Excitable Media as Model of Cardiac Arrhythmia

Werner Lehner 23.03.11

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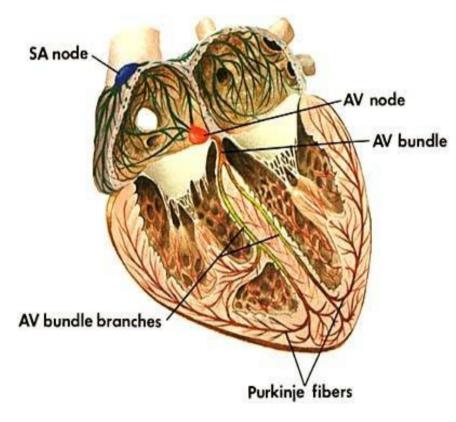
Outline

- Cardiac Arrhythmias
 - Cardiac Conduction System
 - Action Potential
 - Fibrillation
 - Reentry
- Modeling
 - Model
 - BZ-Reaction
 - Finite Element Method
 - Software
- Simulations
 - Plain
 - Sphere
 - Real Geometry

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ISBT

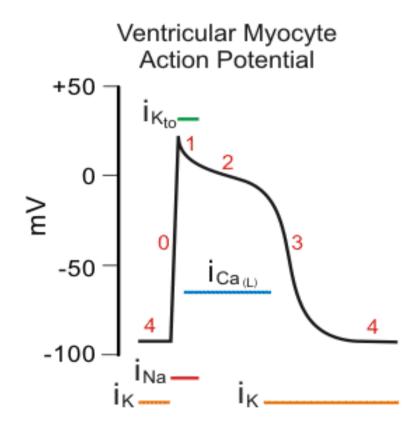
Cardiac Conduction System



- SA-node: primary pacemaker
- AV-node: electrical connection between atria and ventricles
- Av-bundle (His-bundle): transmission to apex
- Purkinje fibers: distribution to ventricular myocardium
- Contraction starts at apex
- AP spreads across ventricles (gapjunctions

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Action Potential

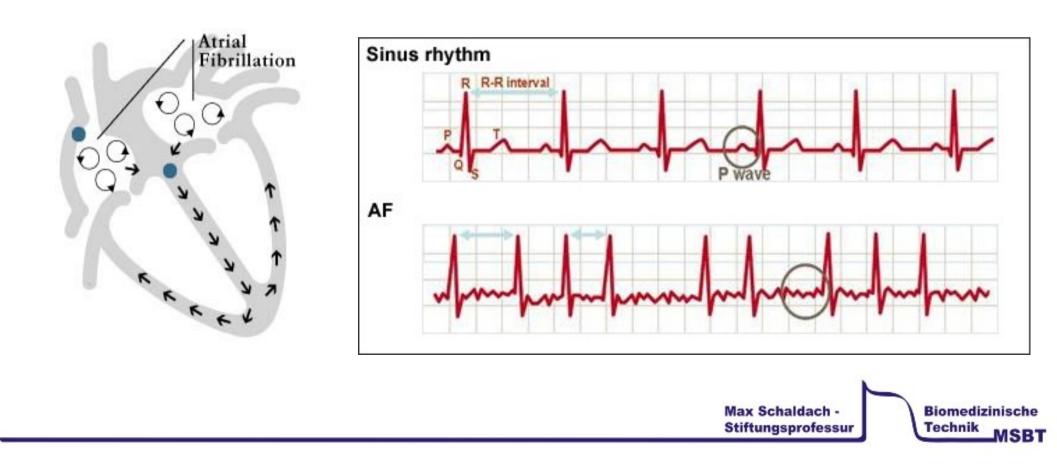


- Rapid change of transmembrane Potential
- Complex system of ion channels
- Resting Potential (4)
- Threshold Potential: ~ -70mV
- Fast depolarization (0)
- Slow repolarization (2,3)
- Effective refractory period (ERP): new AP can not be triggered
- No back-propagation of APs
- All-or-none-characteristic



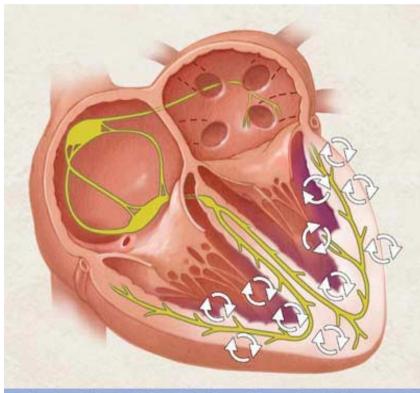
Fibrillation

- Uncoordinated, very rapid activity
- Complex, chaotic electrical patterns
- Atrial (AF), ventricular fibrillation (VF)
- APs propagate on irregular pathways
- Deterioration of atrial/ventricular function
- ECG: P-wave replaced by fibrillatory waves (AF)



Fibrillation

- Substrate: inhomogeneities in electrophysiol. properties of myocardium (refractory period, conduction velocity, excitability)
- Fractionation of wavefronts in such areas (multiple wavelet mechanism)
- Formation of self-perpetuating impulses (reentry mechanism)

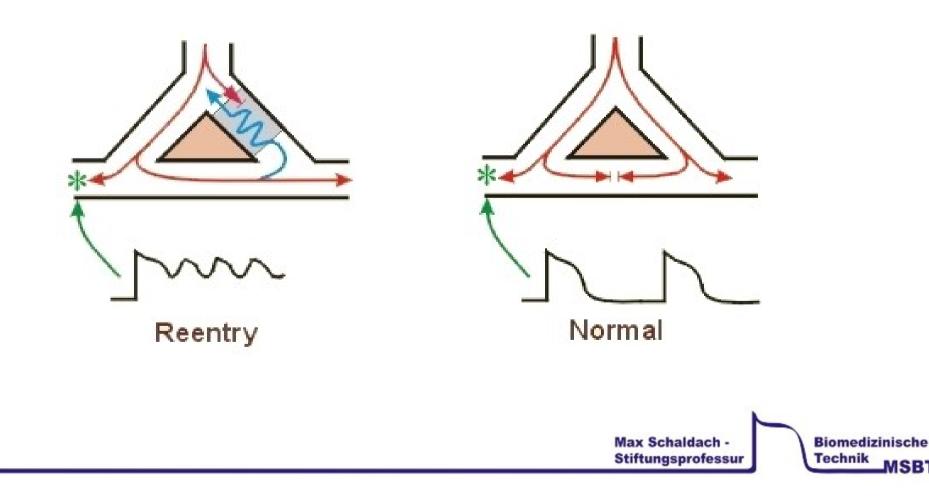


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Reentry

- Obstacle: anatomic (myocardial infarction), functional (refractory area)
- Unidirectional conduction block
- Wavelength = conduction velocity x refractory period
- Reentry possible if pathway >= wavelength



SBT

Reentry



"Leading circle": source of spiral waves

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Therapy

- Drugs (change electrophysiol. parameters, AF)
- Ablation (AF): destruction of reentry pathways
- Defibrillation (VF): depolarization of the whole myocardium by internal or external high-energy shock
- No mild therapy available (VF)

Termination of (multiple) Reentry by local low-energy Stimulation?

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Modeling

- Goals:
 - Simulation of reentrant-excitation
 - Simulation of electric stimulation
 - Stimulation-algorithm for termination of reenrty
- ToDo:
 - Choice of appropriate mathematical Model
 - Method for numerical solution
 - Software-framework
 - Simulations

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Biomedizinisch

Models

- Cell-based models:
 - Model for membrane potential of a single cell
 - Emulation of ion-channel functionality
 - e.g. Hodgkin-Huxley-Model:

$$C_{Mem} \frac{dV_{Mem}}{dt} = I_{Na} + I_{K} + I_{Ext} + I_{L}$$

- in most cases very complex (>30 PDEs)
- Tissue = network of many single cells \rightarrow numerical effort



Models

- Excitable media
 - Model for system of many cells
 - Reaction-diffusion system
 - AP described by activator-inhibitor system ("reaction"):

$$\frac{\partial u}{\partial t} = f(u, v) \quad , \quad \frac{\partial v}{\partial t} = \varepsilon g(u, v)$$

• AP-propagation ("diffusion")

$$\dot{u} = D \nabla^2 u$$

- U: membrane potential ("activator")
- V: repolarization current ("inhibitor")
- AP-characteristics: ref. period, no damping, all-or-none

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• Starobin-Starmer:

$$\frac{\partial u}{\partial t} = D \nabla^2 u + \lambda(u) u - m - v \quad , \quad \frac{\partial v}{\partial t} = \varepsilon (\gamma u - v)$$

• Panfilov:

$$\frac{\partial u}{\partial t} = D \nabla^2 u + \lambda (u, v) u - m - v \quad , \quad \frac{\partial v}{\partial t} = \varepsilon (u, v) (\gamma u - v)$$

• Tyson-Fife (BZ-reaction)

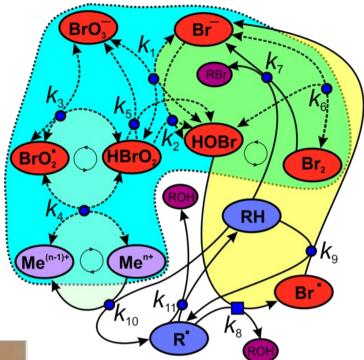
$$\frac{\partial u}{\partial t} = D \nabla^2 u + u - u^2 - (fv + \phi) \frac{u - q}{u + q} \quad , \quad \frac{\partial v}{\partial t} = \varepsilon (u - v)$$

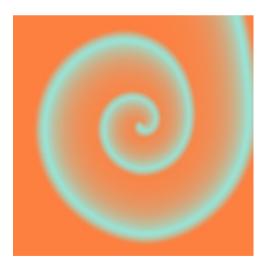
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ASBT

BZ-Reaction

- Nonlinear chemical oscillator
- High complexity
- Well-known example of a excitable medium
- Good correspondence between simulation and experiment
- Simplification of complex systems to few variables is feasible







simulation

experiment

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Finite Element Method

• Strong form (e.g Poisson-eq):

$$-\nabla^2 u = f$$

- Division of domain $\boldsymbol{\Omega}$ in M subdomains (meshing)

$$\Omega = \sum_{j=1}^{M} \Omega_{j}$$

• Weak form:

$$\int_{\Omega} v \nabla^2 u = \int_{\Omega} v f \longrightarrow \int_{\Omega} \nabla v \nabla u = \int_{\Omega} v f$$

• Sobolov space H:

$$v \in H_0^1(\Omega) : \int_{\Omega} [(v')^2 + v^2] d\Omega < \infty$$

$$v, u \in H_0^1(\Omega) : \int_{\Omega} v \nabla^2 u = \int_{\Omega} v f \quad \forall v \quad \frown \quad -\nabla^2 u = f$$

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 Ω_{i}

node n_k

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Finite Element Method

• Basis of H, e.g. linear:

$$\Phi_{i}(x, y) = \alpha_{i} + \beta_{i} x + \gamma_{i} y \quad , \quad \Phi_{i}(n_{k}) = \delta_{ik}$$

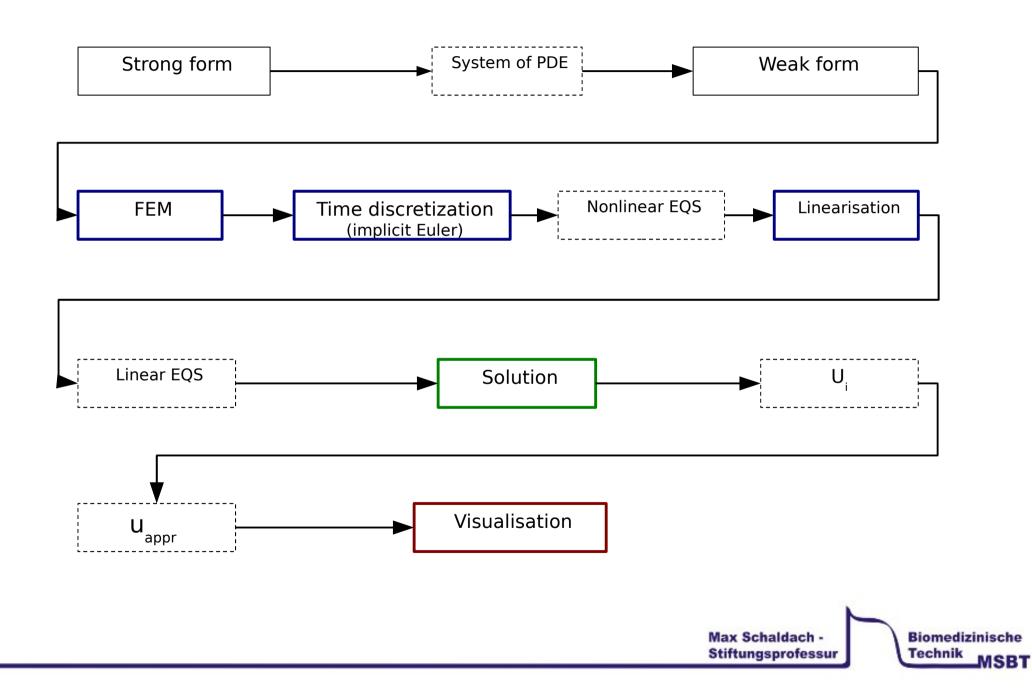
- Approximation of u: $u_{appr} = \sum_{i=1}^{N} u_{i} \Phi_{i}$ • Calculation of u;: $\sum_{i=1}^{N} u_{i} \int_{\Omega} \nabla \Phi_{i} \nabla \Phi_{j} = \int_{\Omega} f \Phi_{j} \longrightarrow \sum_{i=1}^{N} u_{i} A_{ij} = b_{j} \longrightarrow \underline{\underline{A}} \cdot \underline{\underline{u}} = \underline{\underline{b}}$
- Solution of linear equation system $\rightarrow u_{i} \rightarrow u_{app}$

 A_{ii}

Max Schaldach -Stiftungsprofessur $\Phi_{r}(n_{\nu})$

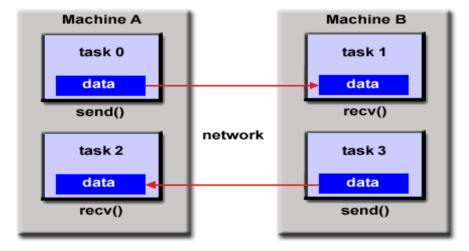
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Numeric Workflow



Software

- Written in C++ under Linux
- Used libraries:
 - PETSC (Portable, Extensible Toolkit for Scientific Computation): LEQ-Solver
 - Hermes3d: FEM
 - VTK/Paraview: Visualization
- Parallelization MPI-2 (SPMD, distributed memory)



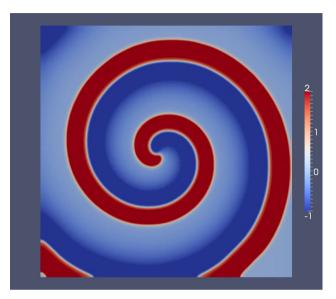
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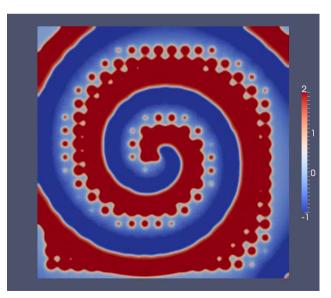
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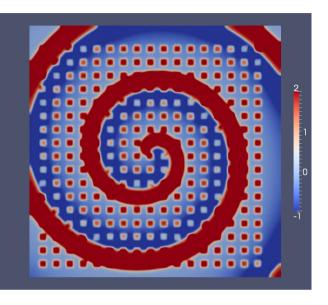
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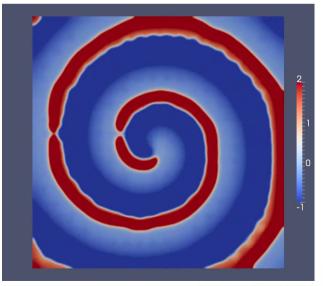
MSBT

Spiral wave on plain domain (600x600x2 nodes), grid stimulation



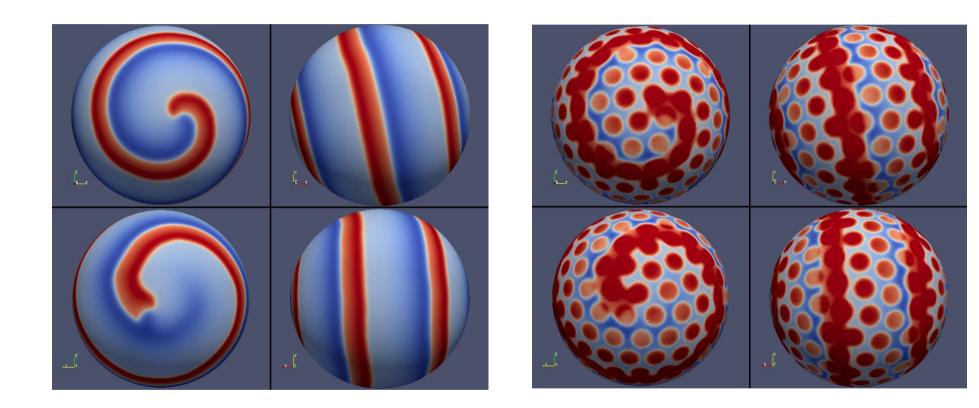




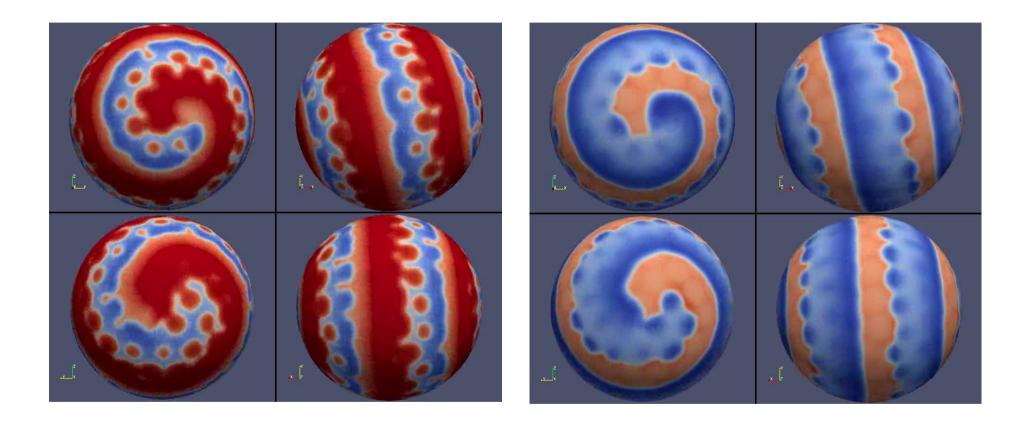


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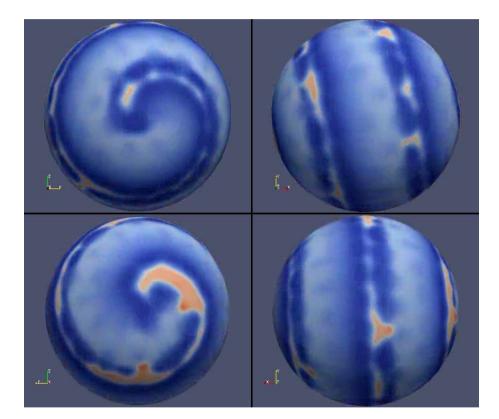
Spiral wave on spherical shell (r=50, d=2, 70000 nodes)

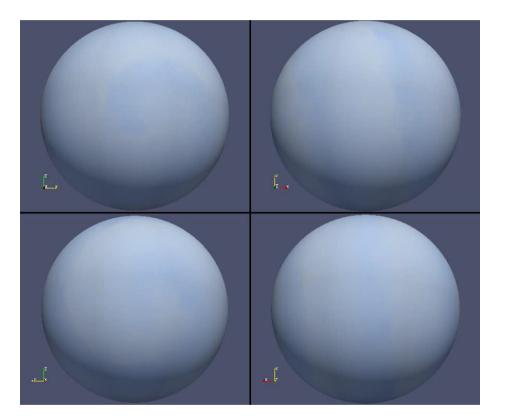


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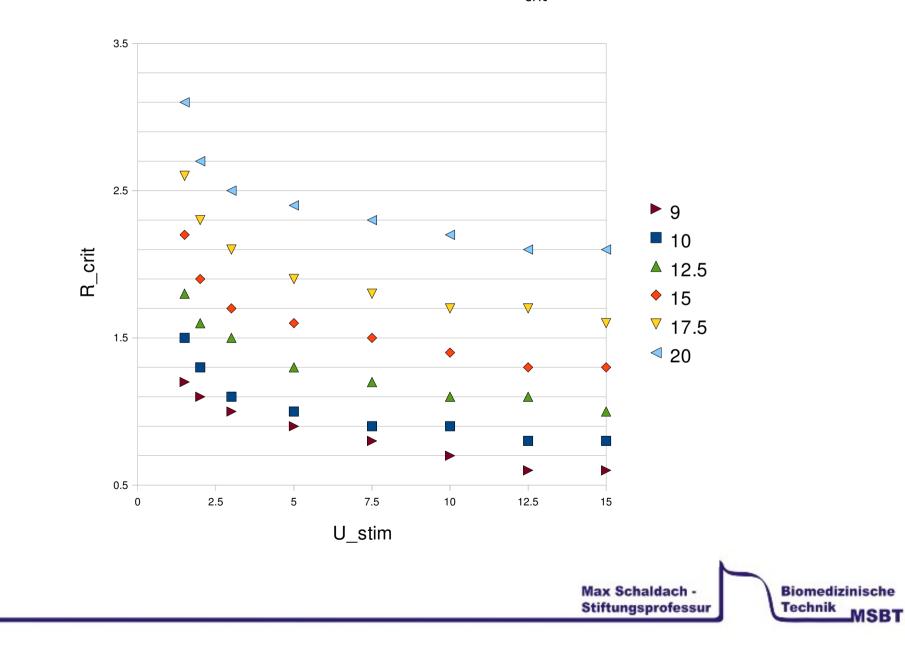
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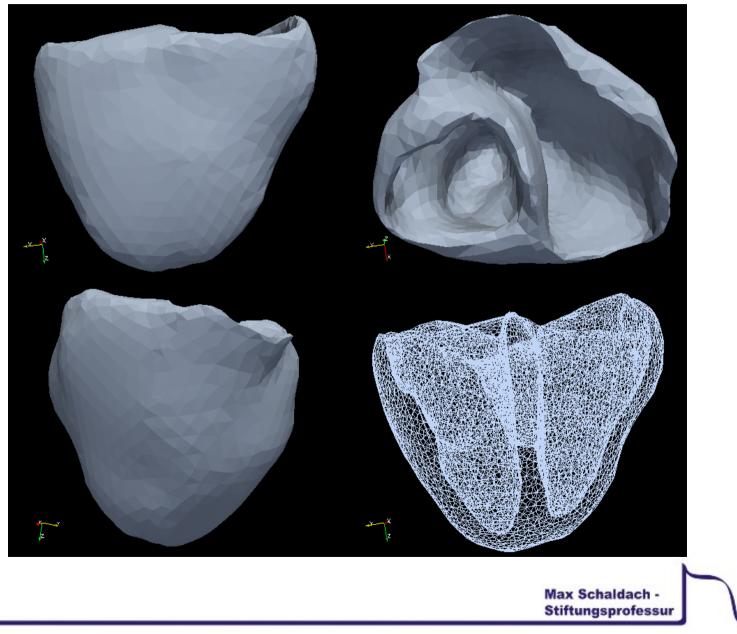


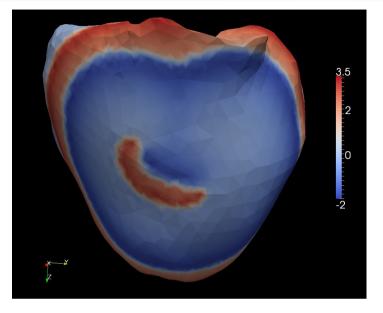
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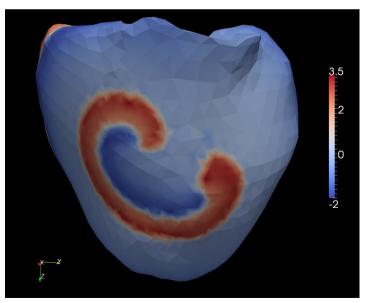
Critical stimulation size $\rm R_{_{crit}}$

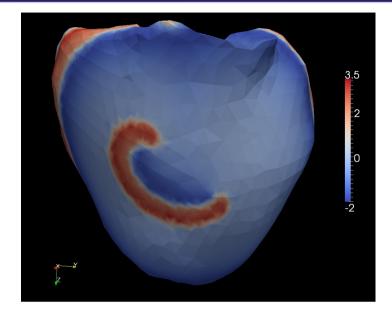


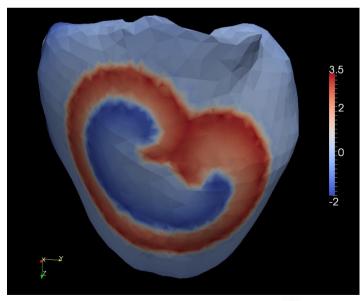
Mesh of human ventricles



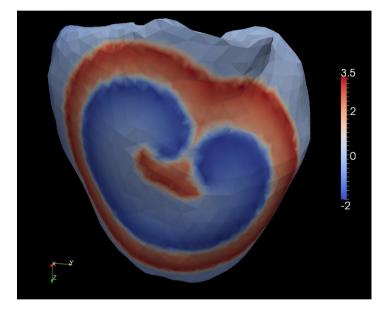


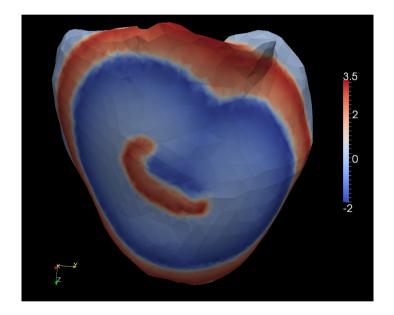






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