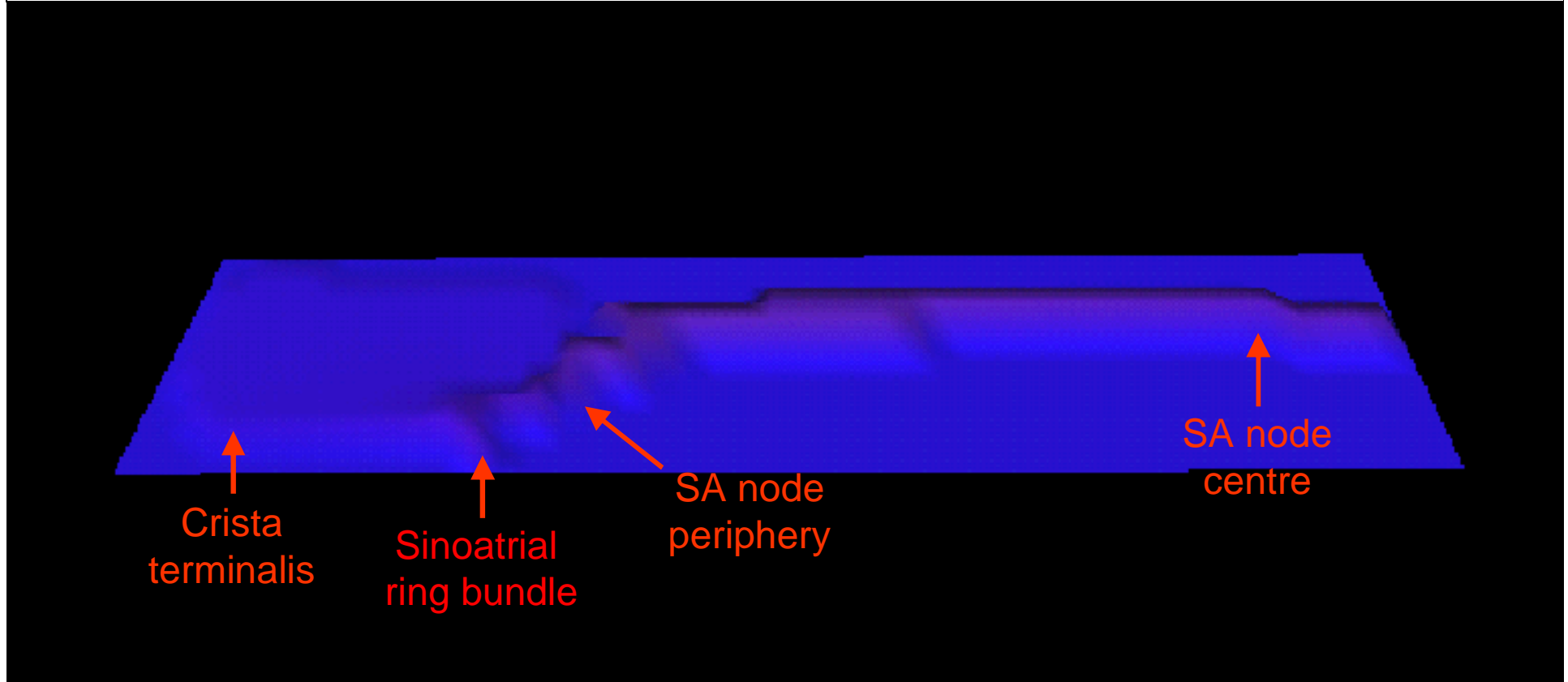
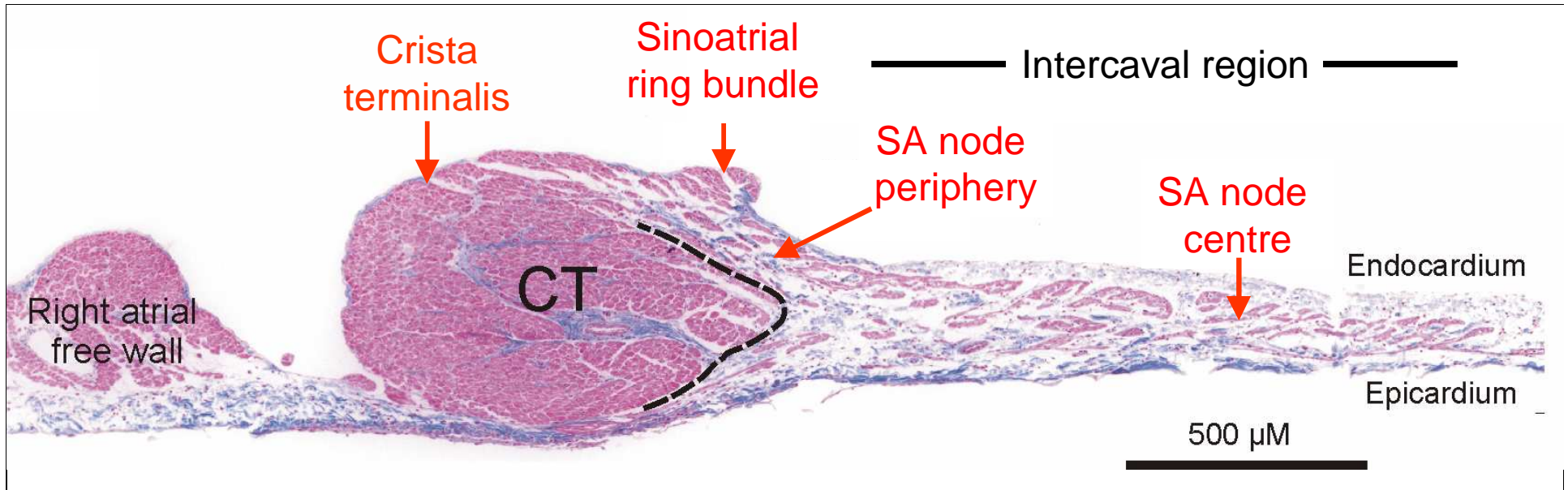


Experiment

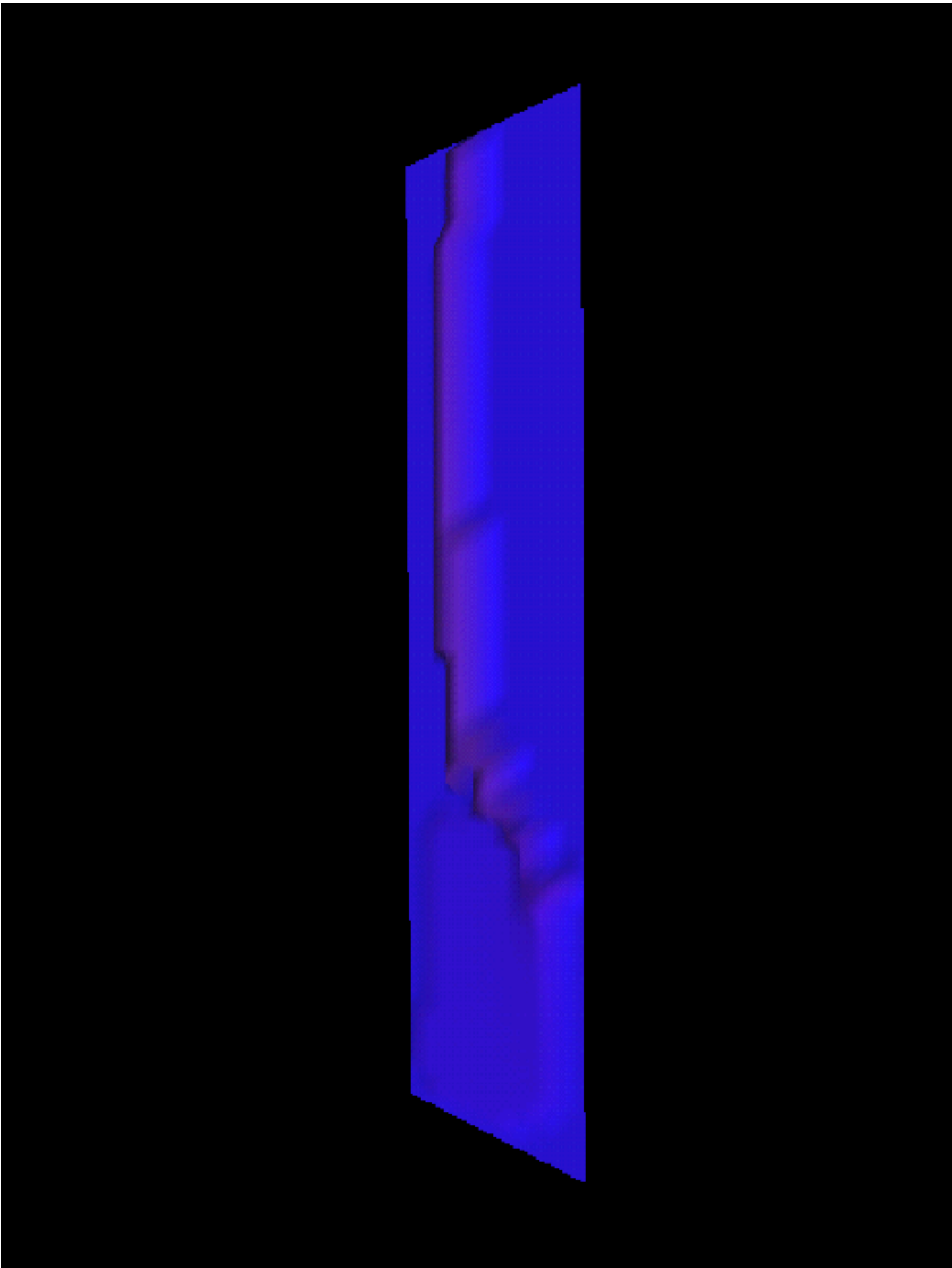


# Model of sinoatrial node of rabbit

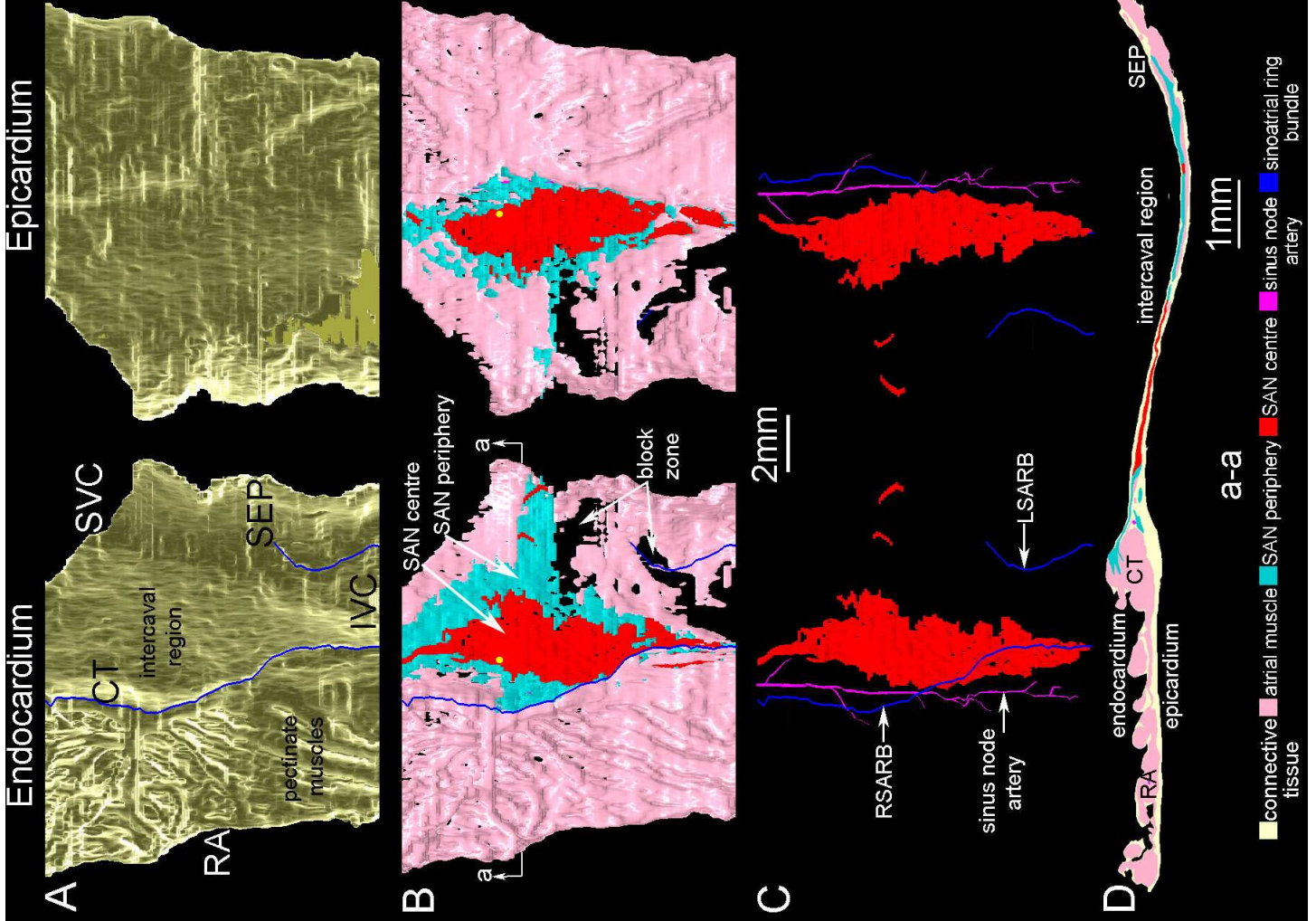
Halina Dobrzynski, Jue Li and  
Henggui Zhang





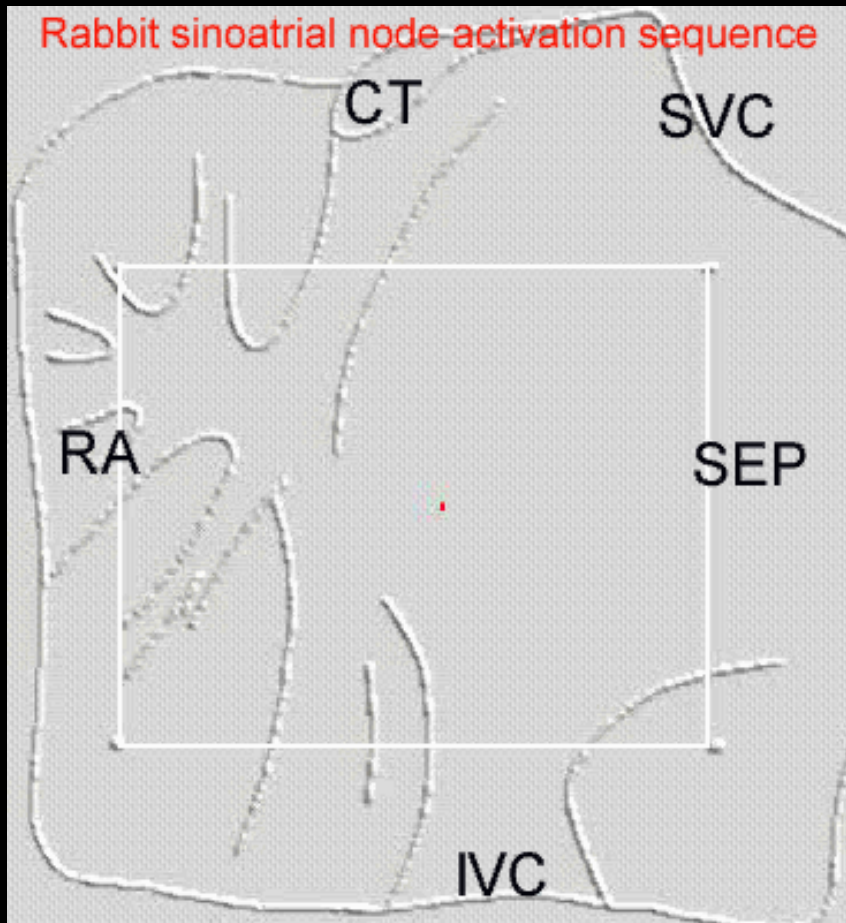


# SA node

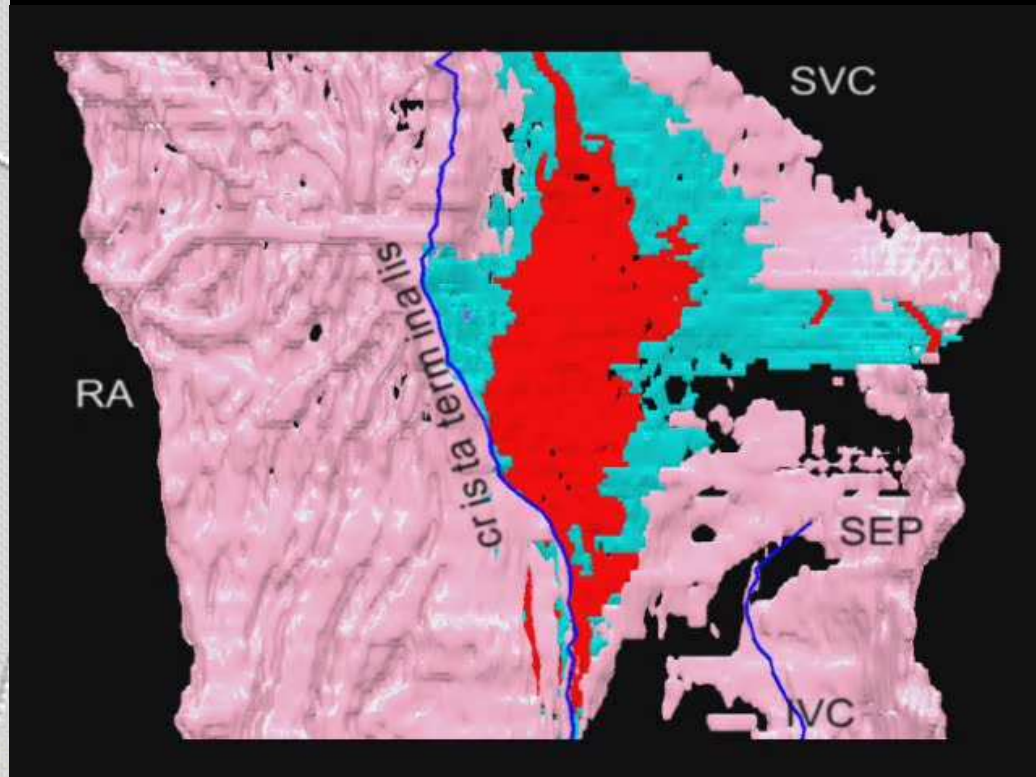


# SA node activation sequence

Experiment



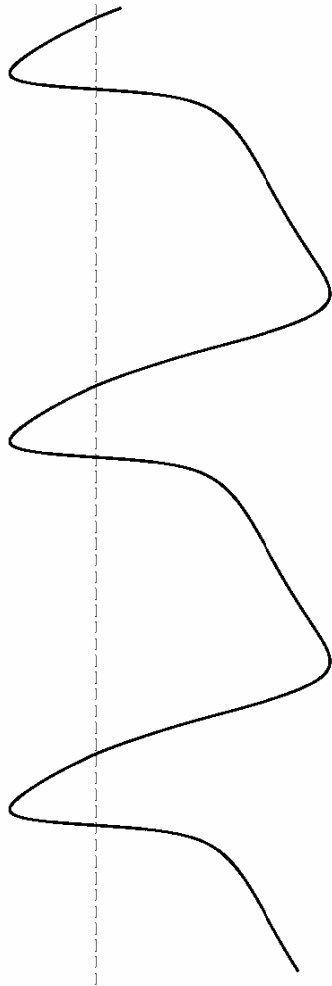
Model (cellular automaton model)



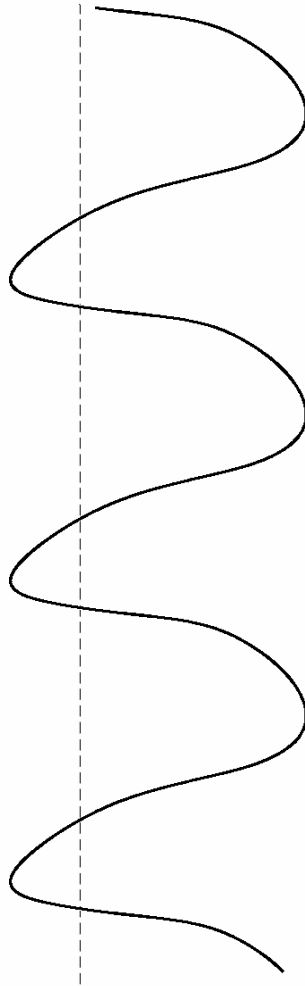
# Models of action potentials of rabbit

Shin Inada, Jules Hancox and  
Henggui Zhang

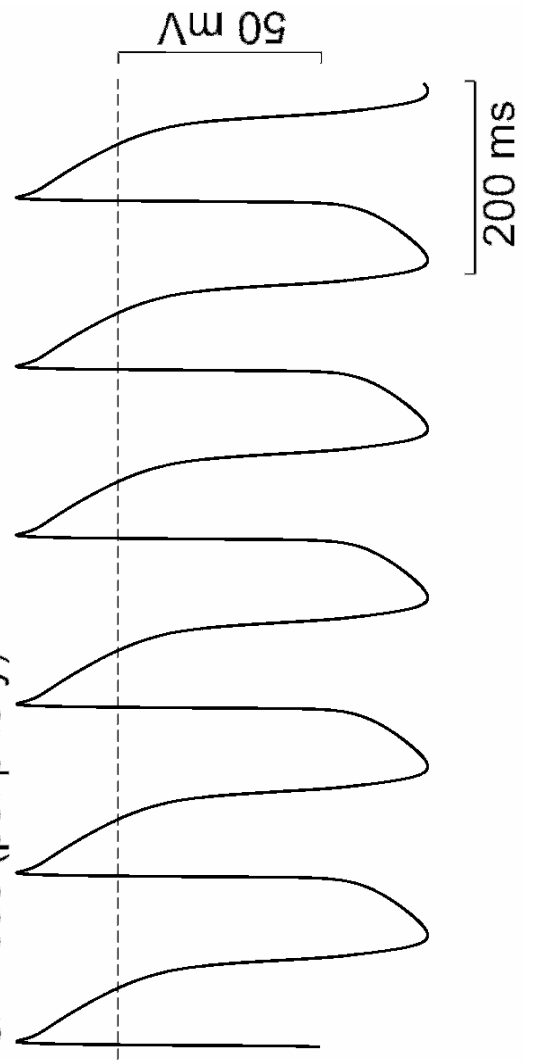
AV node

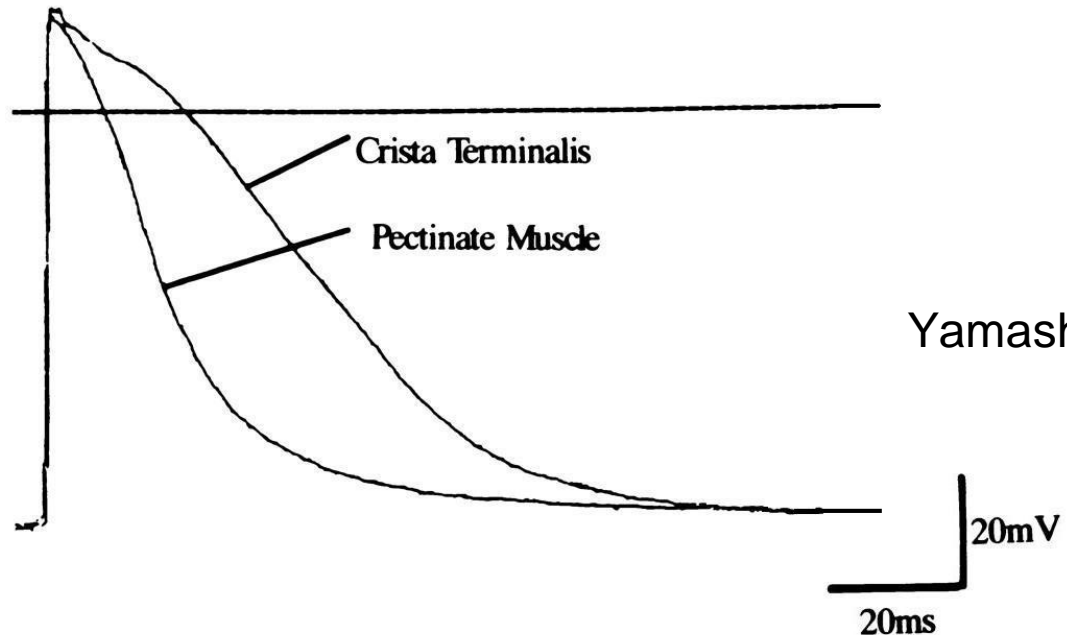


SA node (centre)

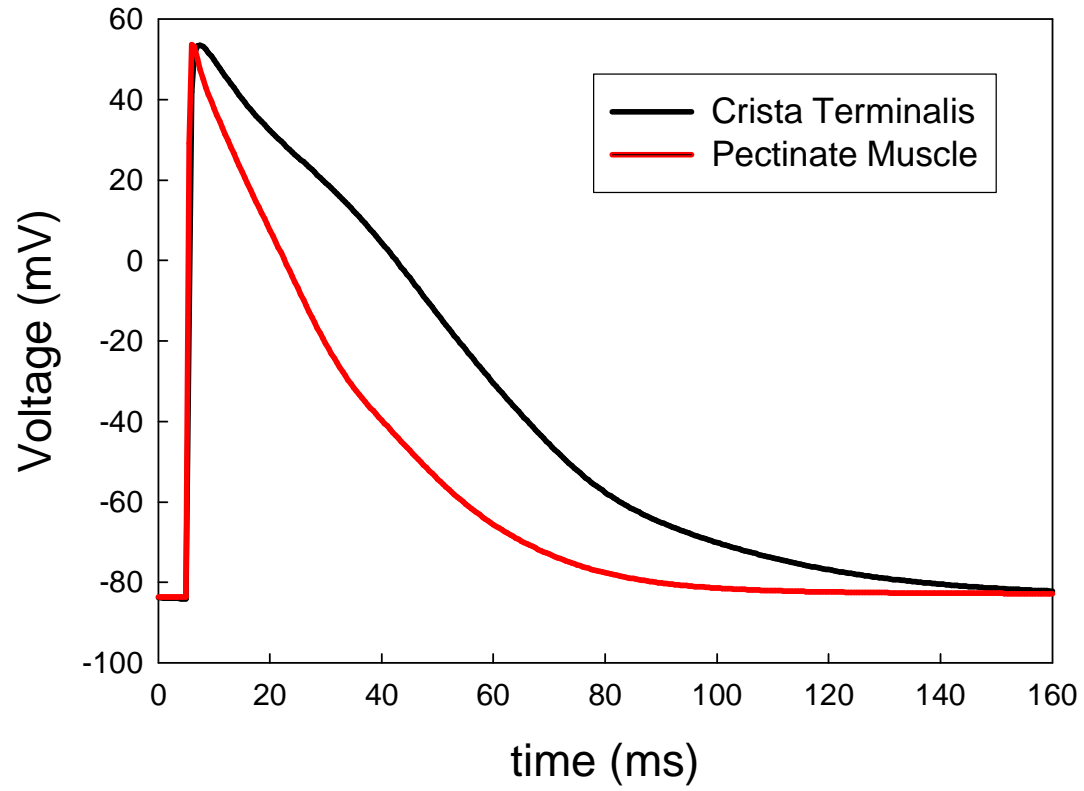


SA node (periphery)





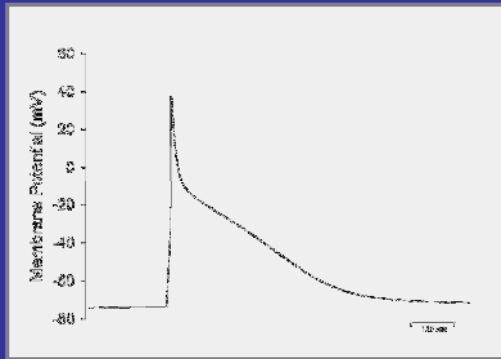
Yamashita et al., 1995



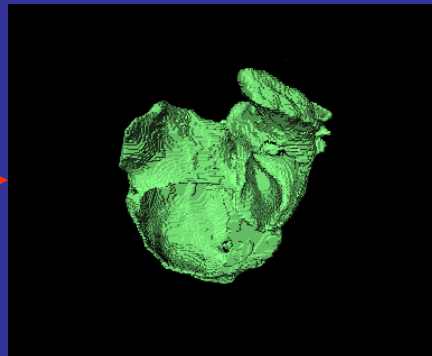
# Reconstruction of Clinical Electrophysiology of Human Atrium

# Development of Hierarchical Models of Human Atrium

- Chronic human atrial fibrillation (AF) and AF remodelling  
(BHF: 2004 – 2007; Collaborator: Prof. CJ Garratt, Manchester Royal Infirmary)
- Effects of beta-blocker on human atrial excitation  
(Collaborators: Dr. A Workman and A Rankin, Glasgow Royal Infirmary)



Cellular Electrical  
model



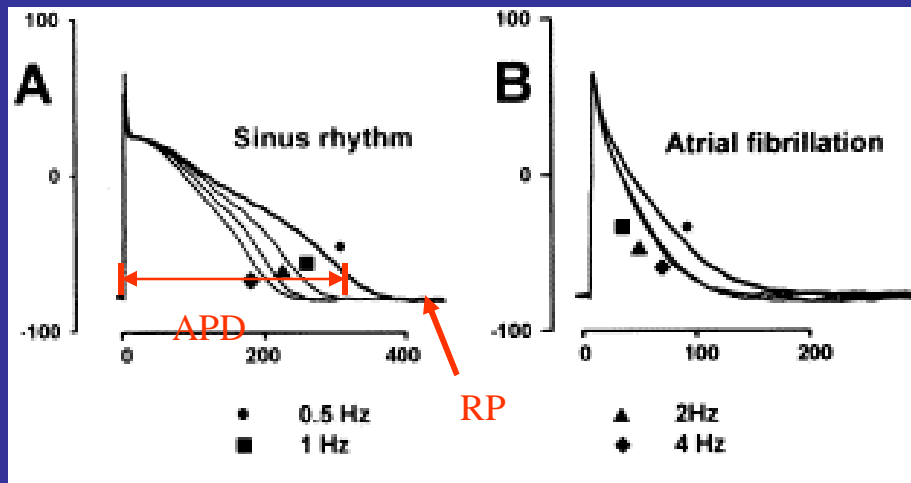
Anatomic  
Geometrical  
Model

- Anatomic model of atrium
- Ionic mechanisms underlying
  - ❖ initiation
  - ❖ terminationof atrial arrhythmias
- Complexity association between distortion of chambers and arrhythmias
- Actions of anti-arrhythmia drugs



# Simulation of AF-induced Ion Channel Remodelling

# Chronic AF-induced changes in the electrical activity of human atrial myocyte



	RP (mV)	APD <sub>90</sub> (ms) (0.2 Hz)
SR	-76.3±2.2	319 ±48
AF	-78.9 ±2.9	134 ±12
Chang	-2.6 ↓	-58% ↓

SR: sinus rhythm

AF: atrial fibrillation

RP: resting potential

Bosch et al. (1999)

Cardiovasc Res 44: 121-131

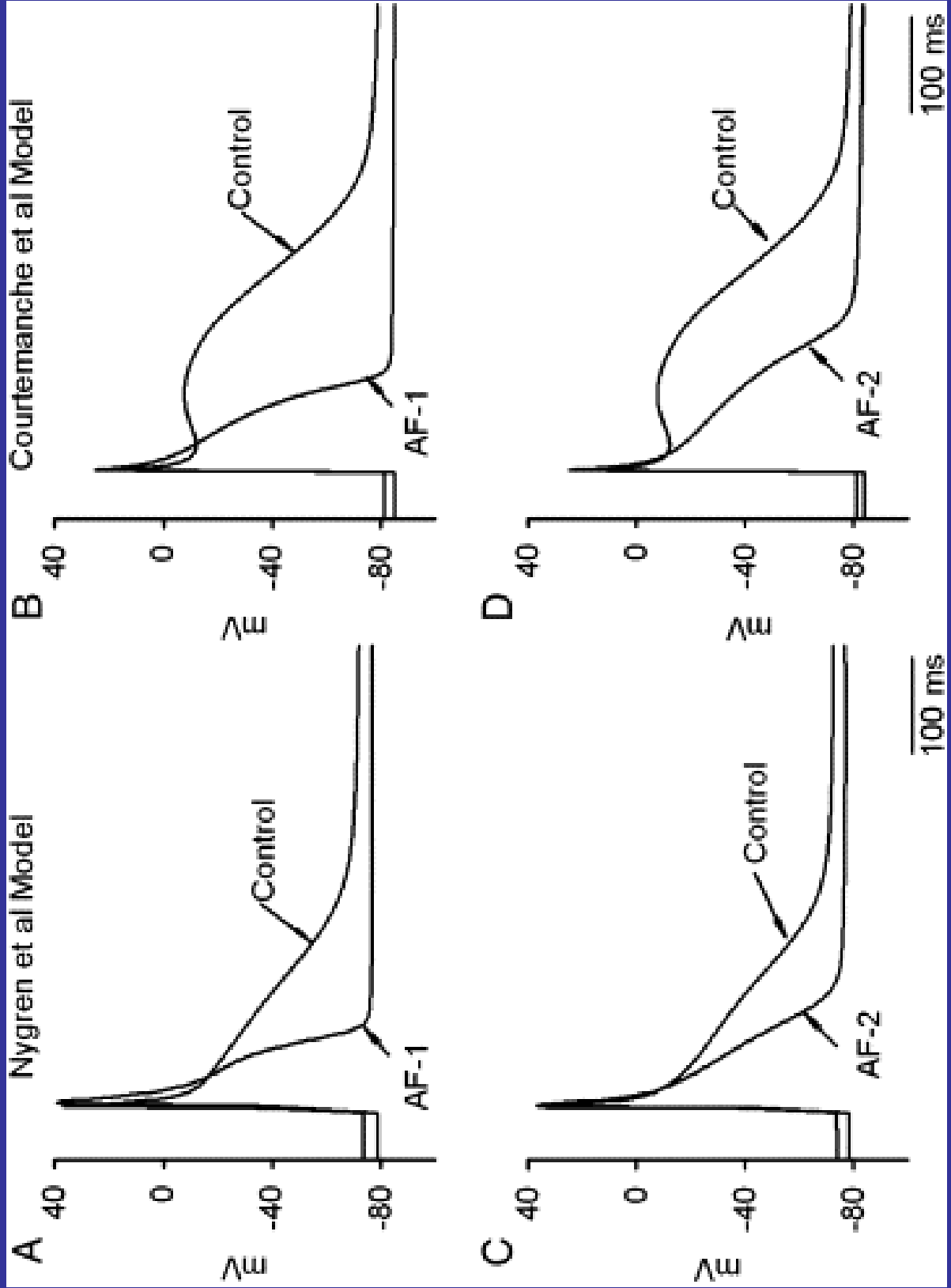
## Chronic AF-induced remodelling of ionic channels

	Regulation	Current density	Channel kinetics
$i_{Ca,L}$	Down ↓	↓ 74%	$\tau_{inact,fast}$ ↑ 62%
$i_{to}$	Down ↓	↓ 85%	Activation curve shifted by 16 mV
$i_{K,sus}$	--	--	--
$i_{Na}$	--	--	Inactivation curve shifted by 10 mV
$i_{K,1}$	UP ↑	250% (-90 mV) ↑ 235% (-20 mV)	--

Bosch et al. (1999) Cardiovasc Res 44: 121-131

# RESULTS

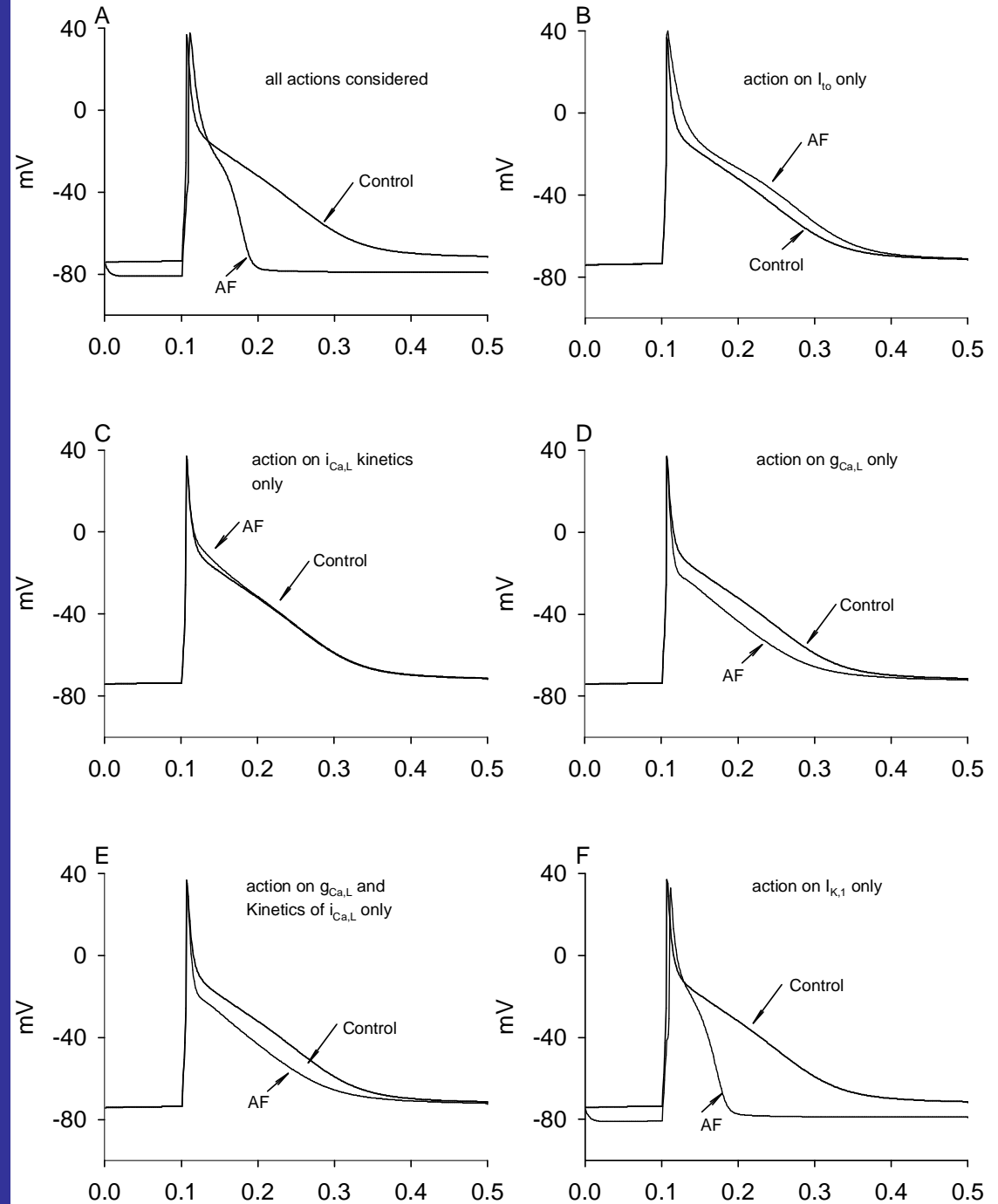
## Simulation of AF-remodelling



## Role of each individual AF remodelled ionic channels

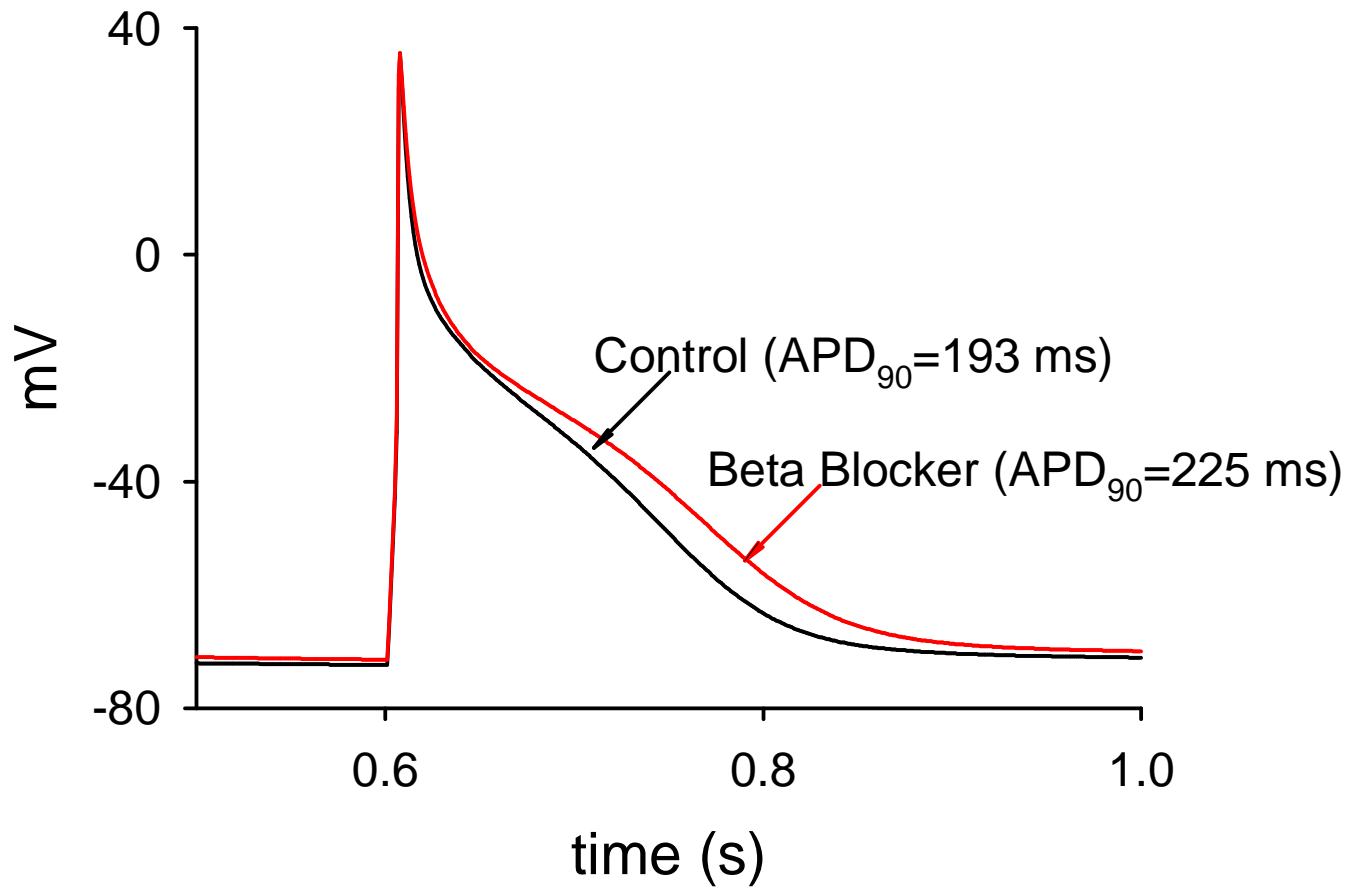
**Exclusive method:**  
*all other actions are removed while only the interested action is considered.*

Zhang et al.  
Cardiovascular Research (2005)



# Effects of BB Induced Ionic Channel Remodelling on Human Atrial AP & Conduction

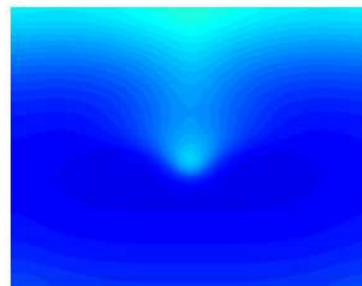
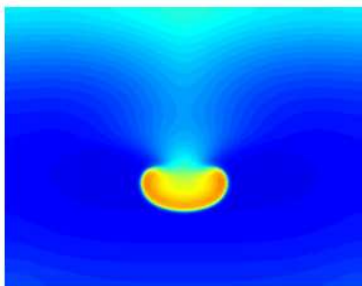
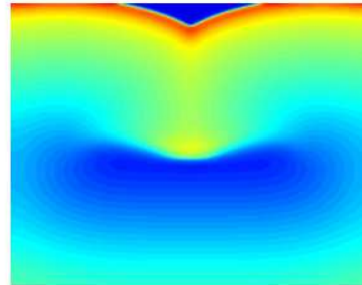
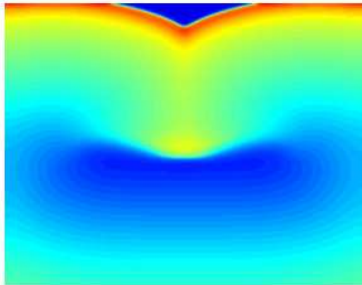
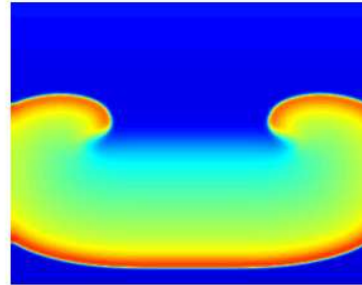
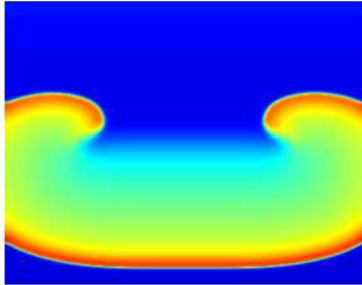
# Effects of BB on normal human atrial APs



# Control

L = 55.04 mm

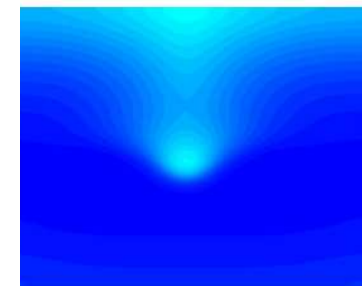
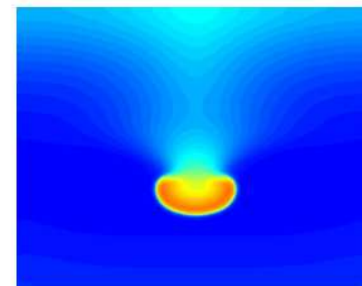
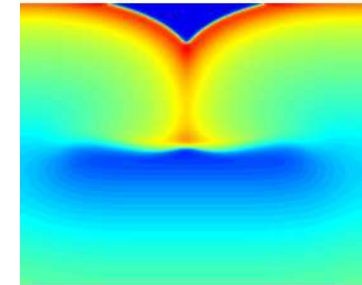
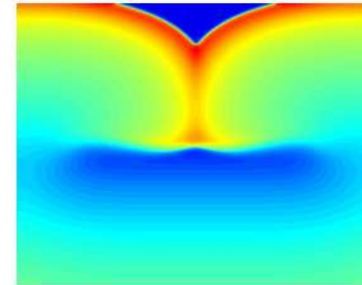
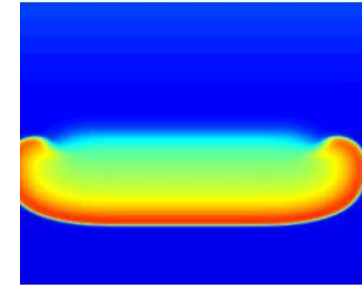
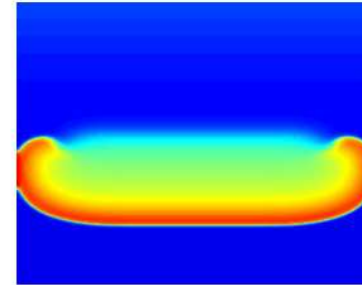
l = 54.62 mm



# Beta Blocker

L = 67.2 mm

l = 66.88 mm





# Reconstruction of Electrophysiology of Ventricle

# Current projects on ventricle modelling

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- Short QT syndrome and gene mutation of potassium channels  
(Collaborator: Prof. JC Hancox, Bristol) (Zhang & Hancox Biochem, Biophys Res Commun 2005)
- Acute ischemia  
(Collaborators: Prof. JC Hancox, Bristol)
- Exercise remodelling  
(Dr. E White, Leeds)
- Effects of volatile anesthetics  
(Dr. S Harrison, Leeds)

# Sub-cellular modelling

## Intracellular cardiac Ca<sup>2+</sup> handling

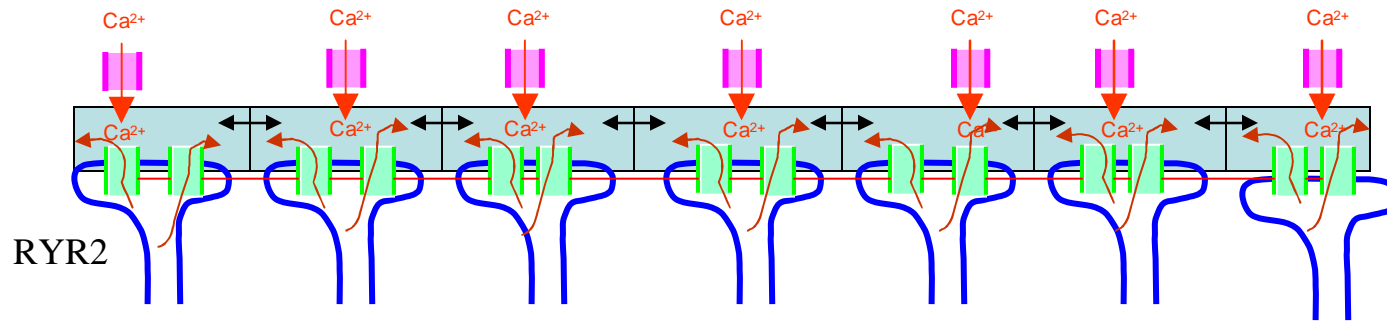
Modelling team

Dr. H Zhang  
(Manchester)

Experimental team

Prof. D. Eisner & Dr. S. O'Neil  
(Manchester)

# Model of cluster of RyRs



$$\frac{d[Ca^{2+}]_{sub}}{dt}(x) = J_{Ca, L} + J_{Ca, T} + J_{rel} - J_{NaCa} - J_{Ca, diff} - J_{Ca, spacial\_diff} - J_{buffer, sub}$$

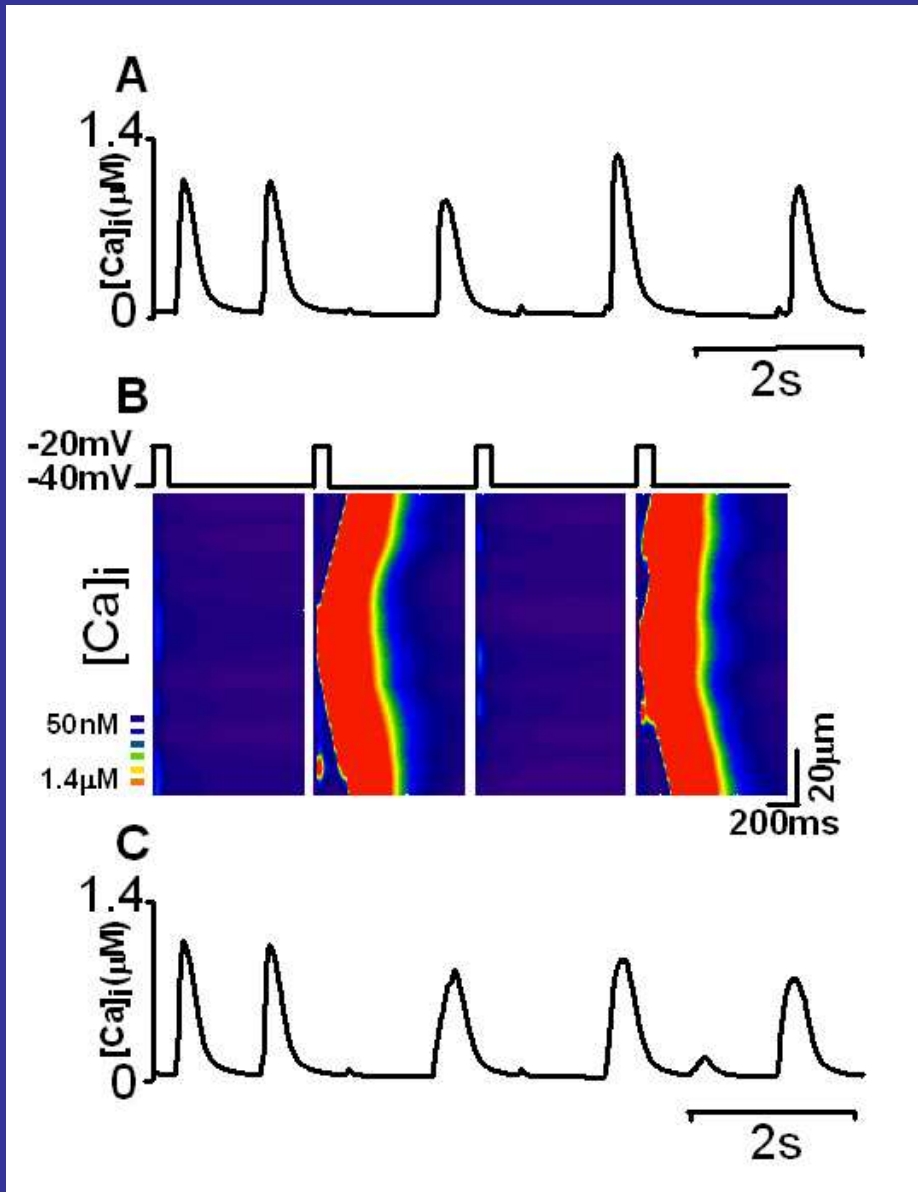
$$\frac{d[Ca^{2+}]_i}{dt}(x) = J_{Ca, diff} + J_{Ca, spacial\_diff} - J_{up} - J_{Ca, P} - J_{buffer, i}$$

$$\frac{d[Ca^{2+}]_{rel}}{dt}(x) = J_{tr} - J_{rel} - J_{buffer, rel}$$

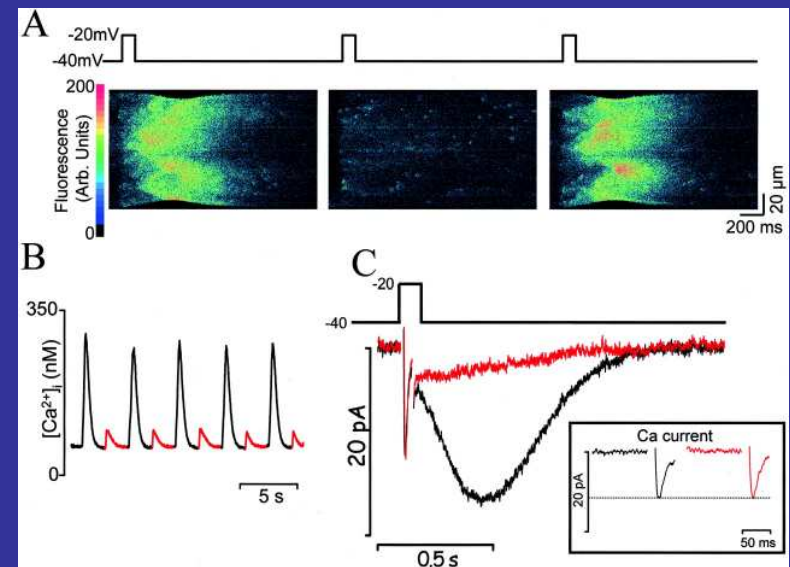
$$\frac{d[Ca^{2+}]_{up}}{dt}(x) = J_{up} - J_{tr} - J_{buffer, up}$$

(1)

# Simulation of intracellular Ca spark and waves



Simulation



Diaz, M. E. et al. *Circ Res* 2004;94:650-656

Experiment