

# SILICOFCM

## In Silico trials for drug tracing the effects of sarcomeric protein mutations leading to familial cardiomyopathy- SILICOFCM project

**Prof. Nenad Filipovic**

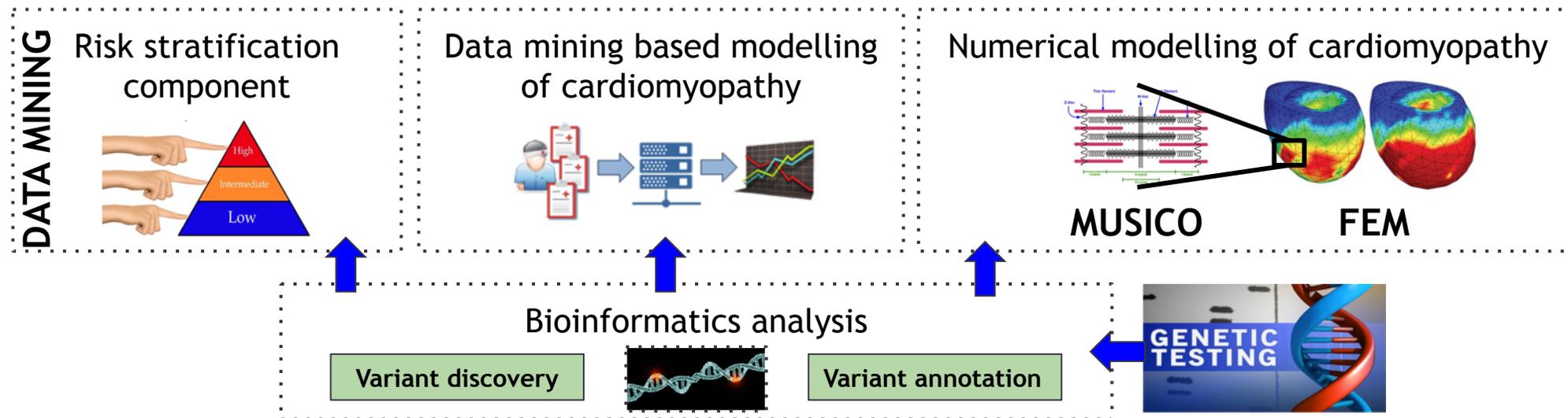
BioIRC Bioengineering Research and Development Center, Kragujevac, Serbia  
University of Kragujevac, Serbia

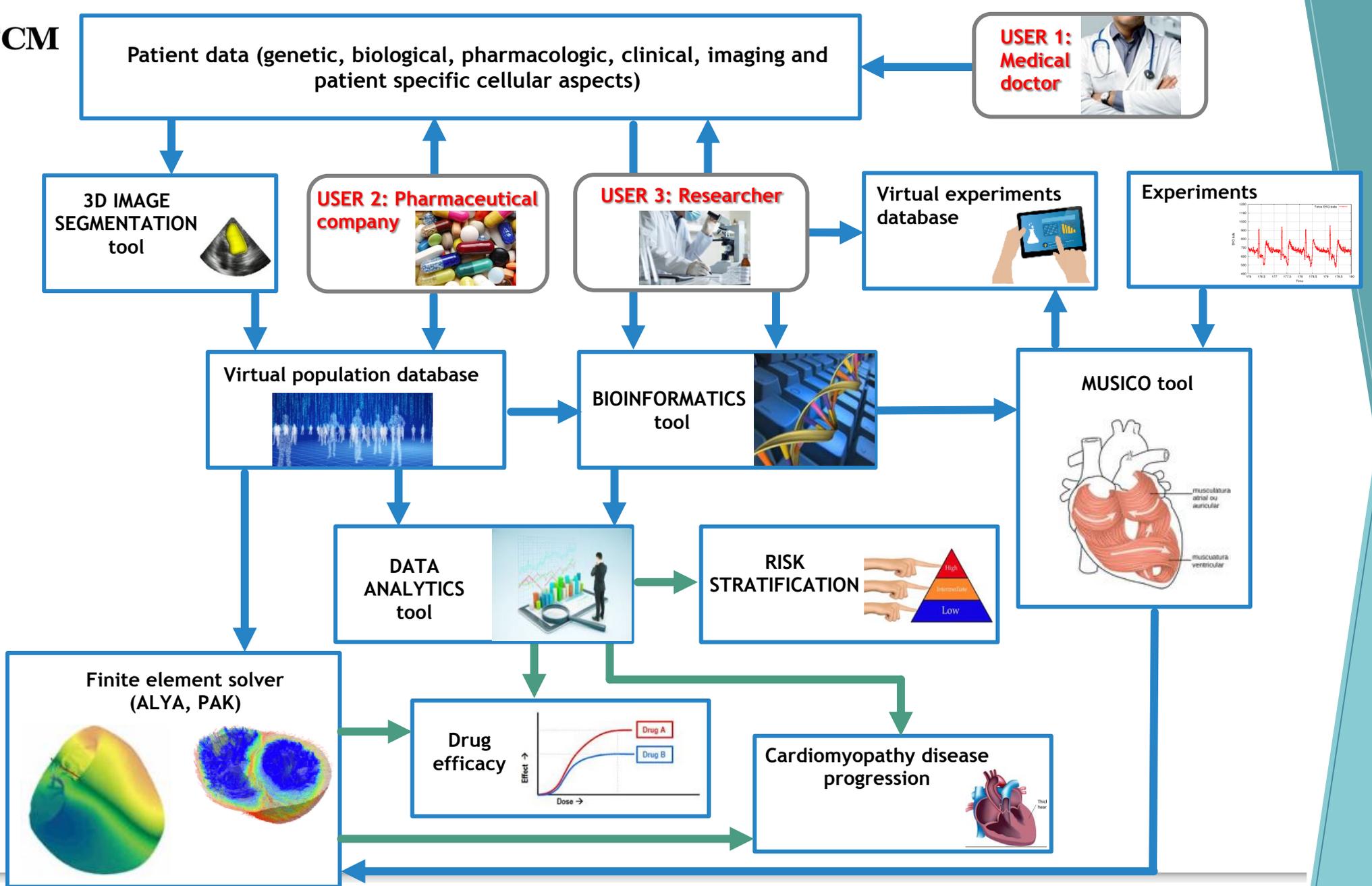
Institute of Engineering and Technology, London: Savoy Place  
September 25-27, 2019

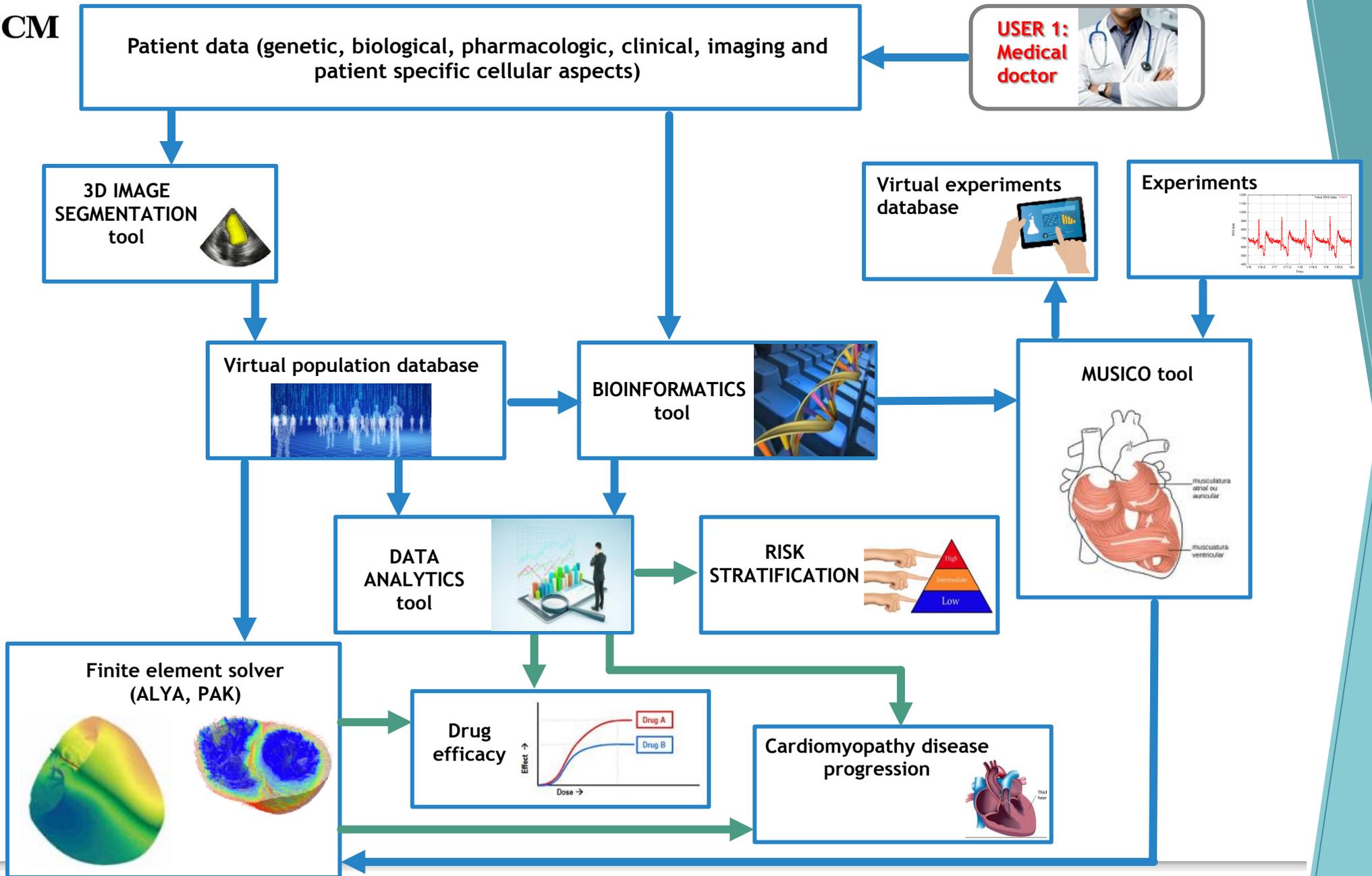


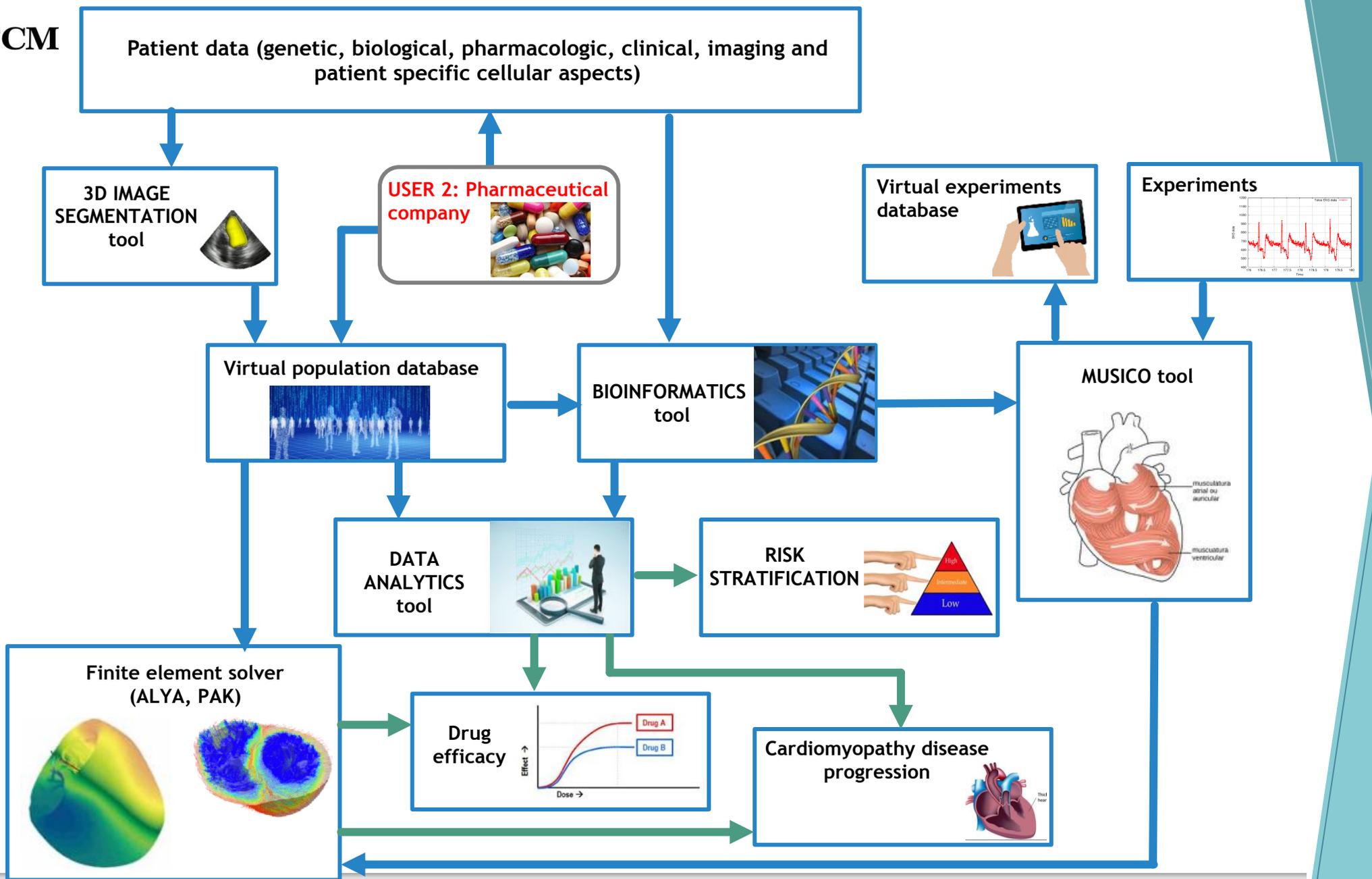
# SILICOFCM Scope

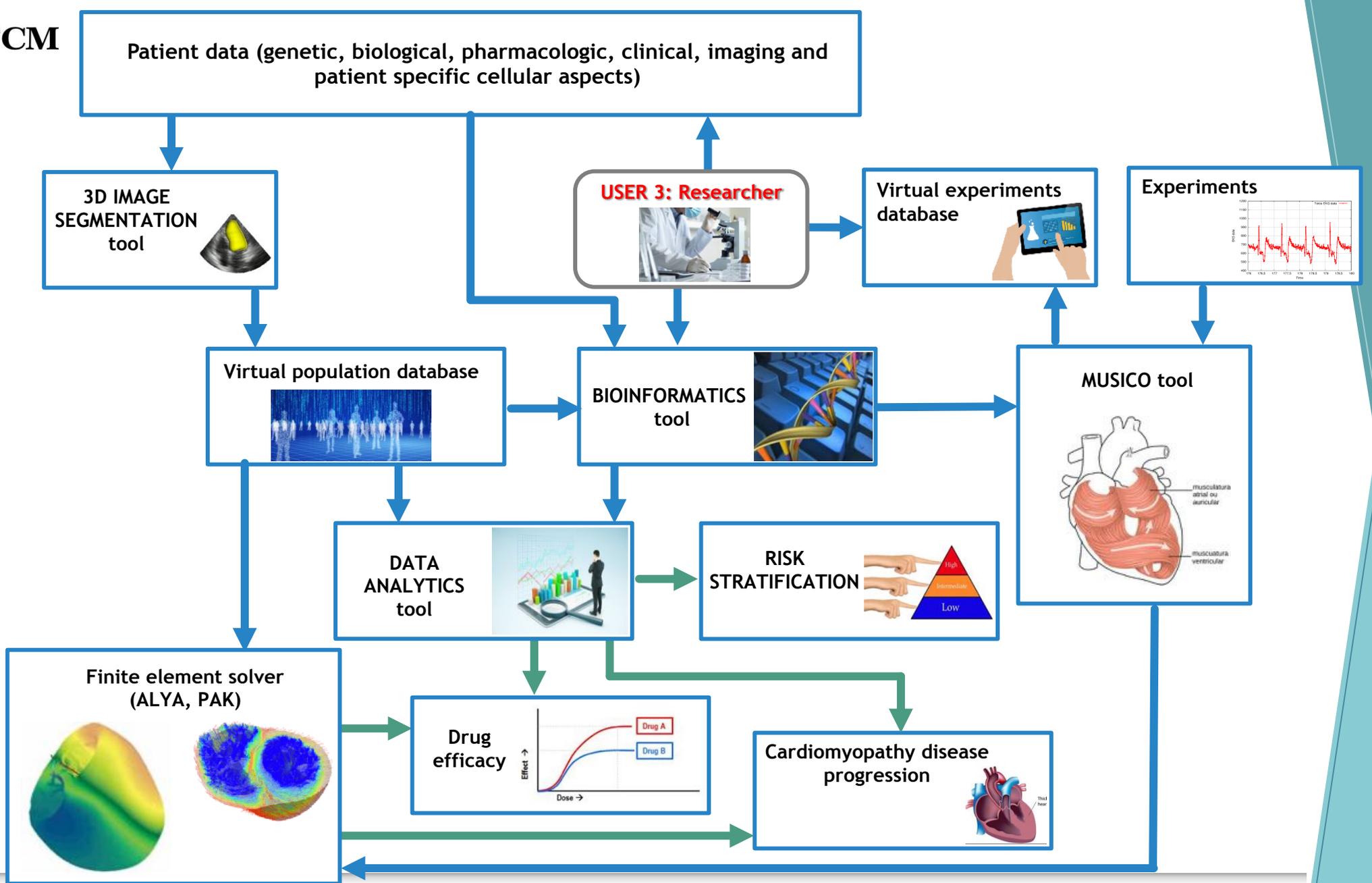
- ▶ SILICOFCM aims to develop a computational platform for *in silico* clinical trials of Familial cardiomyopathies (FCMs) that would take into consideration comprehensive list of patient specific features (genetic, biological, pharmacologic, clinical, imaging and patient specific cellular aspects) capable of **optimizing and testing medical treatment strategy** with the purpose of maximizing positive therapeutic outcome.
- ▶ The SILICOFCM platform is based on the integrated multidisciplinary and multiscale methods for analysis of patient-specific data and development of patient-specific models for monitoring and assessment of patient condition from current through the progression of disease.



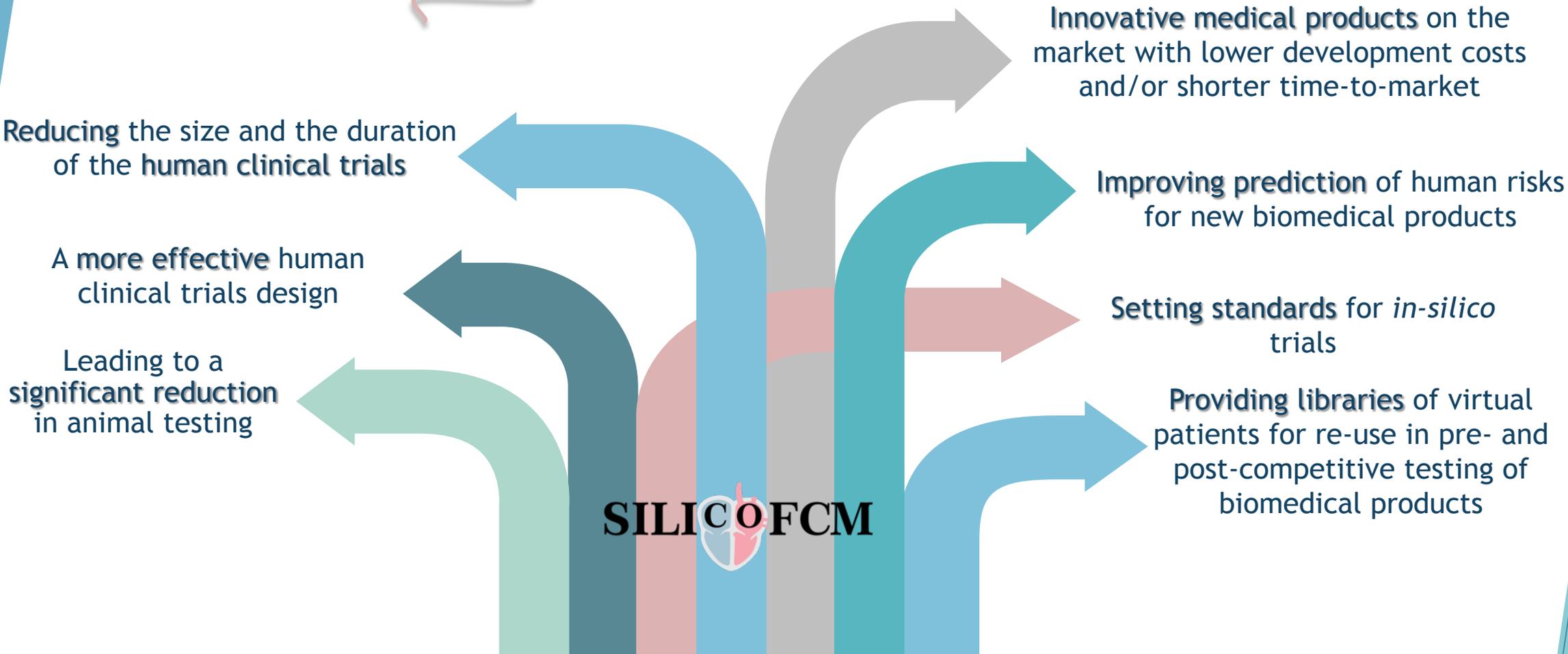
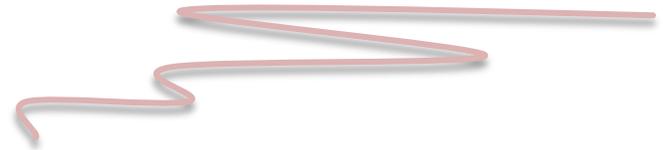




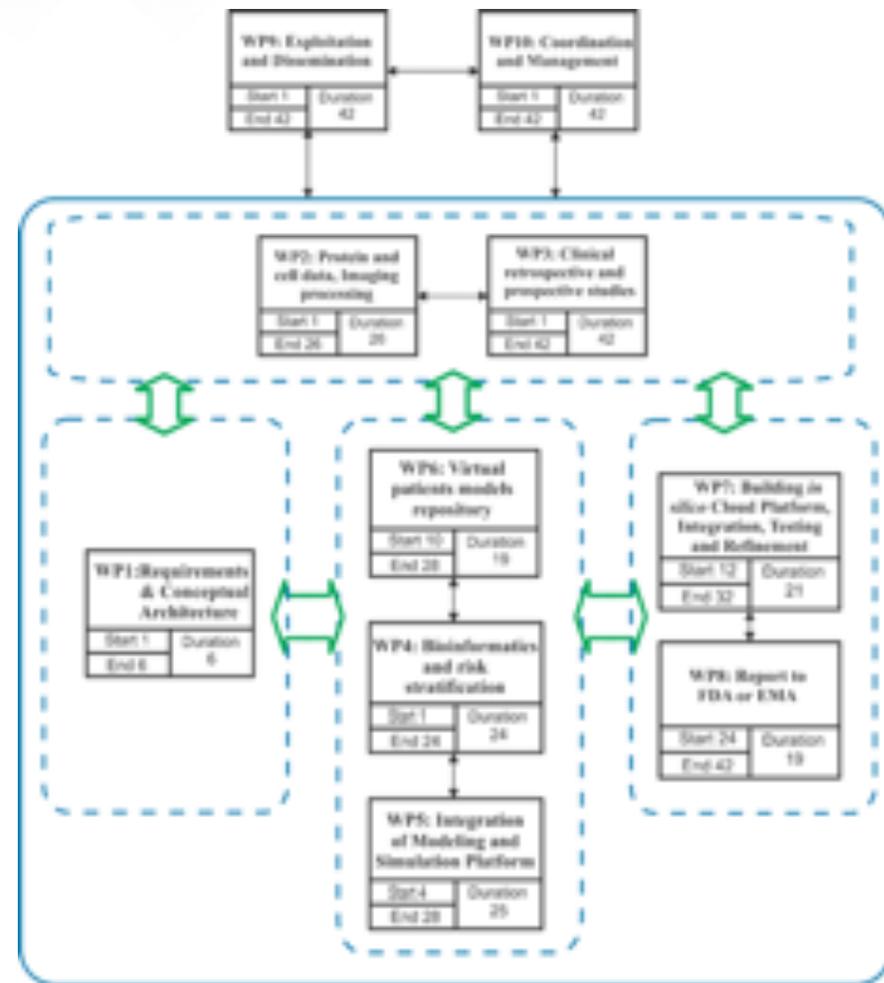
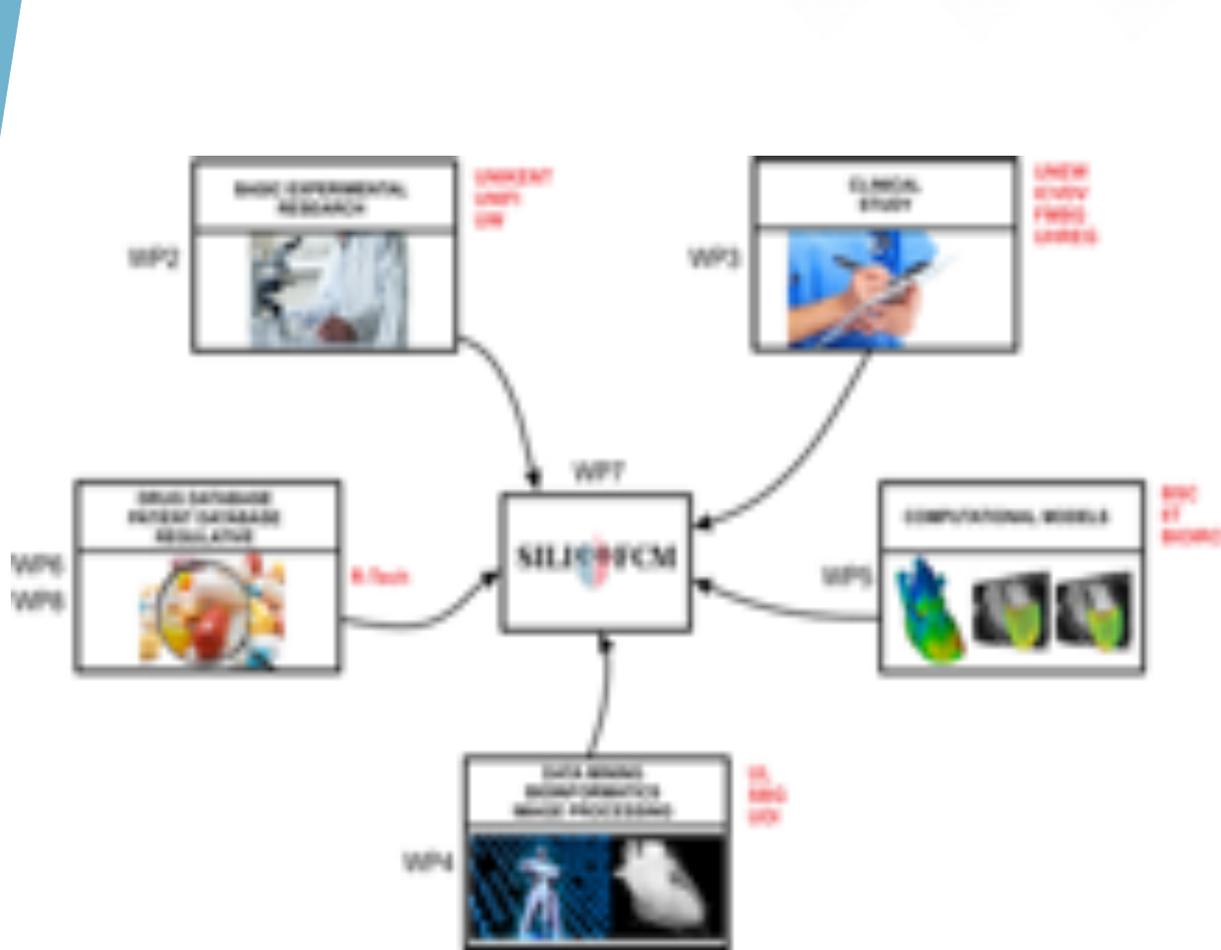




# SILICOFCM Impact



# WPs Interaction

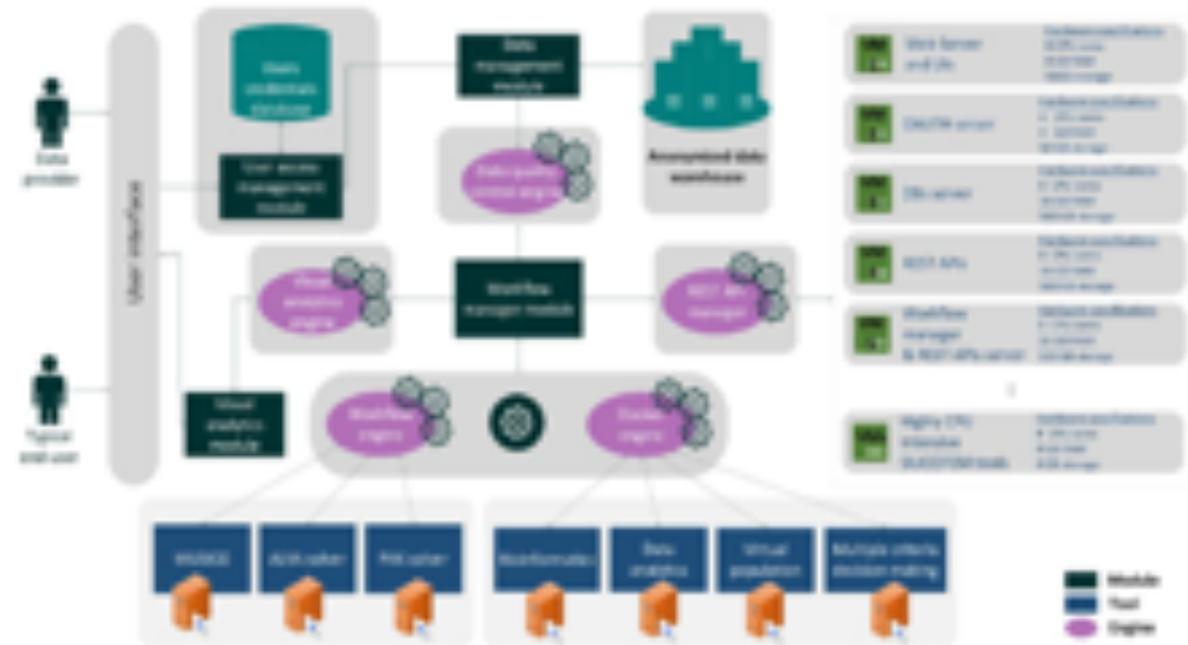
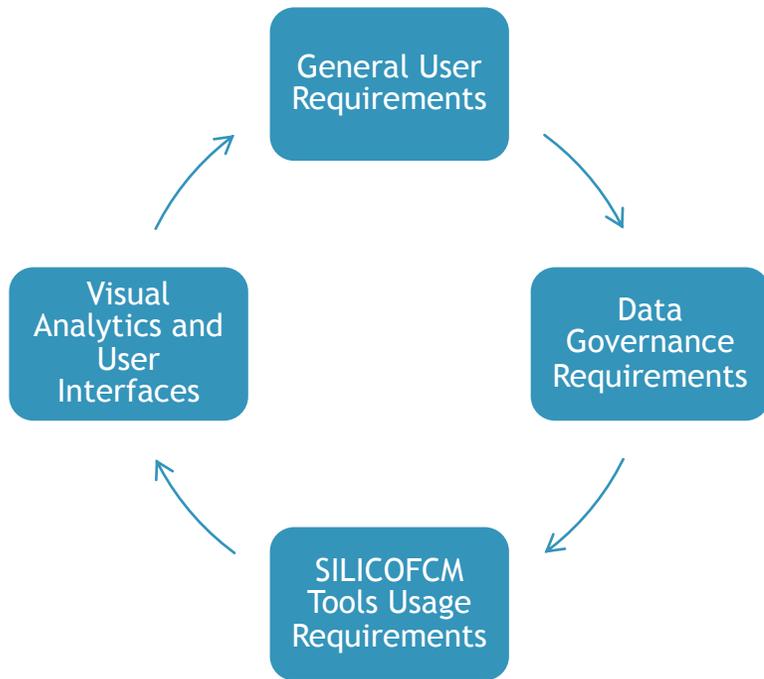


# WP1 Requirements & Conceptual Architecture

Leader BIORIC M1-M6

## Main objective

- ▶ To collect and analyse the state-of-the-art technologies, user requirements and hardware requirements for innovative SILICOFCM Reference Architecture



# WP2 Protein and cell data, Imaging processing

Leader UNIKENT

M1-M26

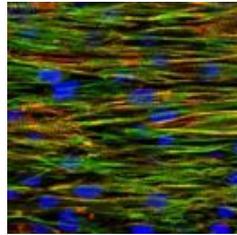
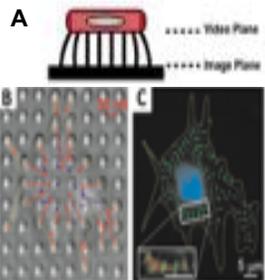
## Main objective

- ▶ To provide protein and cell data as well basic physiological experiments for heart disease
- ▶ To acquire medical images

3D engineered  
Tissue-level  
mechanics  
heart tissue (EHT)



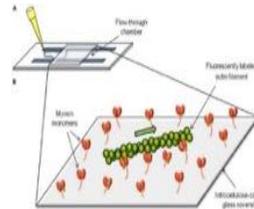
Cell  
Contraction/Relaxation  
(micropost arrays)



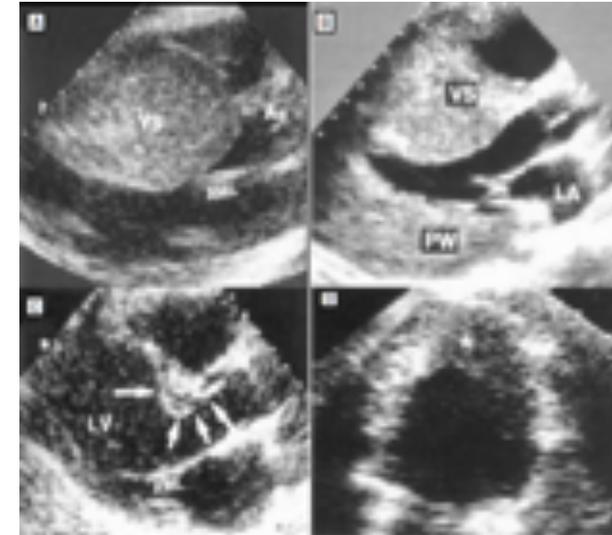
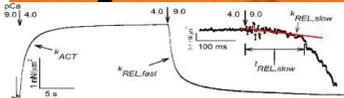
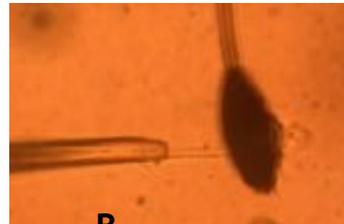
Cell contractility/Ca<sup>2+</sup> transients



Actin-Myosin mechanics and  
energetics (in vitro  
motility/micro-needle assay)



Isolated Myofibrils  
contraction/relaxation kinetics



**WP3** Clinical retrospective and prospective studies

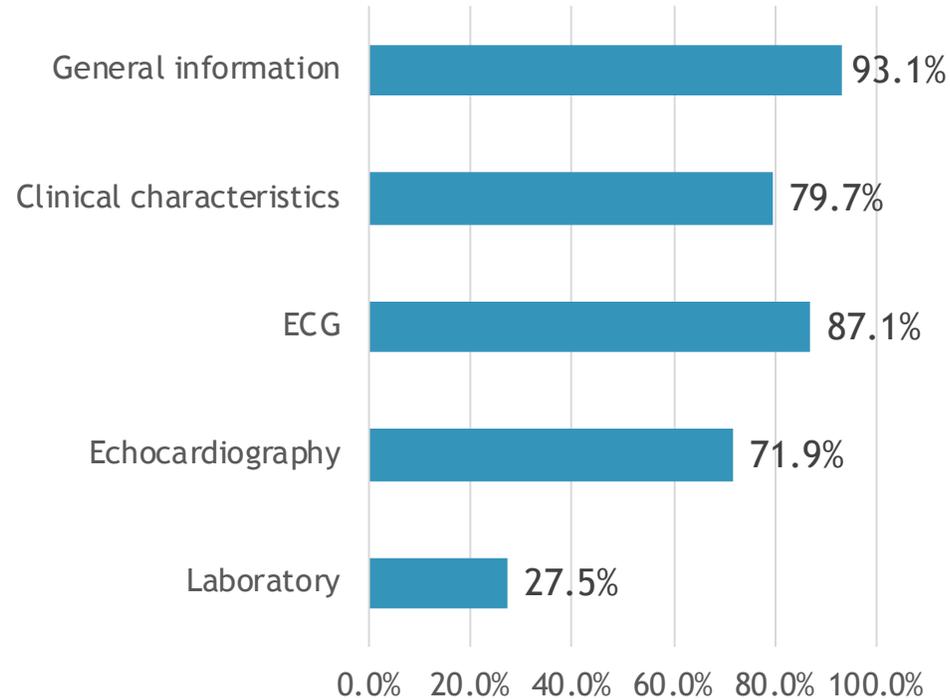
Leader UNEW M1-M42

**Main objective**

- ▶ To create the clinical data for fitting and validation of the mechanical models (WP5), training and testing of the risk stratification tool (Task 4.4) and the data mining model for prediction of cardiomyopathy outcome (WP7)

**Data completeness**

- ▶ ICVDV 90.3%
- ▶ UNIFI 94.4%
- ▶ UNEW 78.7%
- ▶ UHREG 67.9%



WP4

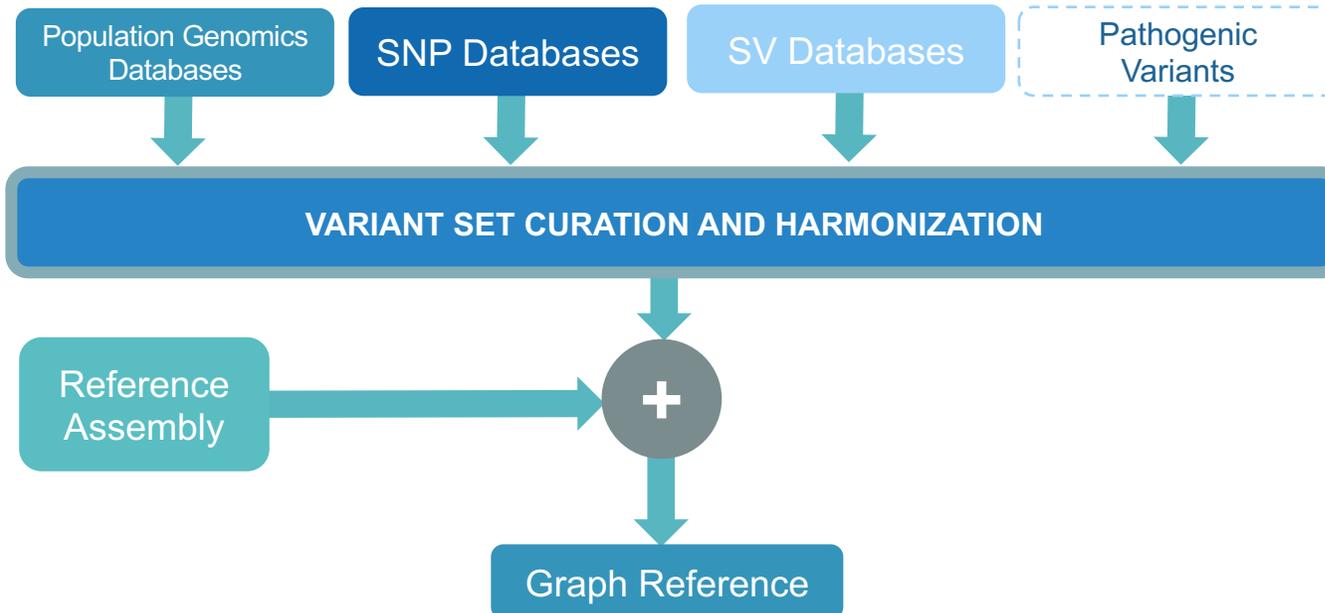
# Bioinformatics and risk stratification

Leader SBG

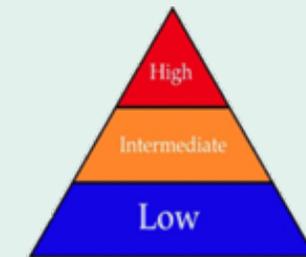
M1-M24

## Main objective

- ▶ To put in place the genomic component in the SILICOFCM platform and integrate it with the SILICOFCM sub-components
- ▶ To develop the cardiomyopathy risk stratification system



Learning performance evaluated with cross-validation ROC analysis approach and other performance measures



risk stratification outcomes for unseen/new patients



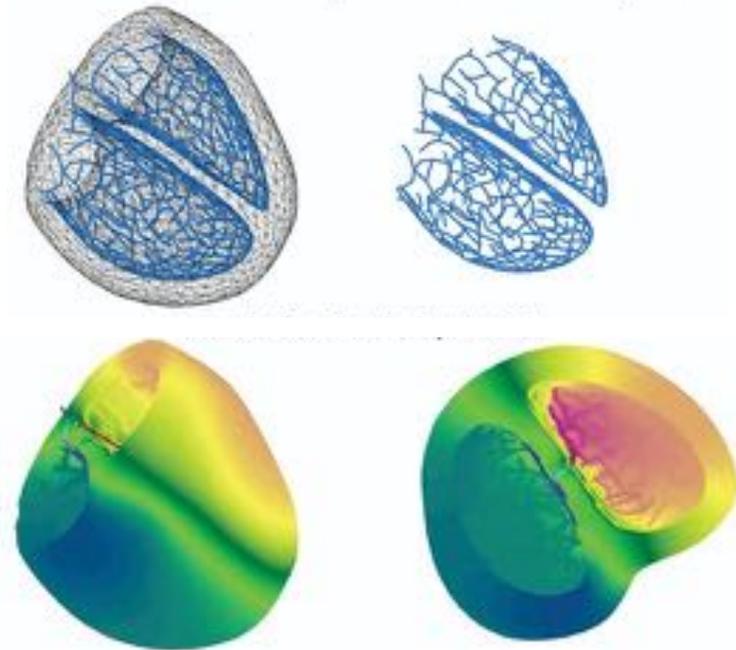
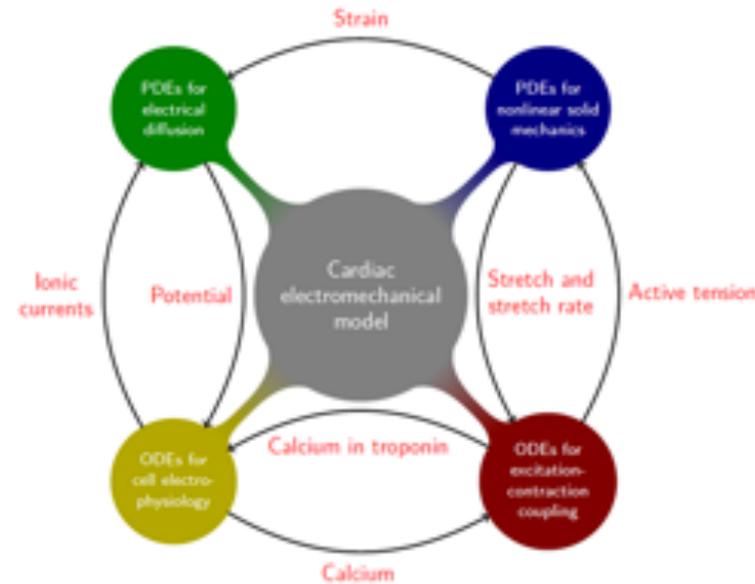
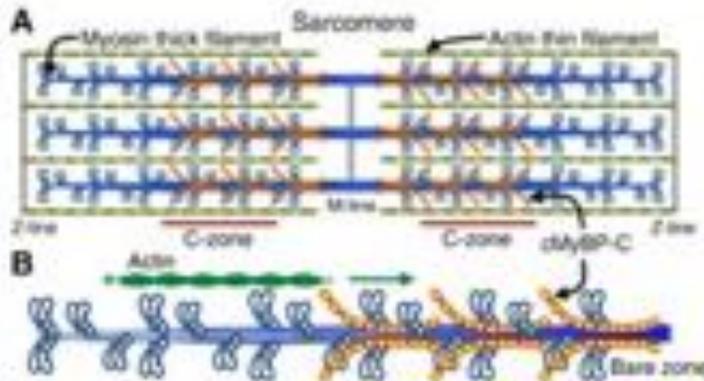
# WP5 Integration of Modelling and Simulation Platform

Leader IIT

M4-M28

## Main objective

- ▶ To link the data from molecular interactions to the whole organ function by coupling Bioinformatics tool, MUSICO and FE solvers (ALYA and PAK)



# WP6 Virtual patients models repository

Leader UL

M10-M28

## Main objective

- ▶ To develop virtual FCM patients models repository and perform pattern identification from heterogeneous data by using data mining algorithms

## SILICOFCM virtual population model - **DRAFT**



WP7

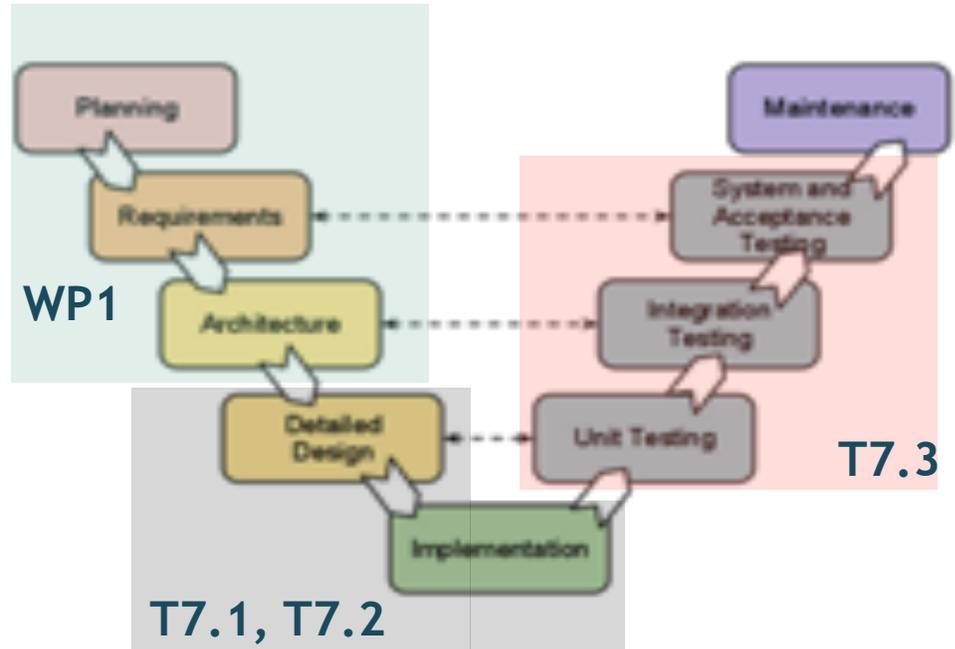
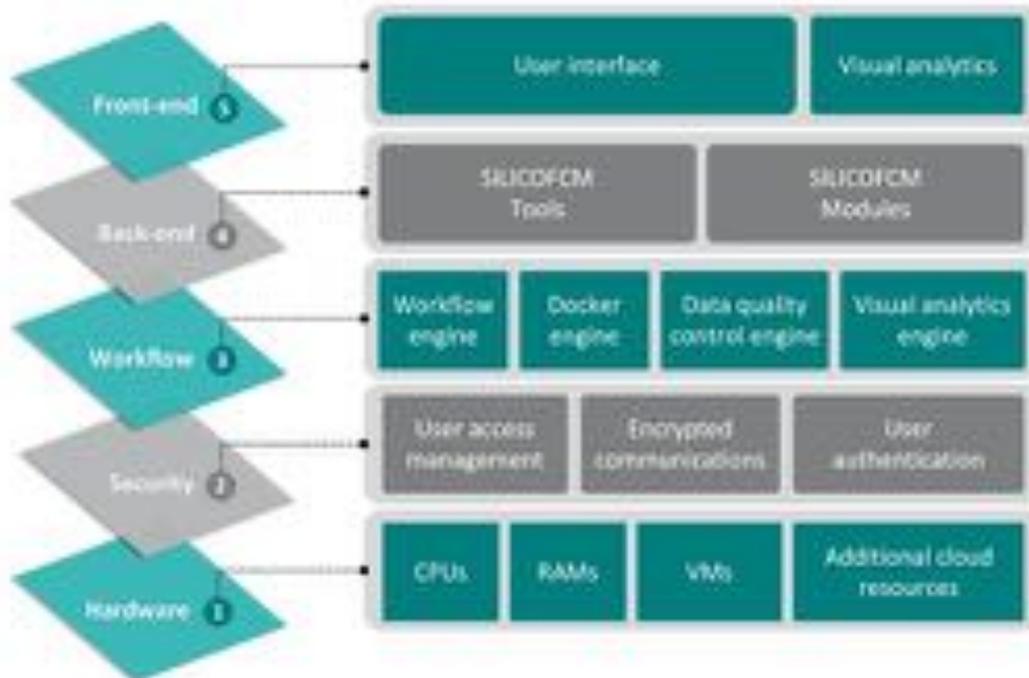
# Building *in Silico* Cloud Platform, Integration, Testing and Refinement

Leader UOI

M12-M32

## Main objective

- ▶ To integrate the SILICOFCM subsystems into the platform guaranteeing a smooth, secure, and standard integration



WP8

# Report to FDA or EMA

M24-M42

Leader R-Tech

## Main objective

- ▶ To perform regulatory approval processes of the project results towards EMA or FDA

### Task 8.1

Development workflow assistant for EMA/FDA approval

Leader BioIRC, M24-M36

### Task 8.2

Set up R&amp;D computation pipelines for drug testing

Leader IIT, M24-M36

### Task 8.3

Interface drug database

Leader UOI, M24-M40

### Task 8.4

Development report tool

Leader R-Tech, M24-M42



WP9

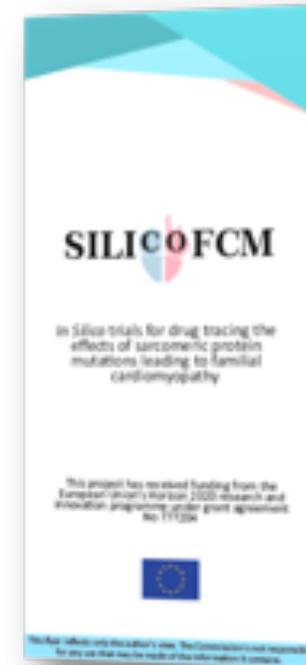
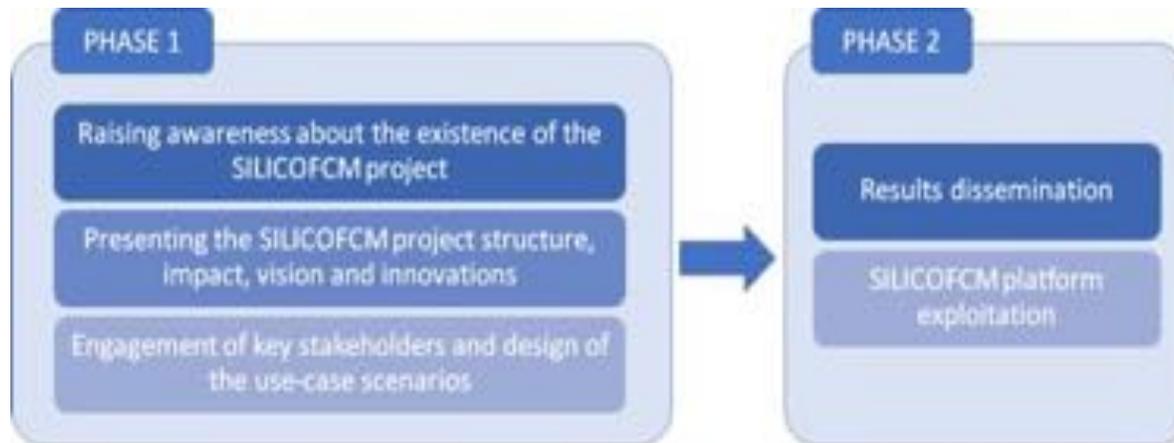
# Exploitation and Dissemination

Leader BioIRC

M1-M42

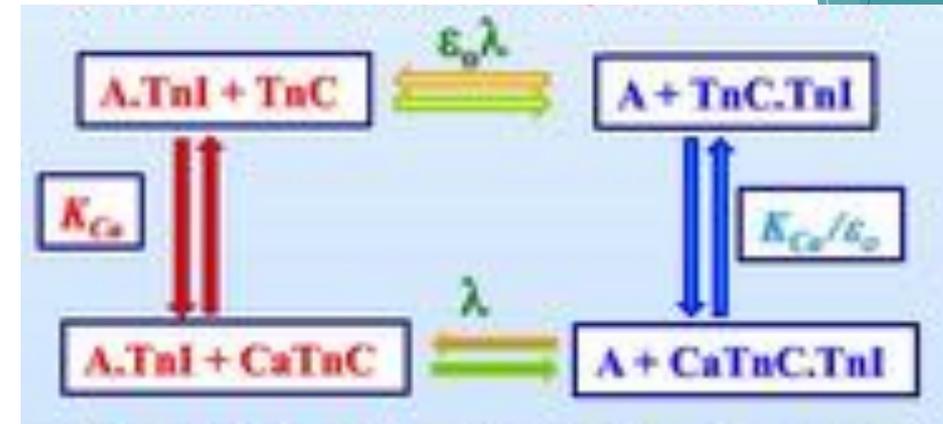
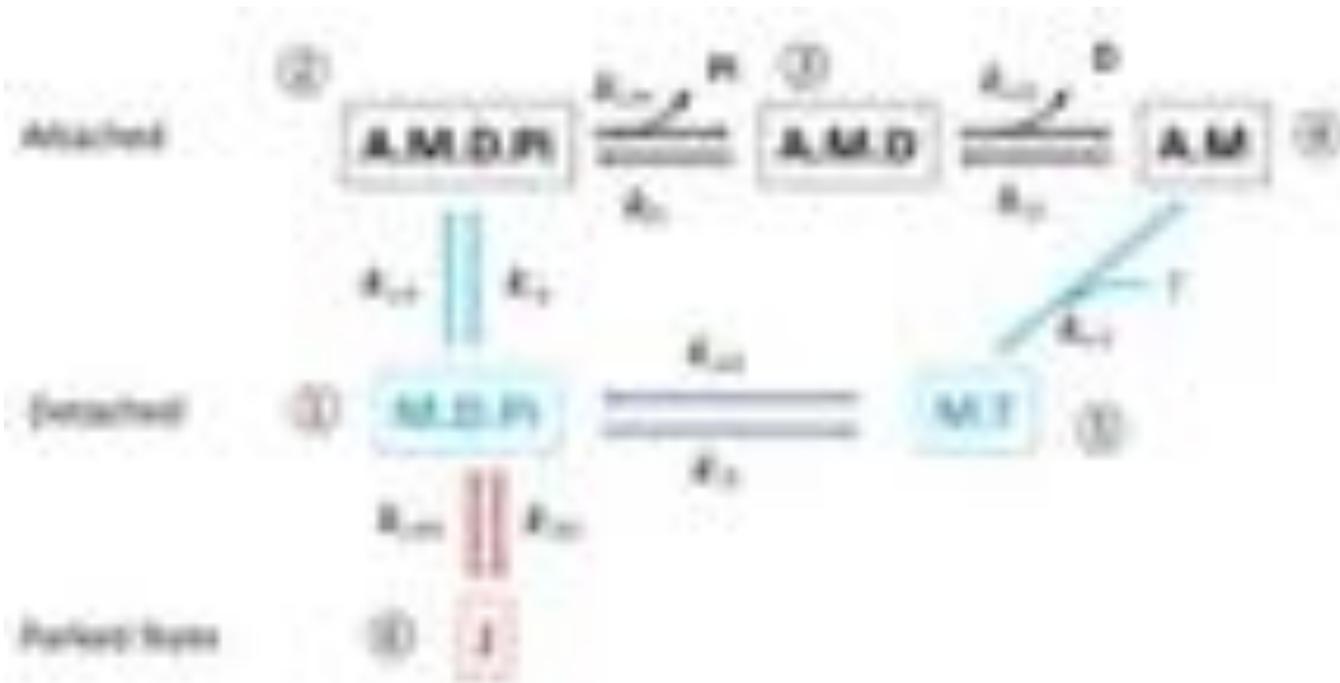
## Main objective

- ▶ To promote the widespread utilisation and exploitation of the project results

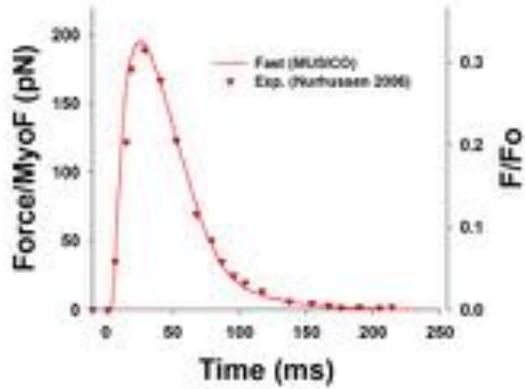


# MUSICO: Six State Crossbridge Model & Calcium Kinetics with TnC

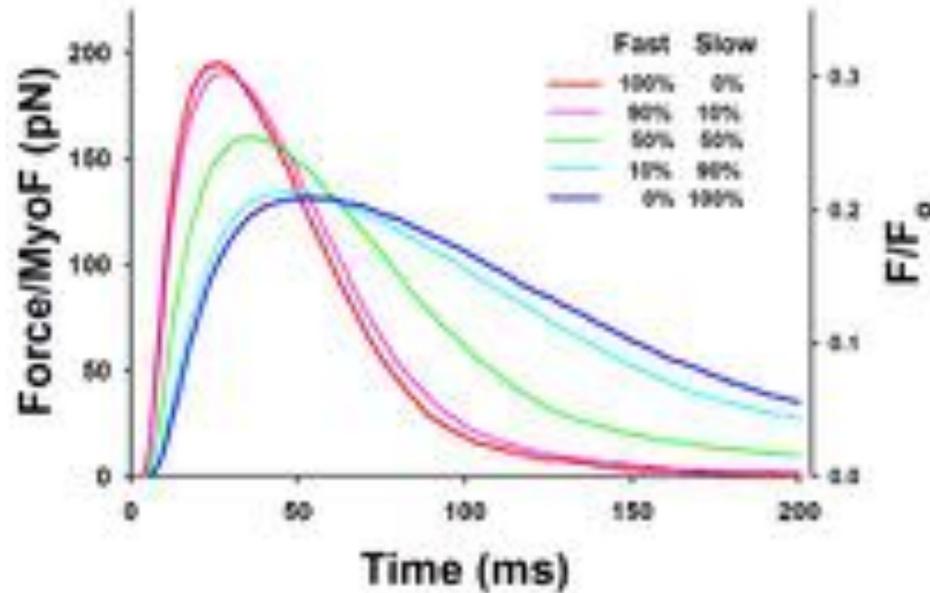
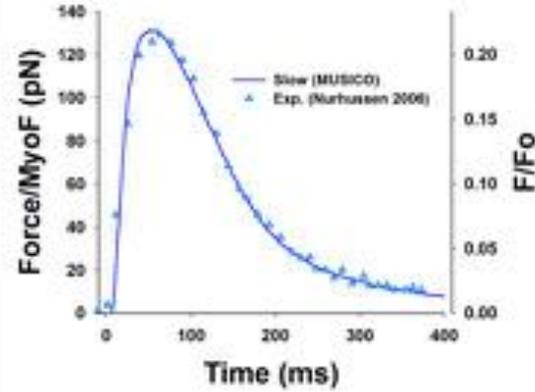
*Cardiac Muscle*



*Fast Twitch Contraction*

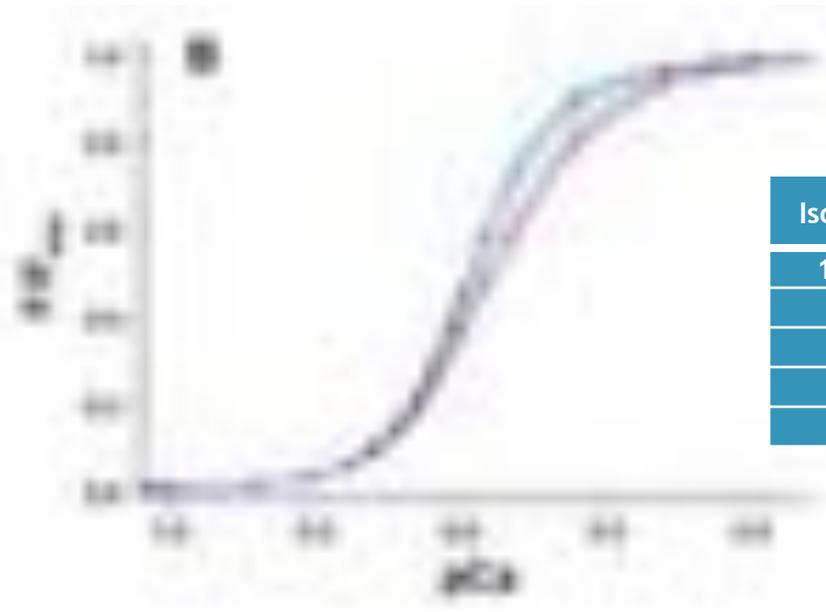
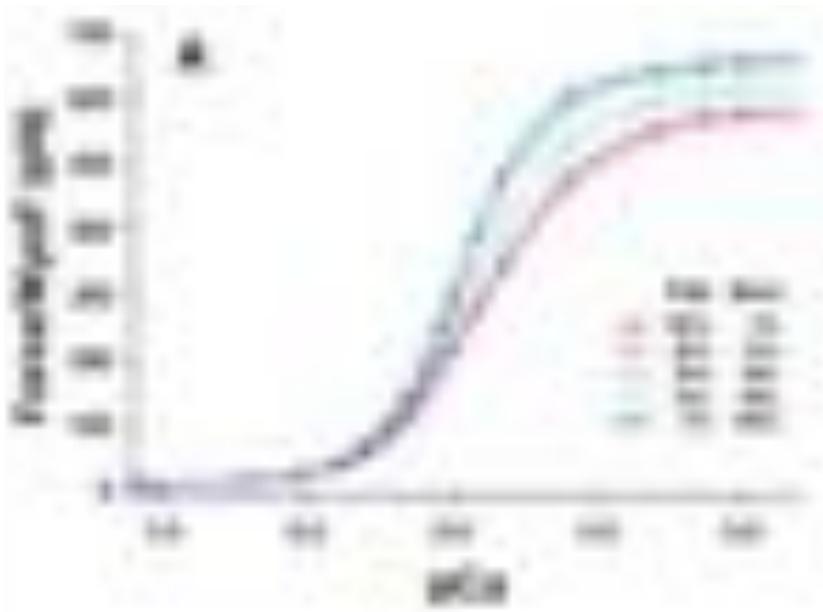


*Slow Twitch Contraction*



- ❖ The calcium transient:
- ❖ Peak of 1.6  $\mu\text{M}$  at 6 ms
- ❖ Decays to low calcium concentrations at about 35 ms.

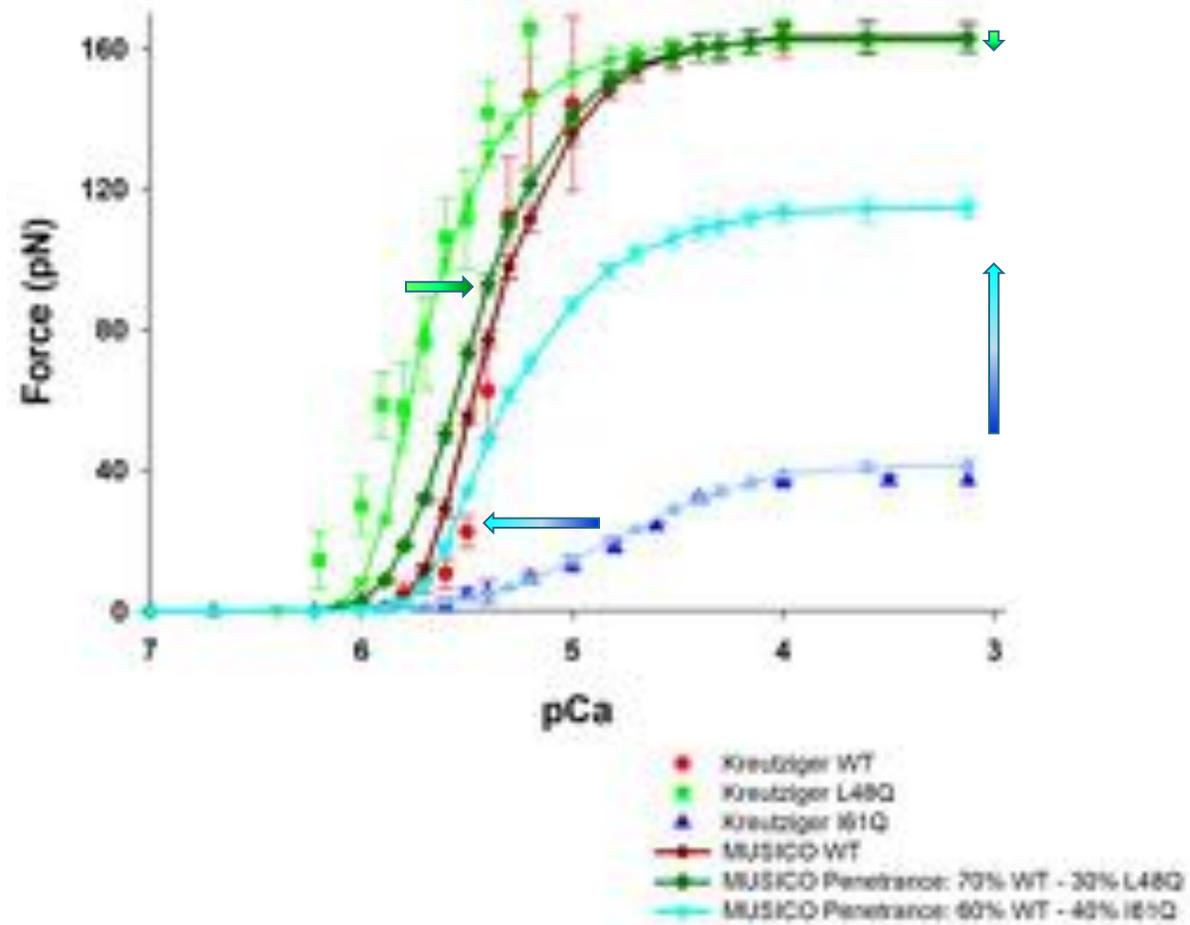
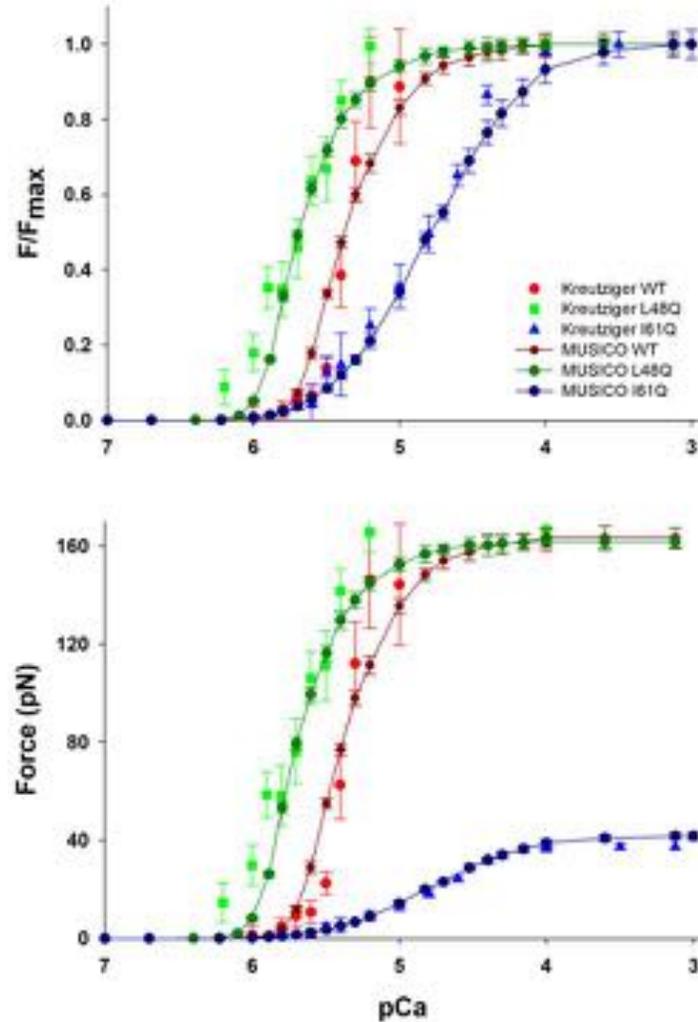
## Effect on Hill Coefficient and Calcium Sensitivity



Isoform Mixture		Hill Coeff	pCa <sub>50</sub>
100% $\alpha$	0% $\beta$	2.302	5.890
90% $\alpha$	10% $\beta$	2.342	5.899
50% $\alpha$	50% $\beta$	2.531	5.933
10% $\alpha$	90% $\beta$	2.807	5.966
0% $\alpha$	100% $\beta$	2.889	5.973

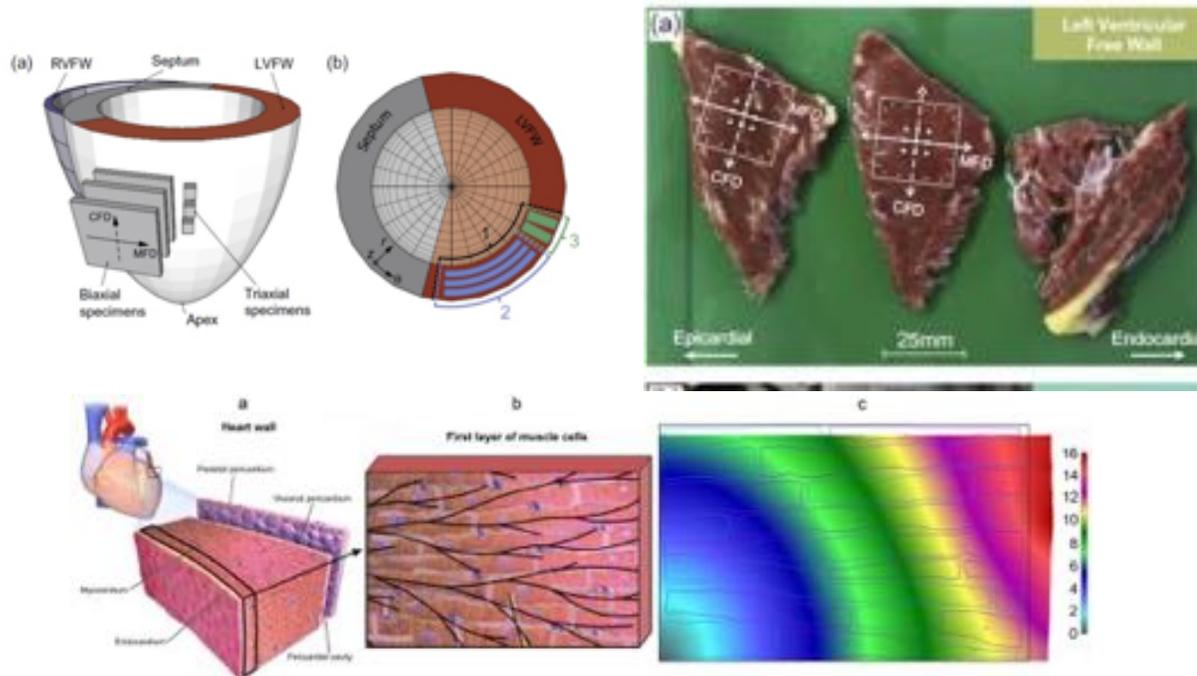
Fit to Kreuziger Experiments - Rat Trabeculae 100% Mutants

Kreuziger - Predicted Penetrance (Rat Trabeculae)

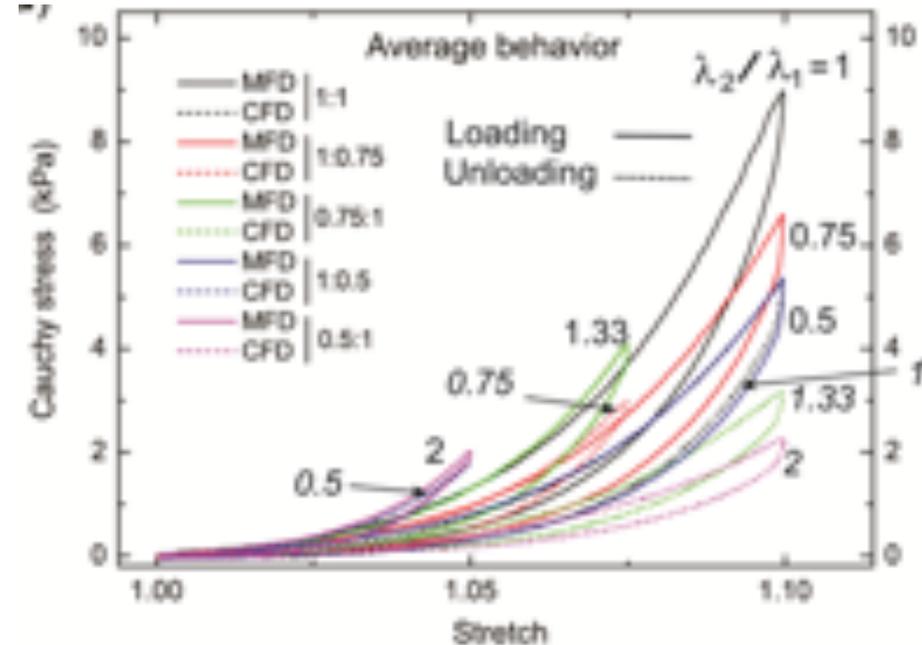


# Upgrade FE biomechanical simulation PAK Solver

- Implementation and testing of Heart mechanical model, according to (Holzapfel et al 2015):



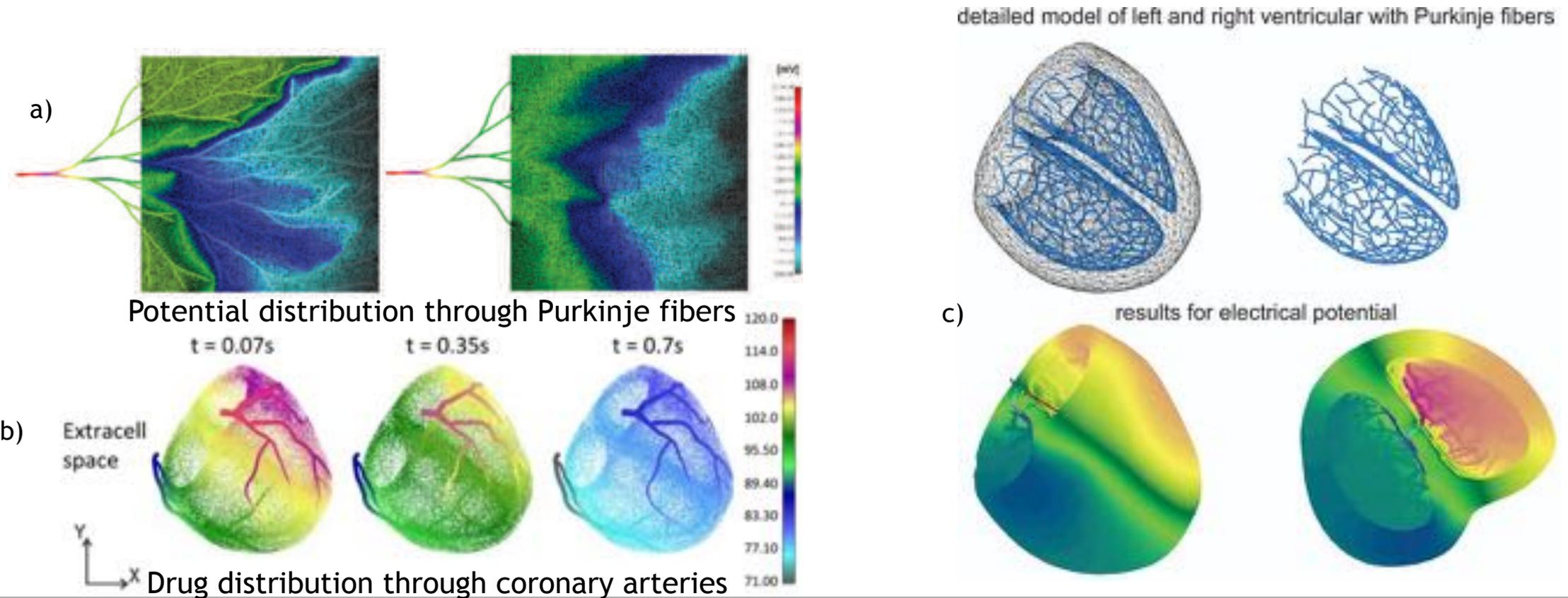
Field of displacements in heart tissue due to uniaxial straining



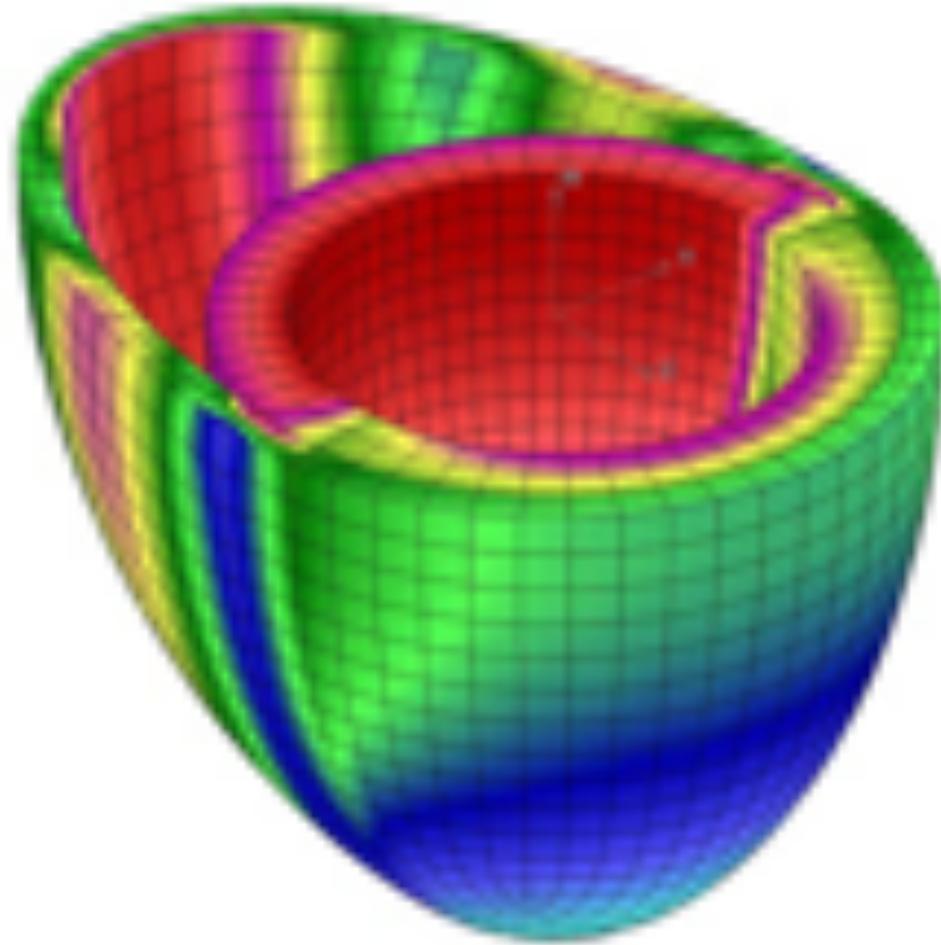
Experimental curves with hysteresis for biaxial loading of myocardium tissue

# Upgrade FE biomechanical simulation PAK Solver

- Application of CSFEM to real heart model (coupled electrophysiology and mechanics)

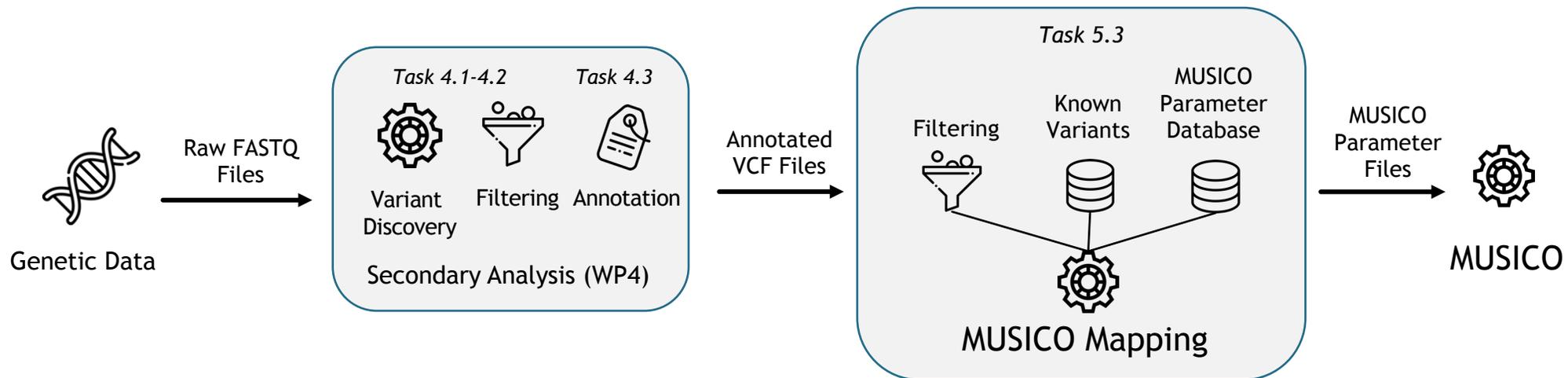


# Fluid-solid interaction-PAK

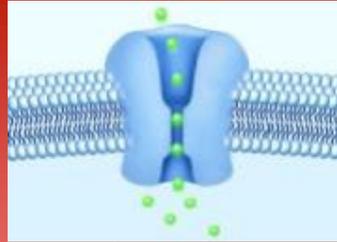


# Linking bioinformatics and MUSICO

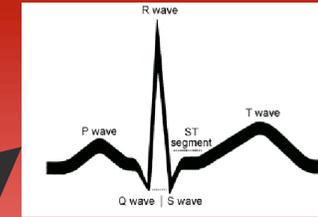
- ▶ The identified variants as part of Task 4.2 will be annotated with the annotation pipeline developed as part of Task 4.3.
- ▶ The annotated variants will be then filtered once more to reduce the variant call sets to variants of interest to MUSICO platform.
- ▶ The filtered mutations of interest then will be converted to a set of physiological parameters which will be the input for the MUSICO Platform.
- ▶ The associated tools were prepared during first 12 months, and corresponding databases will be populated and tested with known variants and experimentally determined associated parameters in the upcoming 6 months.



## Electrophysiology

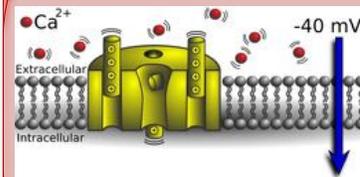


ion channels



ventricles

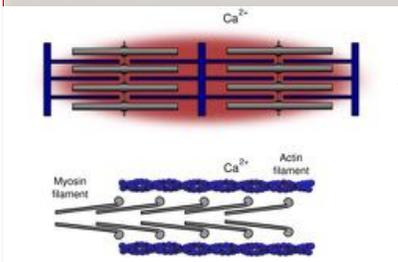
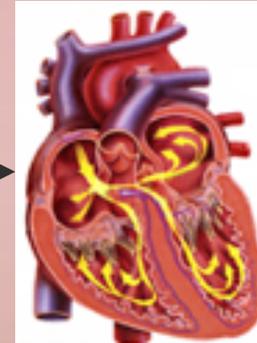
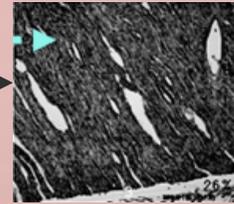
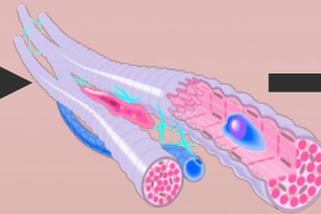
electrocardiogram



Ca<sup>2+</sup> ion channels

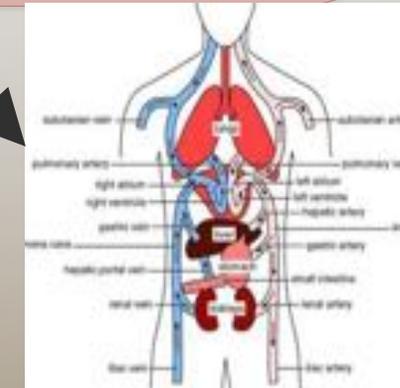
whole cell

myocardium



sarcomere

## Biomechanics



circulation

# Fluid-Electro-Mechanic Cardiac Model - The Heart as a Multi-Physics Coupled System



Electrophysiology:

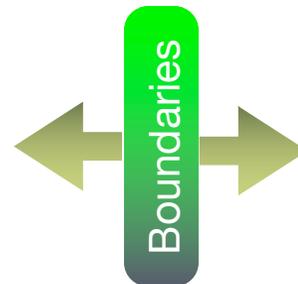
Linear anisotropic (fibers) diffusion + non-linear source terms

Rogers-McCulloch, O'Hara-Rudy, Ten Tusscher-Panfilov, Fenton-Karma,...



Electro-mechanical coupling, via Ca<sup>+</sup> transient:

Hunter & McCulloch 1998, Land-Niederer 2017, Rice-Winslow 2006



Large deformations + non-linear, orthotropic material models:  
Holzapfel and Ogden 2009

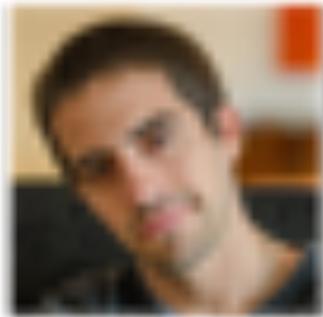
ALE + Immersed Boundaries

Navier -Stokes for Incompressible Flow

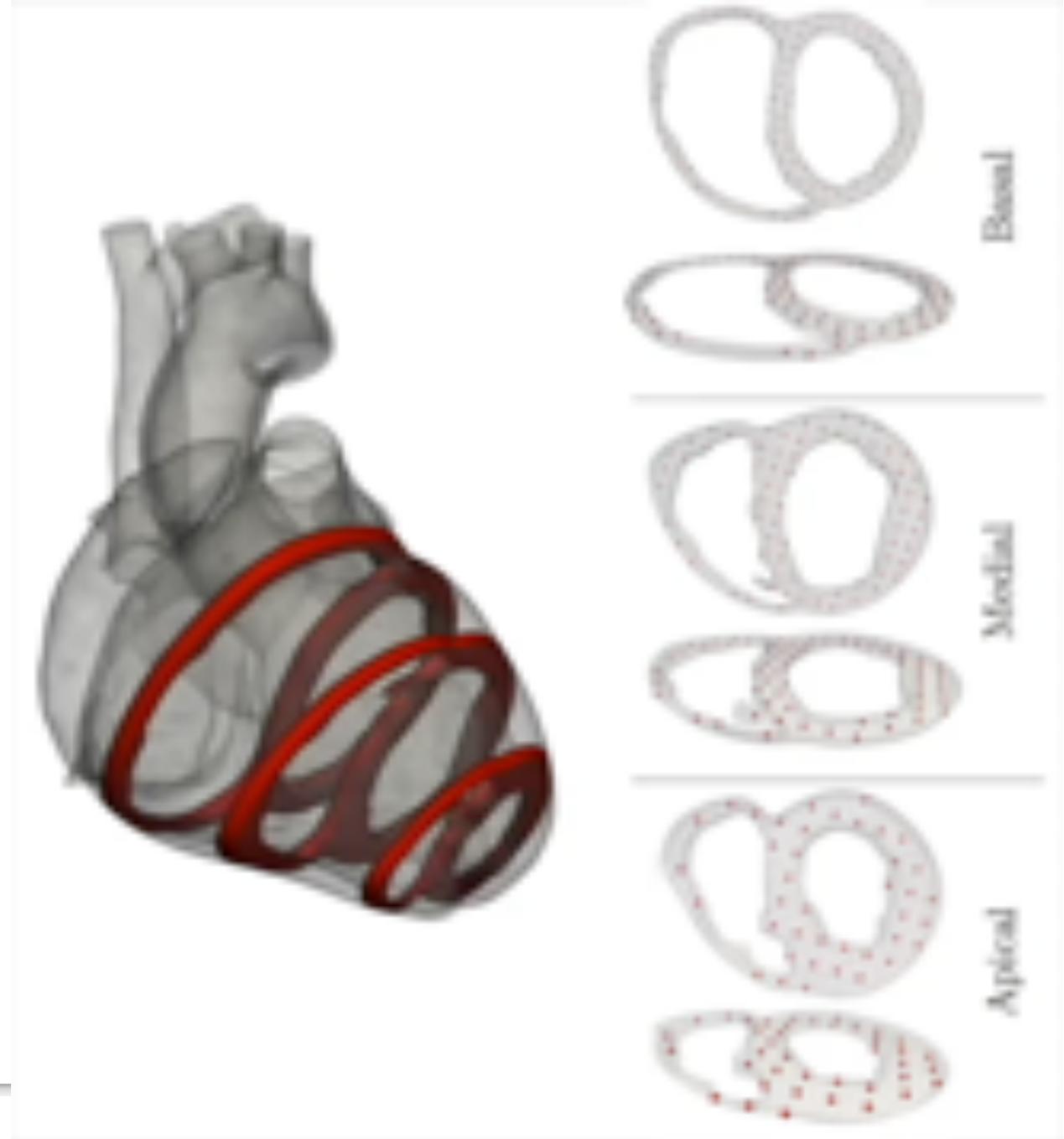
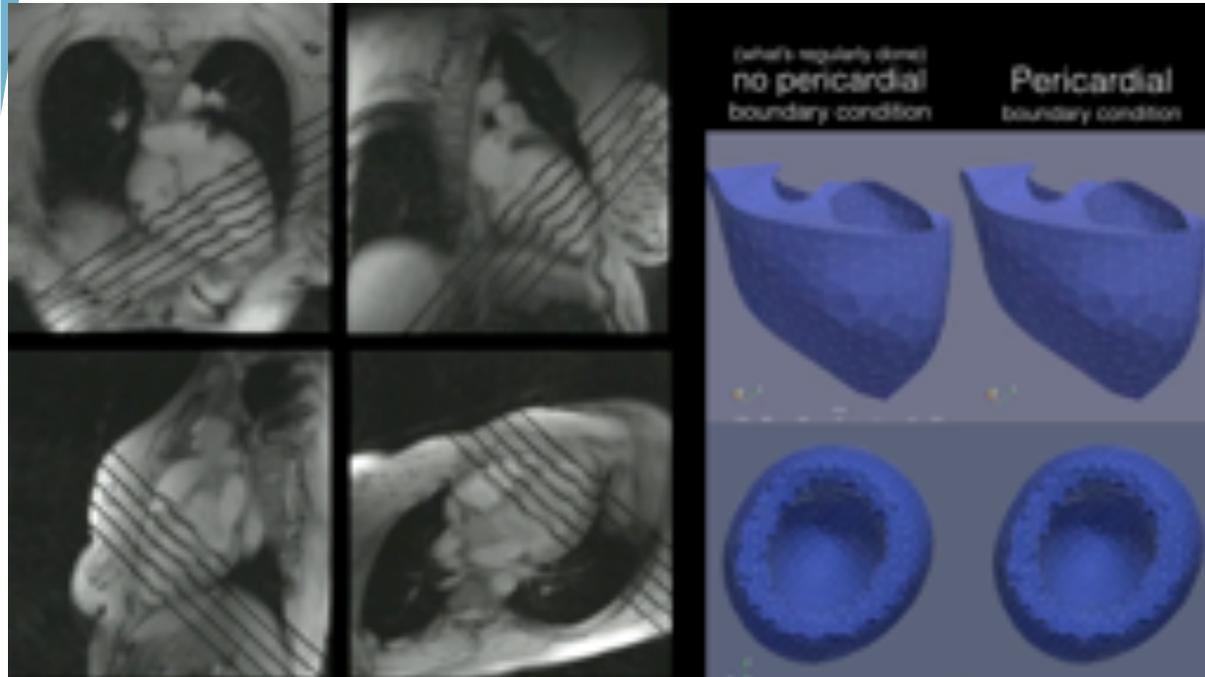


# Fully Coupled Electro-Mechanic-Fluid simulation

Number of elements: 4M total  
240 cores, 12 hrs, 400 ms



# Boundary Conditions and Physiological motion



# Human Biventricular Geometry Reconstruction

High Resolution MRI of Male and Female Human Hearts

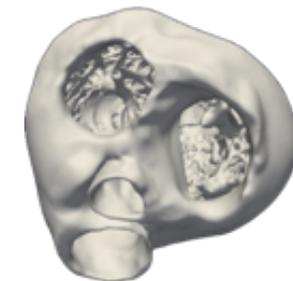
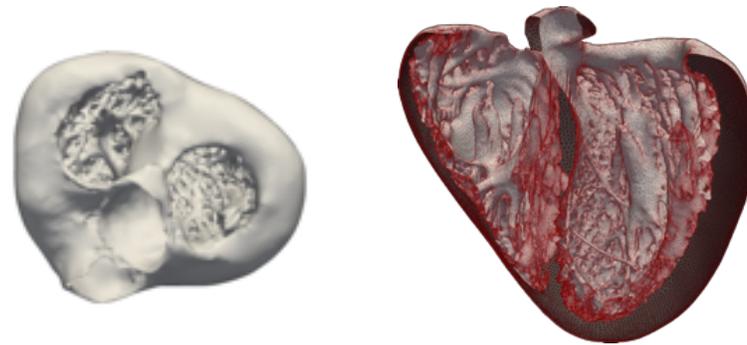
Courtesy of The Visible Heart © Lab



● Male Heart  
● Female Heart

Segmentation and Surface representation

Endocardial structures included are  $\geq 1 \text{ mm}^2$  cross-section

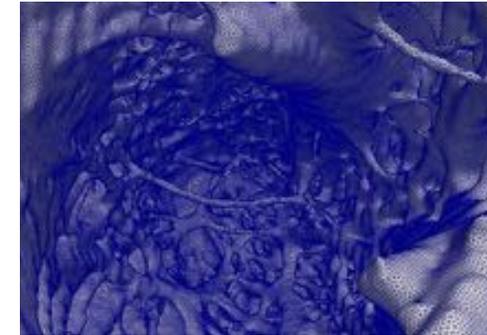


ReMESH

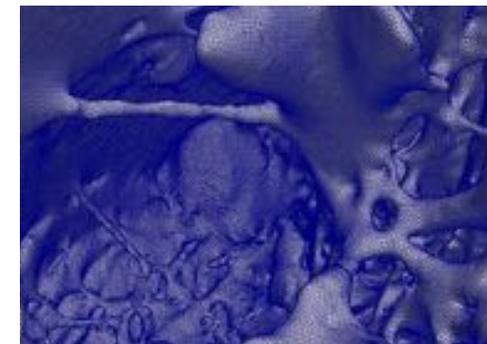


Biventricular Detailed Octree Volumetric Meshes

MAXIMUM ELEMENT SIDE LENGTH: 0.4 mm



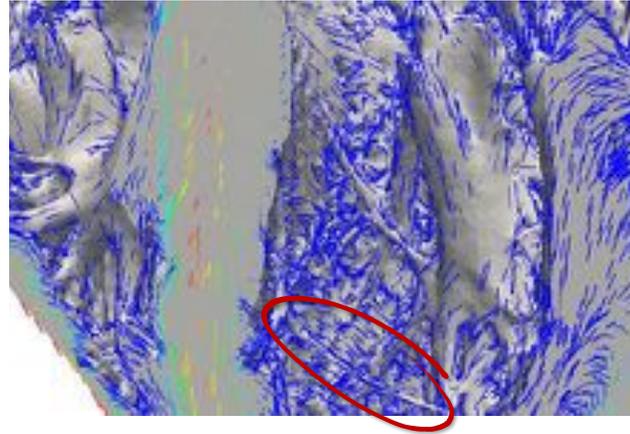
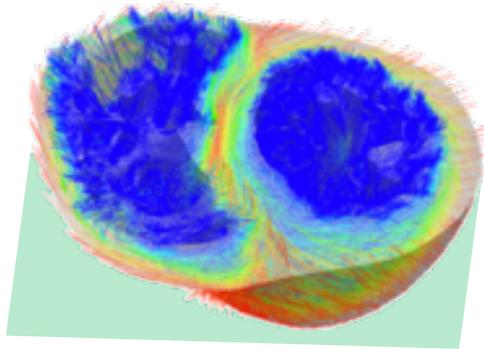
n° elements: 86.318.429  
Volume: 394.2 cm<sup>3</sup>  
n° points: 14.994.563



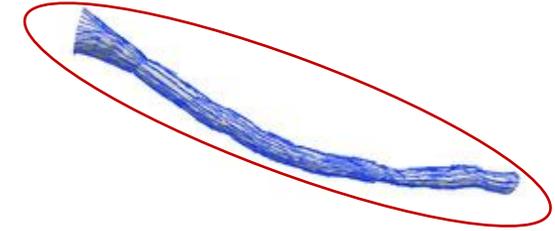
n° elements: 65.501.799  
Volume: 299.2 cm<sup>3</sup>  
n° points: 11.416.445

# Fiber orientations

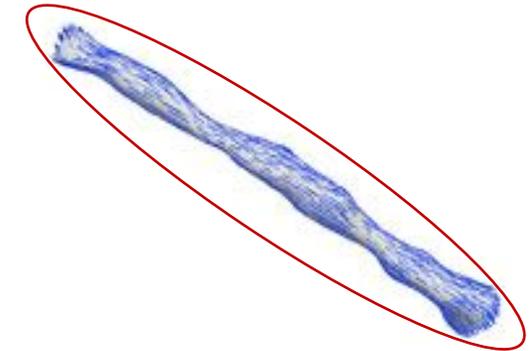
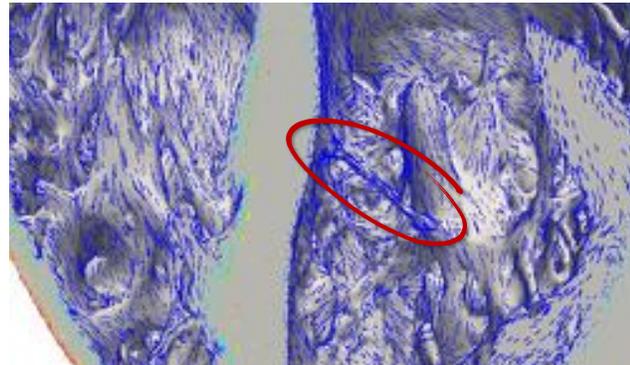
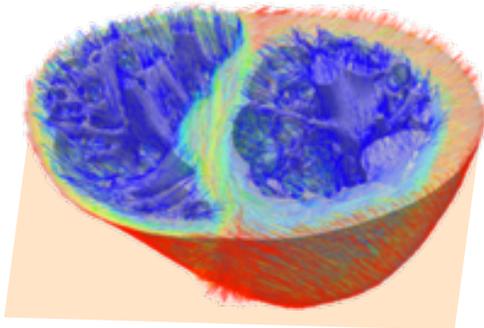
● Male Heart



False Tendons



● Female Heart

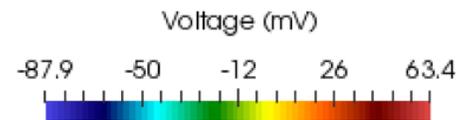
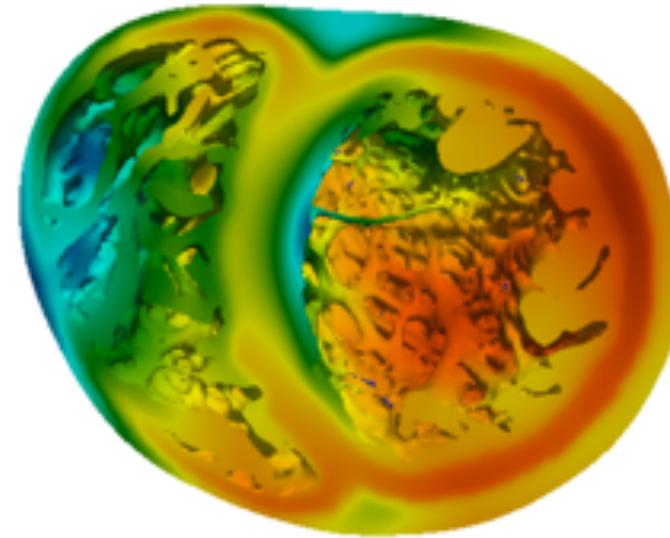
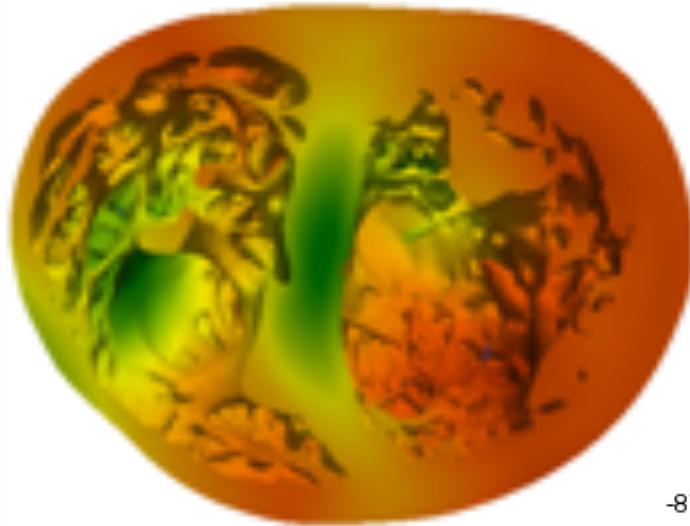
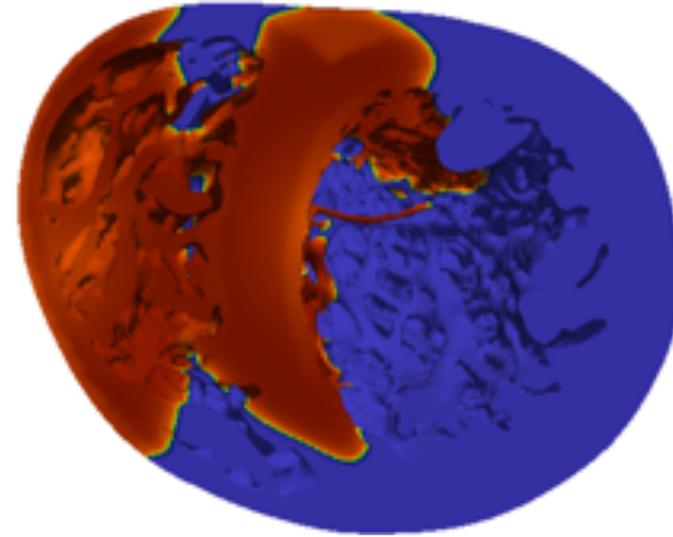
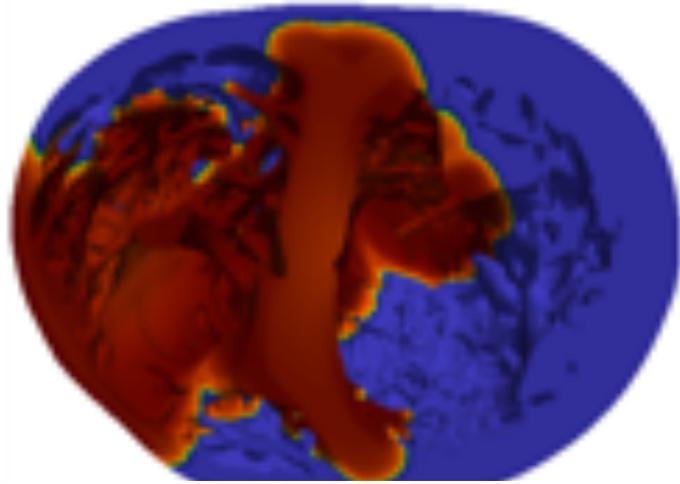


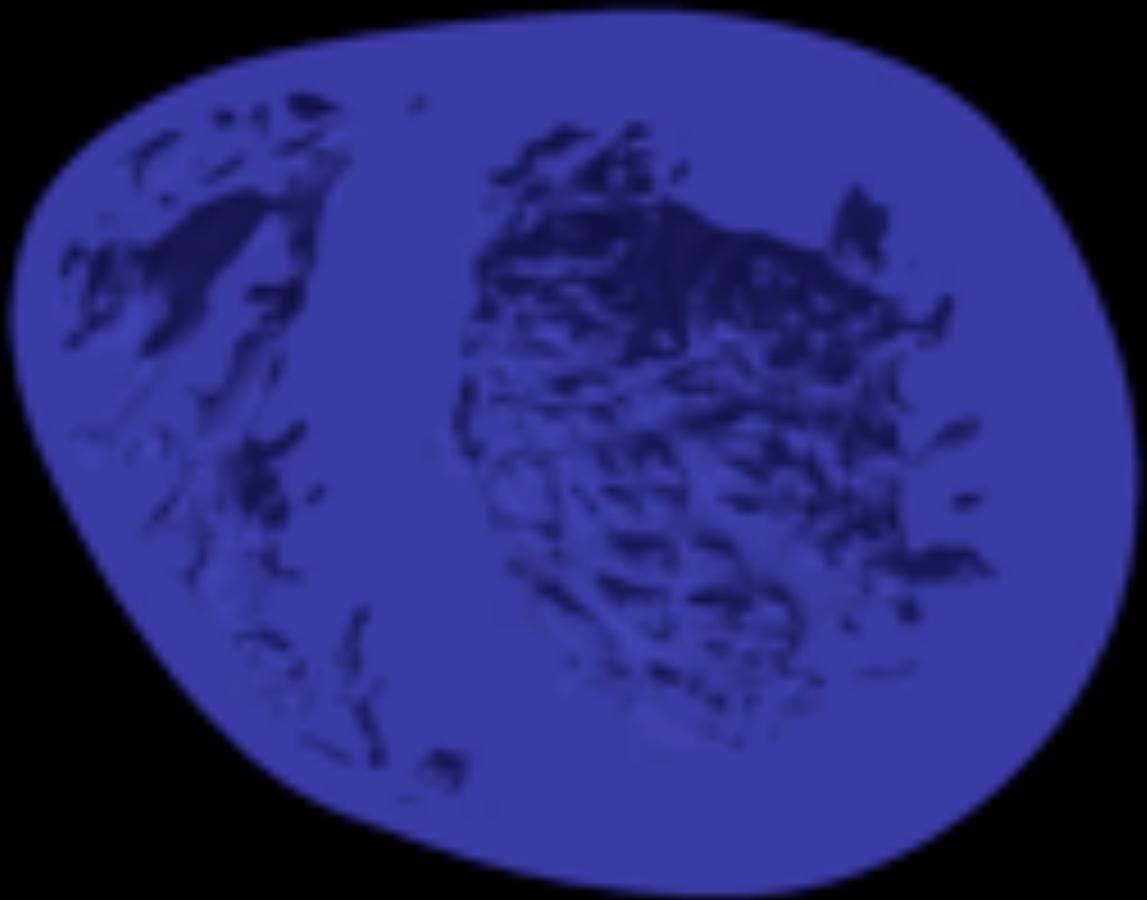
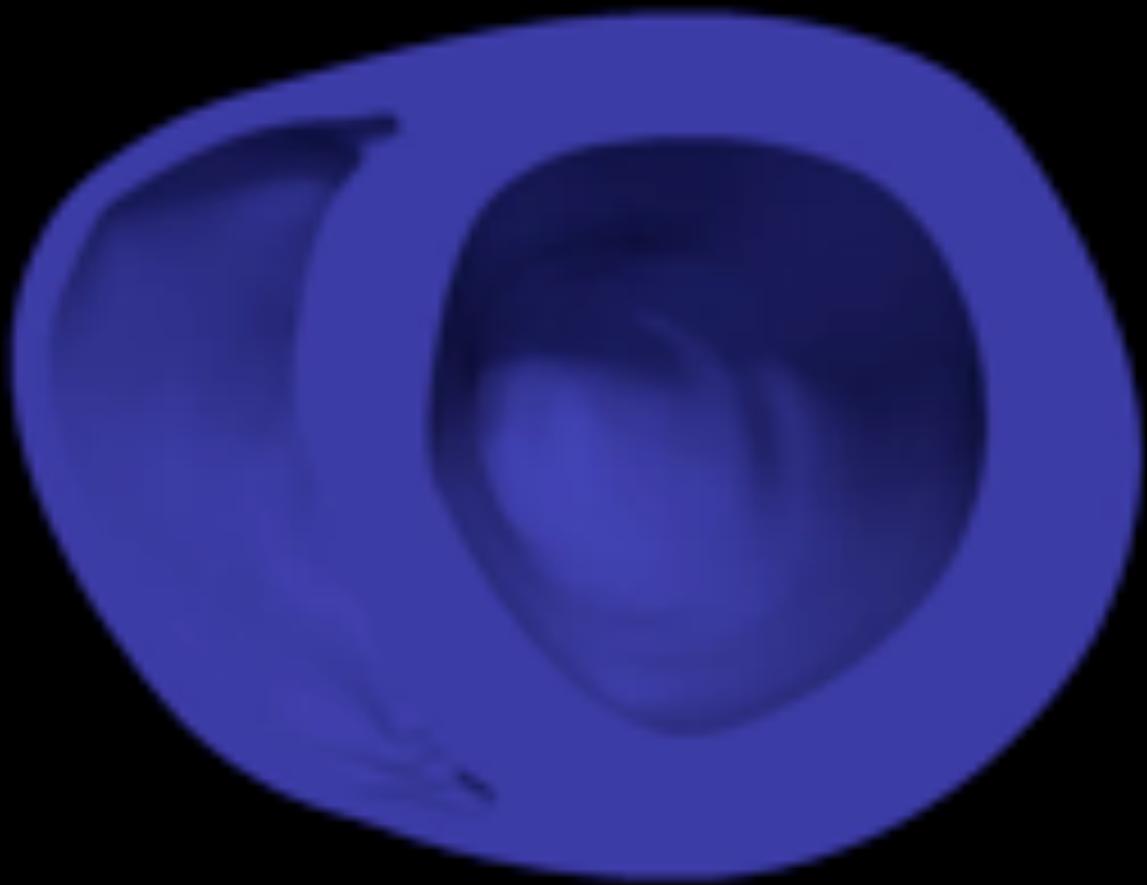
*A Rule-Based Method to Model Myocardial Fiber Orientation for Simulating Ventricular Outflow Tract Arrhythmias. Doste et al. FIMH 2017*



Universitat  
Pompeu Fabra  
Barcelona



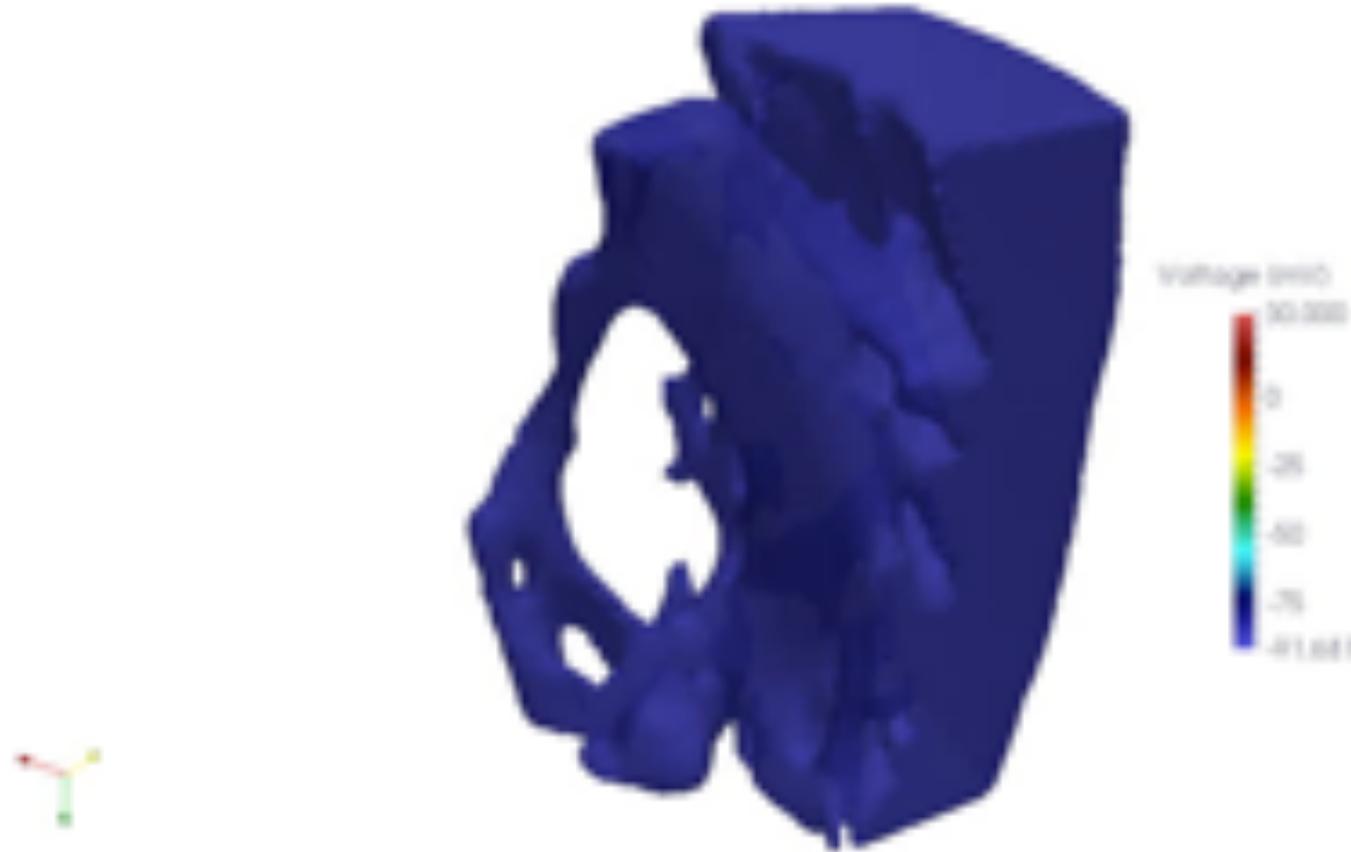




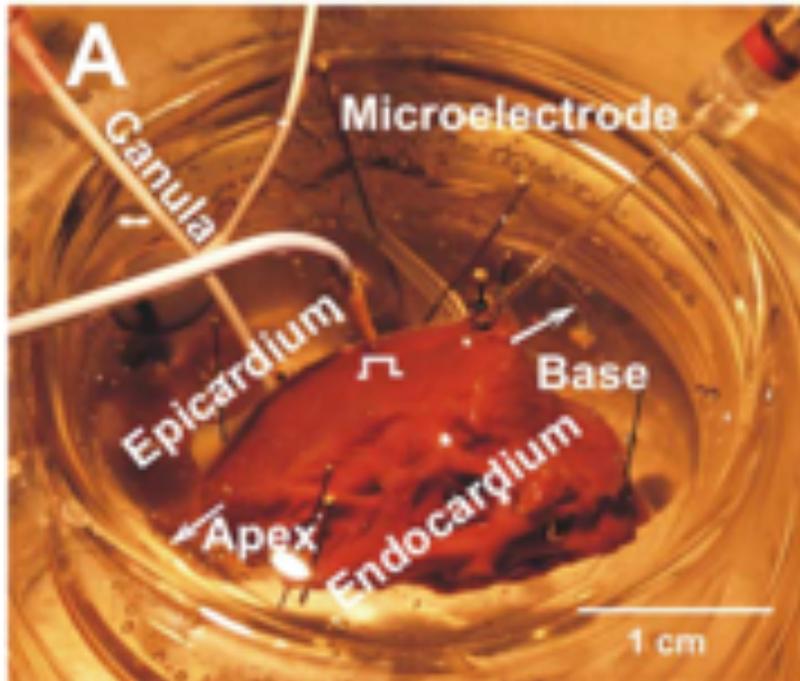
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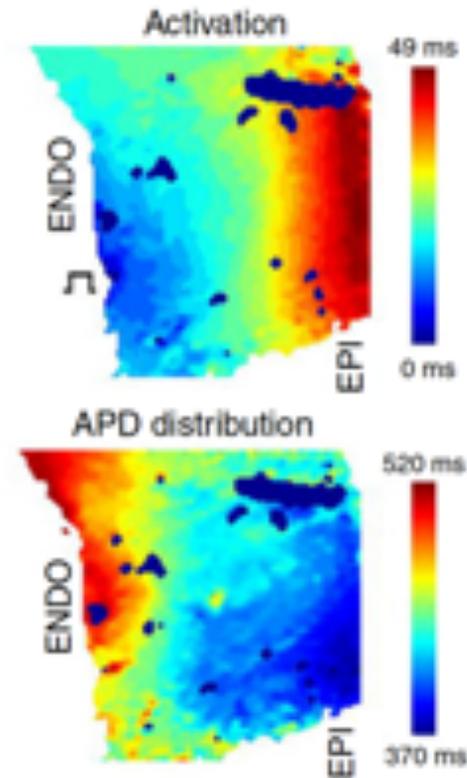
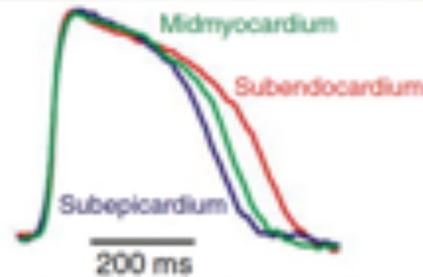
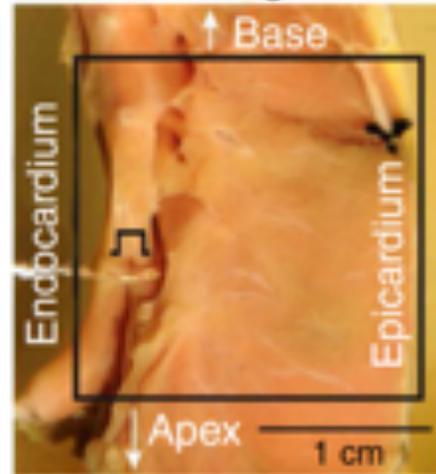
# Impact of endocardial structures on electro-mechanics



# Human Ventricular Wedge Preparation



A Non-Failing heart



# Human Cardiac Wedge Preparation

Number of elements: 13,807,755

Nodal points: 2,480,801

48 cores, 2 hrs

260 ms

