



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

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University of Kragujevac, Serbia

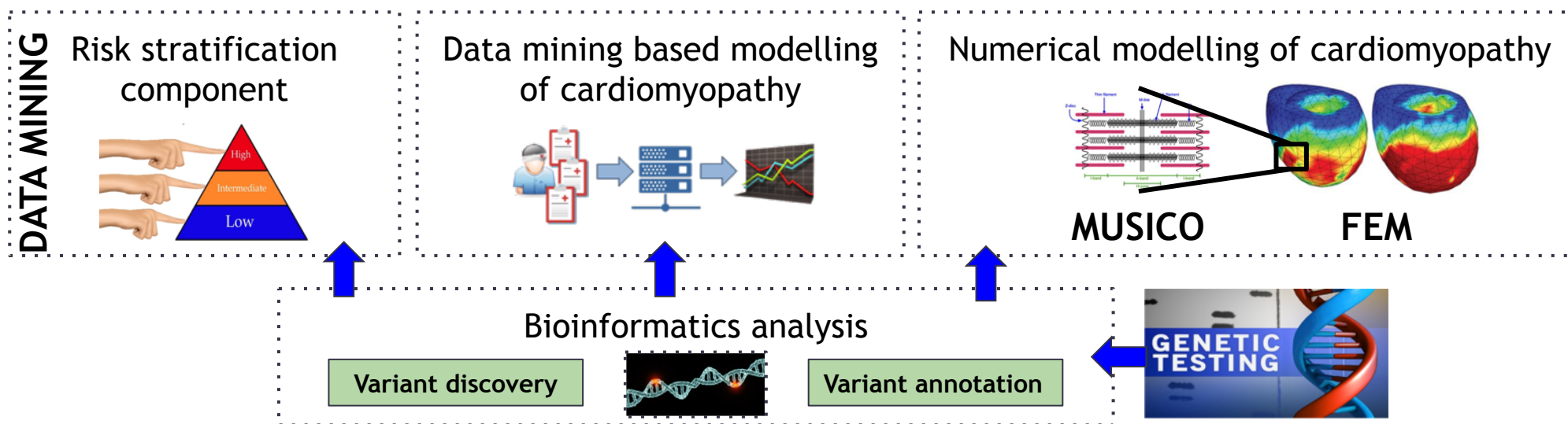
Institute of Engineering and Technology, London: Savoy Place
September 25-27, 2019

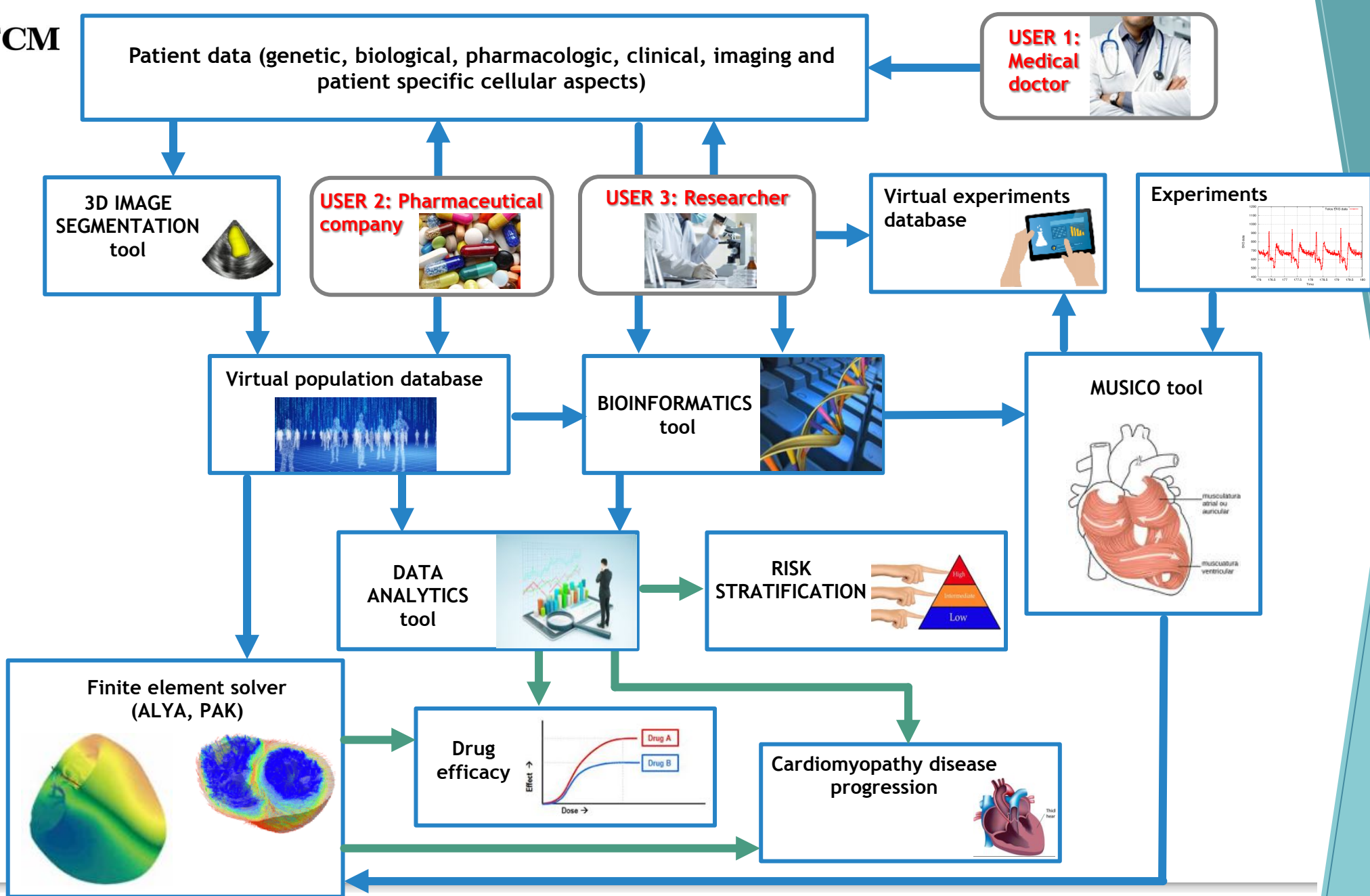


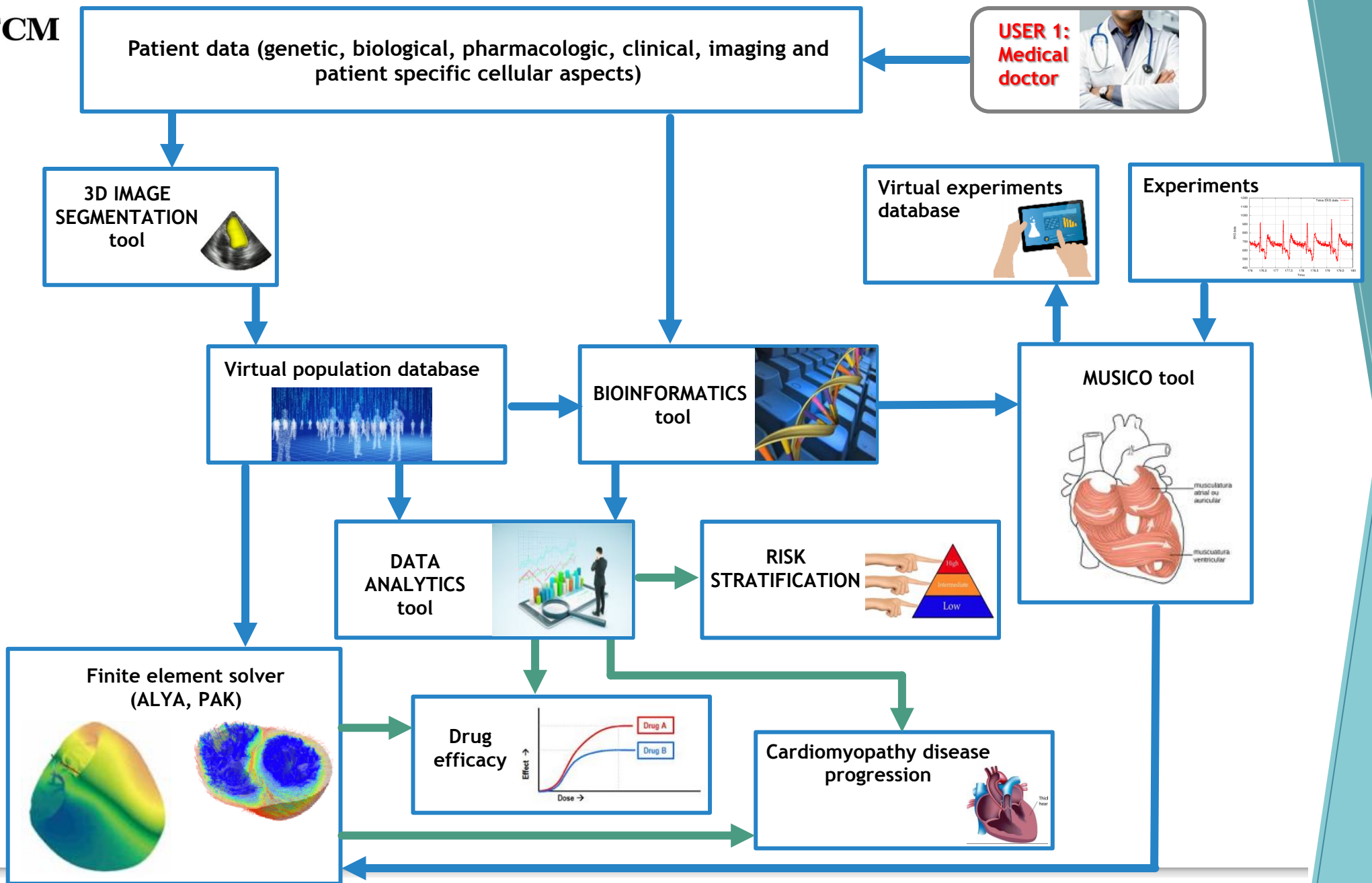
This project has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 777204

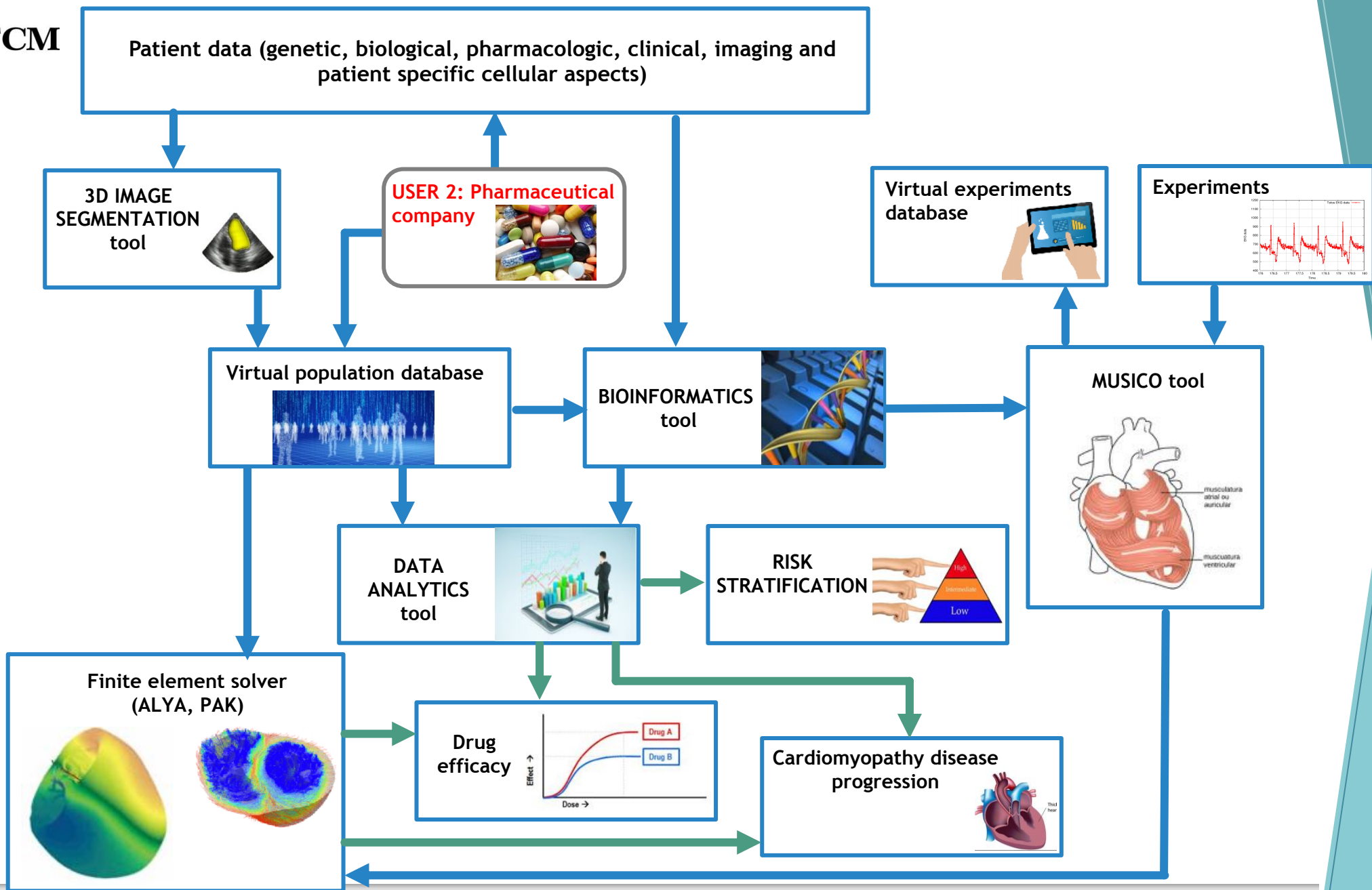
SILICOFCM Scope

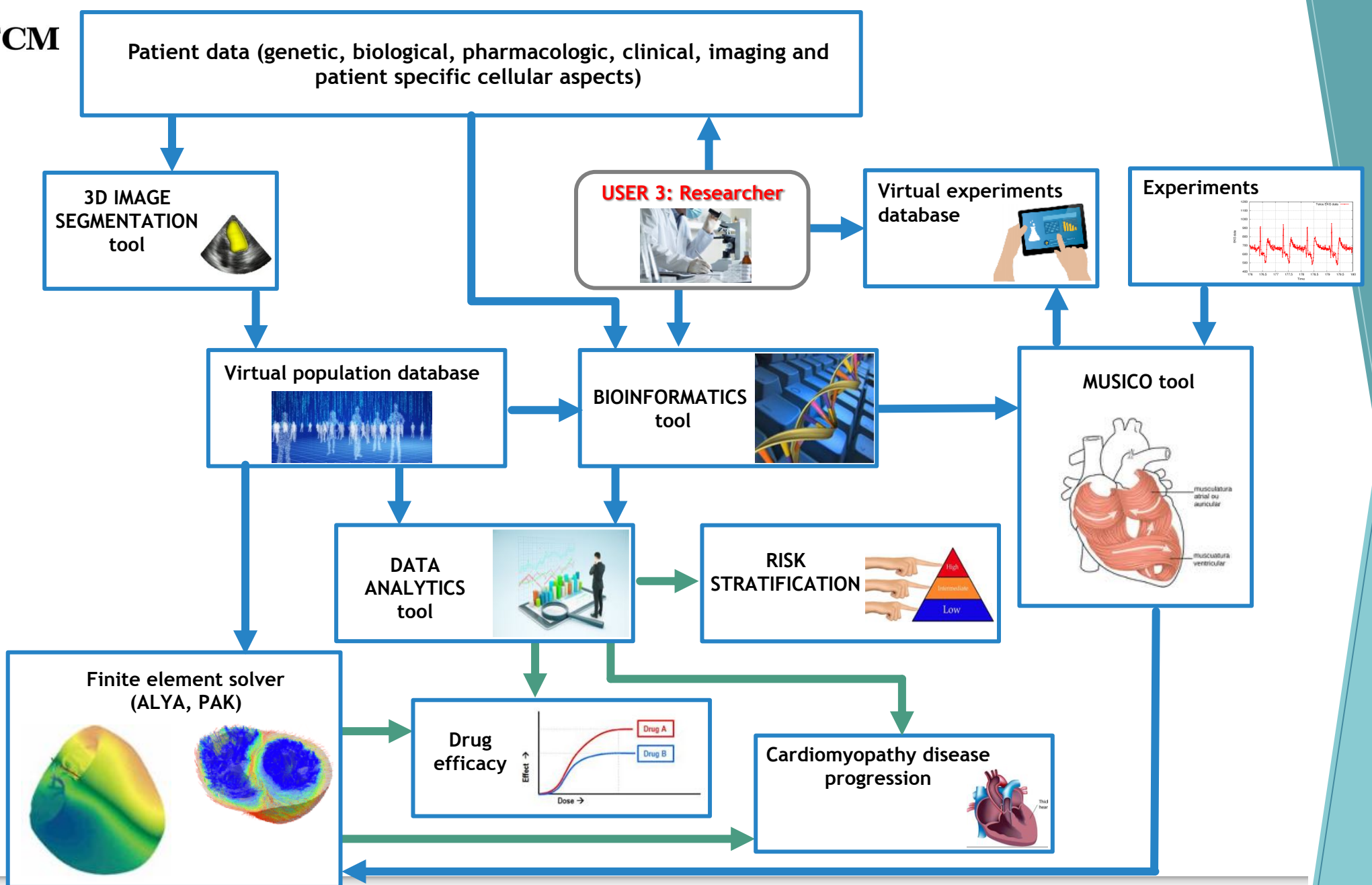
- ▶ SILICOFCM aims to develop a computational platform for *in silico* clinical trials of Familial cardiomyopathies (FCMs) that would take into consideration comprehensive list of patient specific features (genetic, biological, pharmacologic, clinical, imaging and patient specific cellular aspects) capable 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.



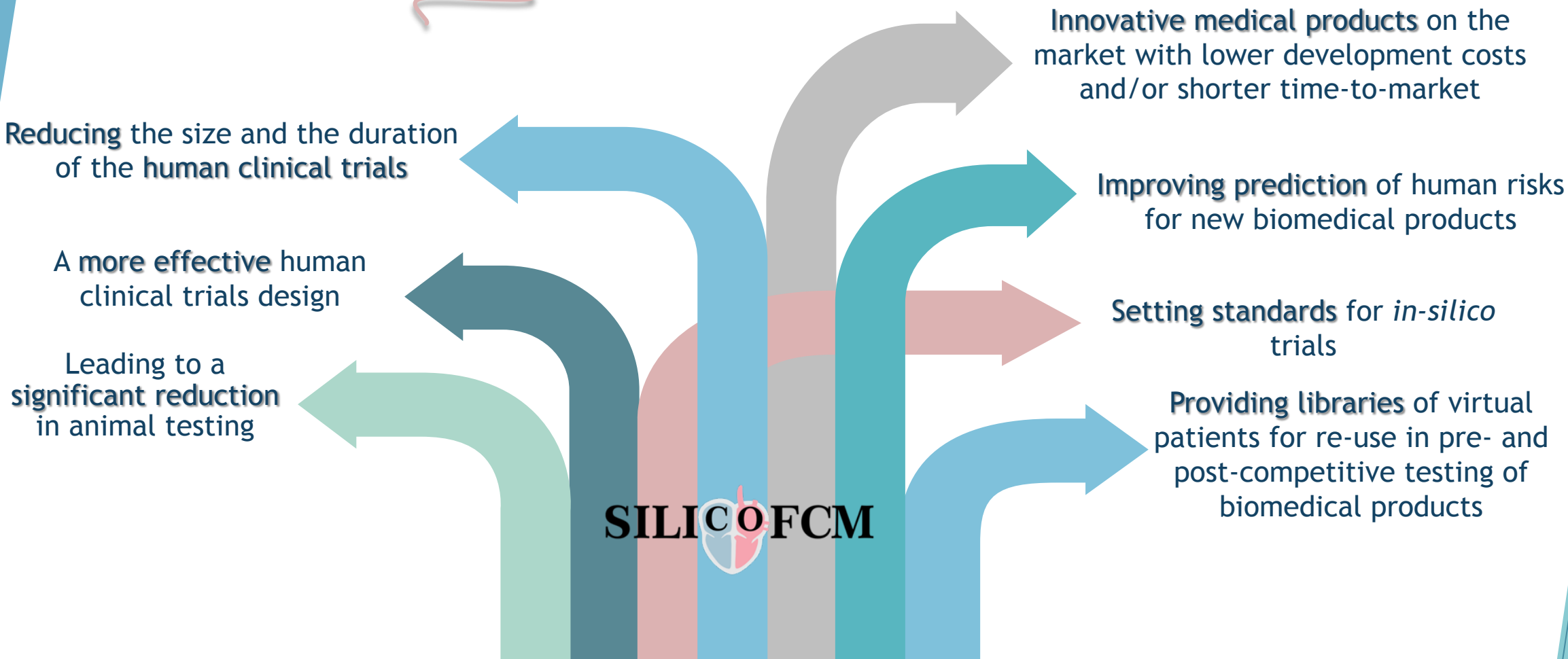
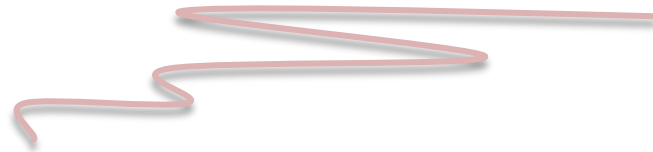




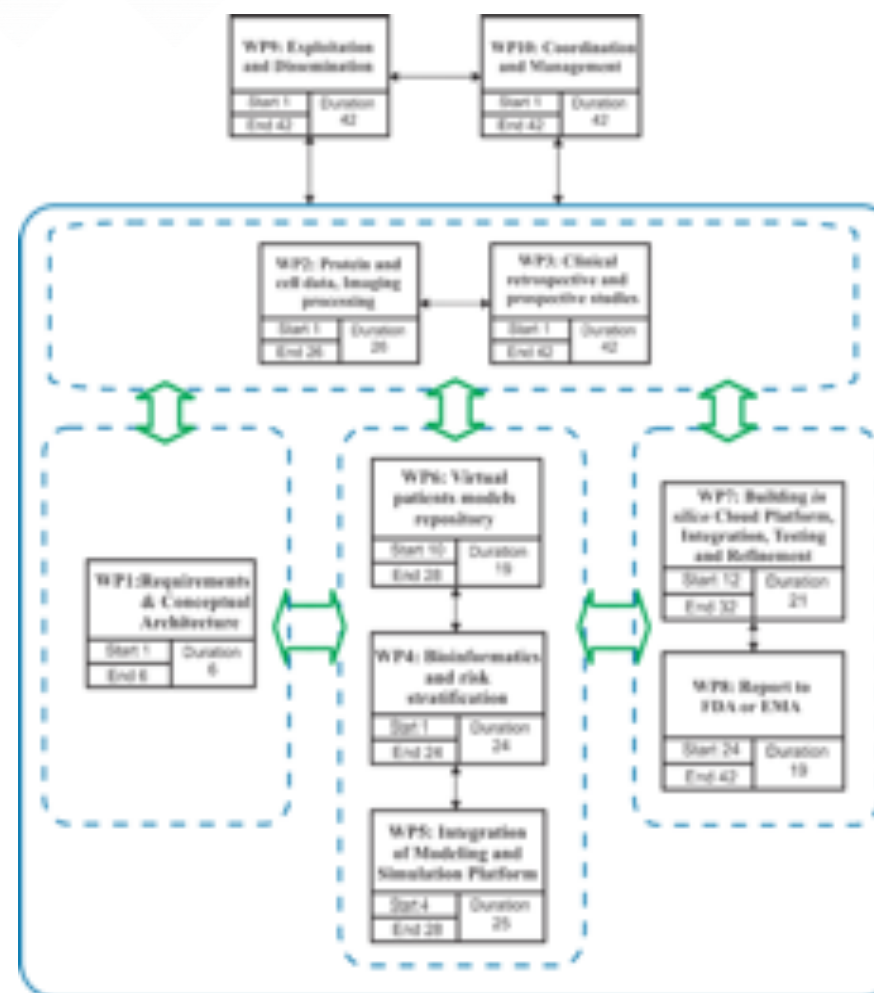
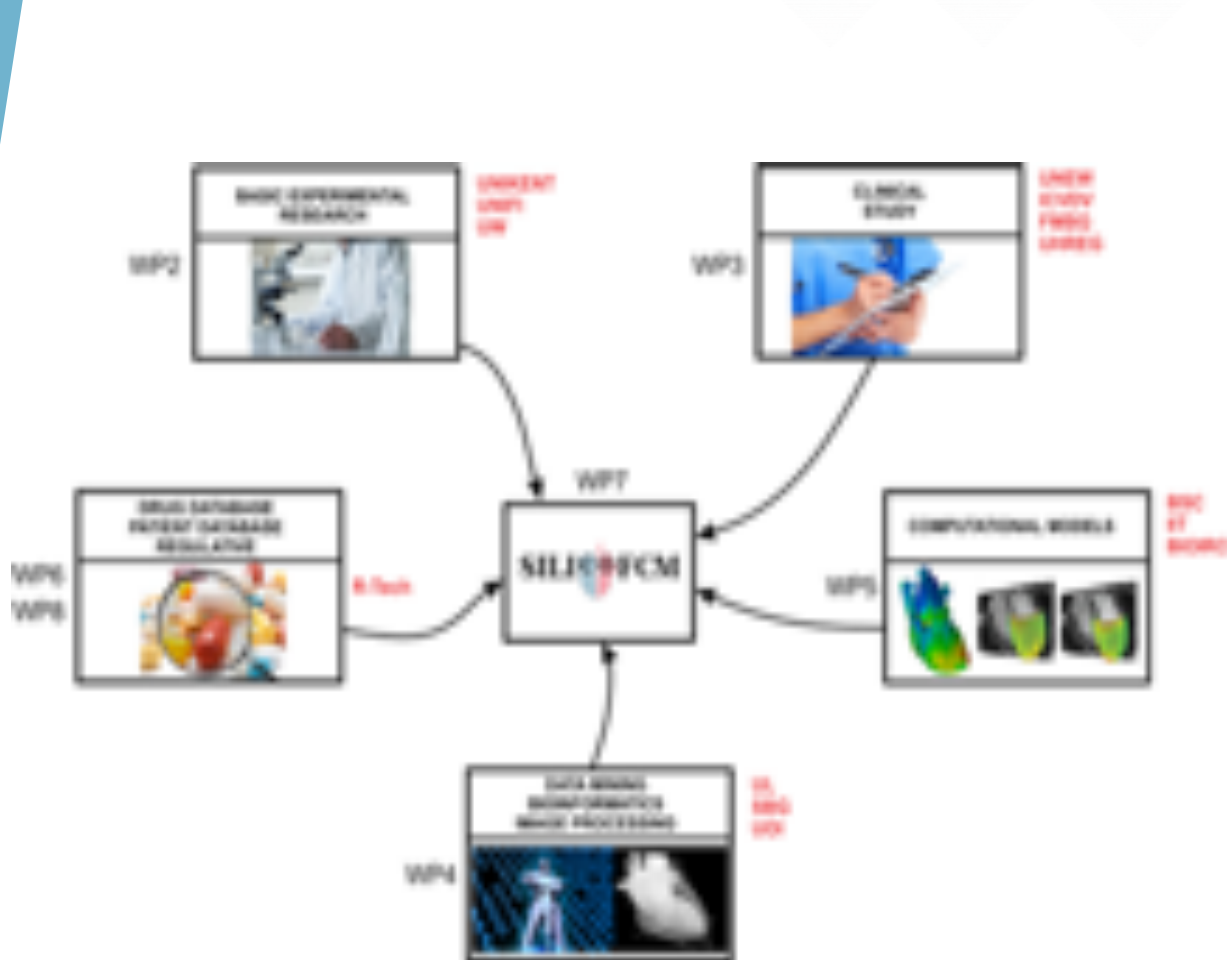




SILICOFCM Impact



WPs Interaction



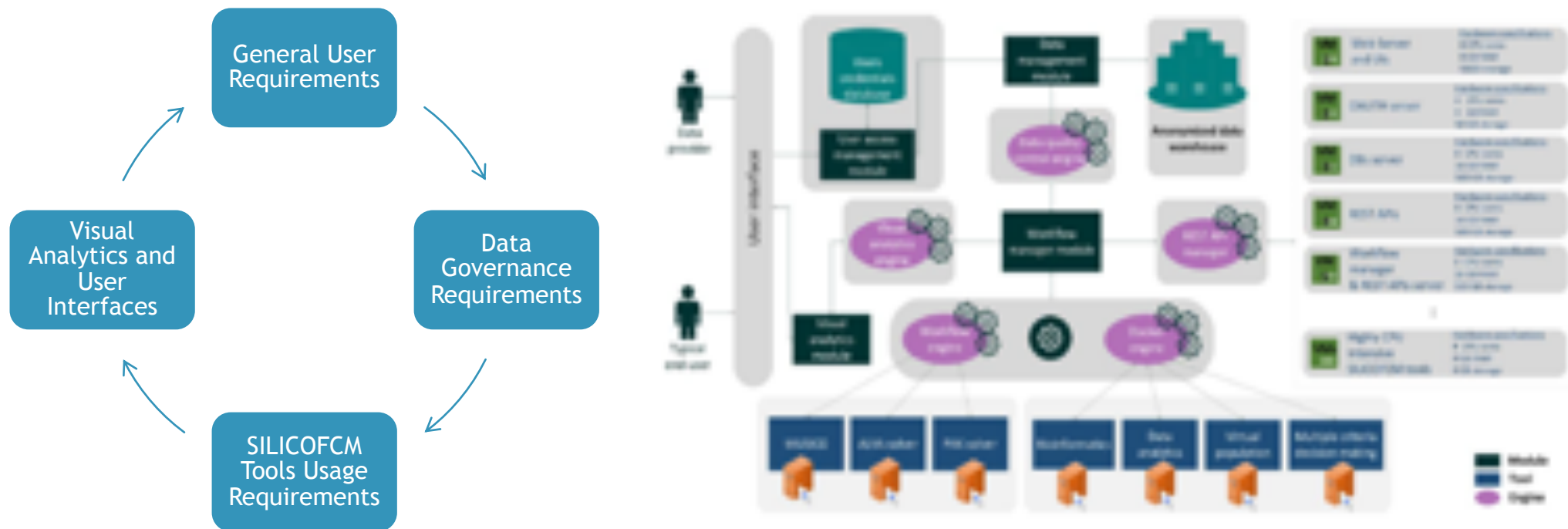
WP1) Requirements & Conceptual Architecture

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

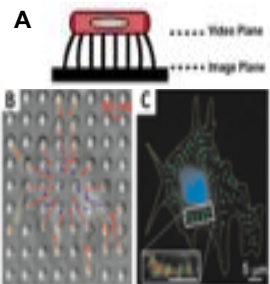
Main objective

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

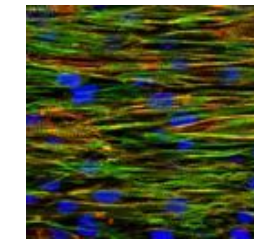
3D engineered
Tissue-level
mechanics
heart tissue (EHT)



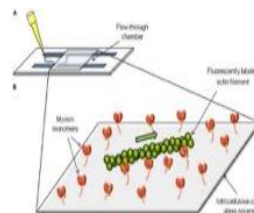
Cell
Contraction/Relaxation
(micropost arrays)



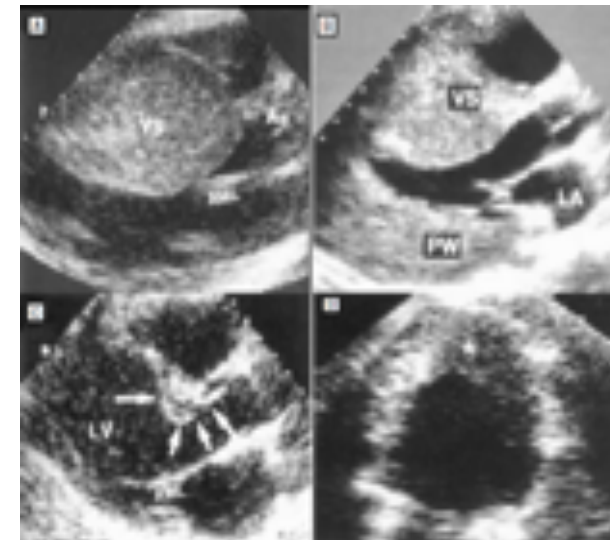
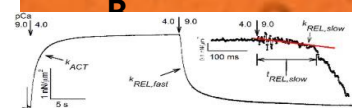
Cell contractility/ Ca^{2+} transients



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



Isolated Myofibrils
contraction/relaxation kinetics



WP3

Clinical retrospective and prospective studies

Leader UNEW

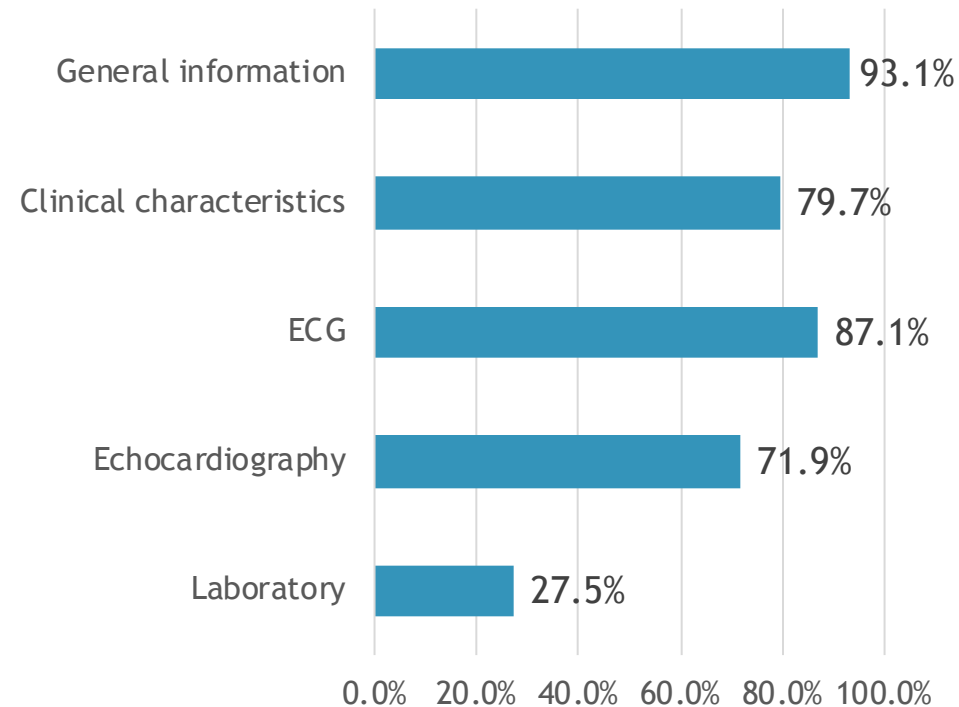
M1-M42

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

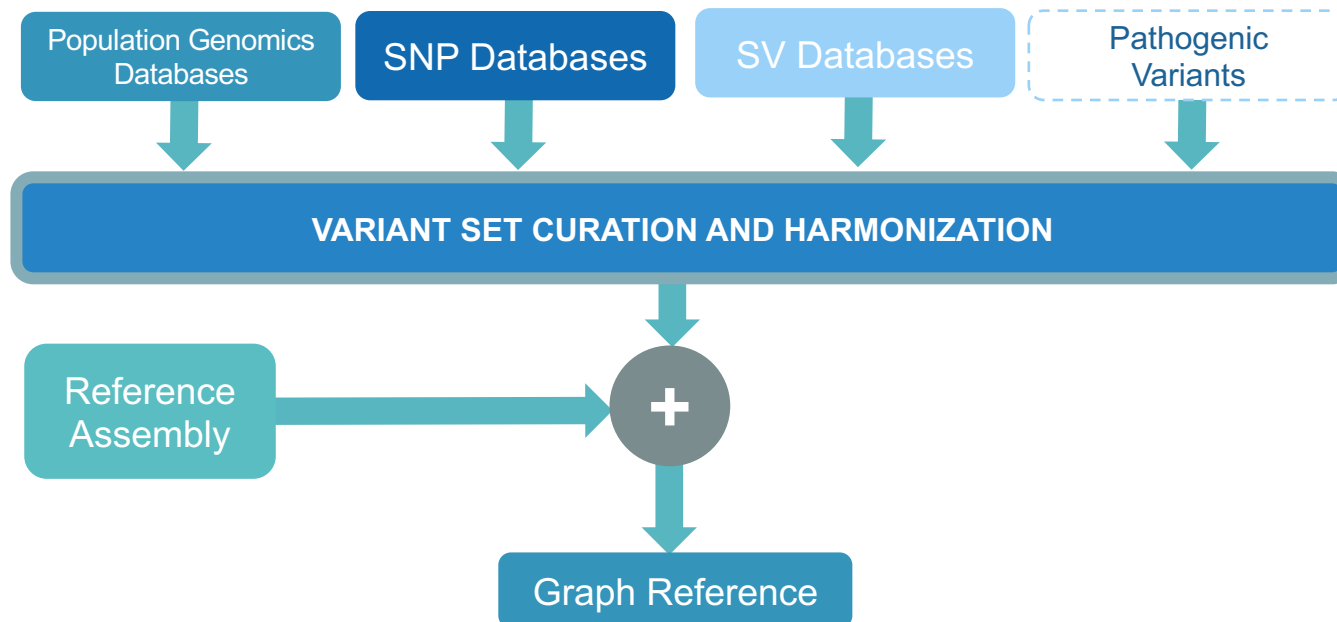
Bioinformatics and risk stratification

Leader SBG

M1-M24

Main objective

- ▶ To put in place the genomic component in the SILICOFM platform and integrate it with the SILICOFM 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

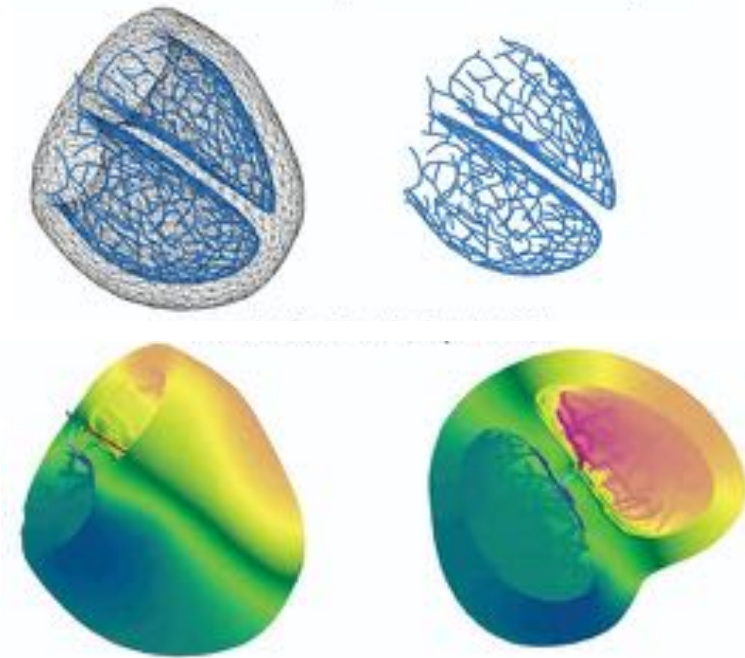
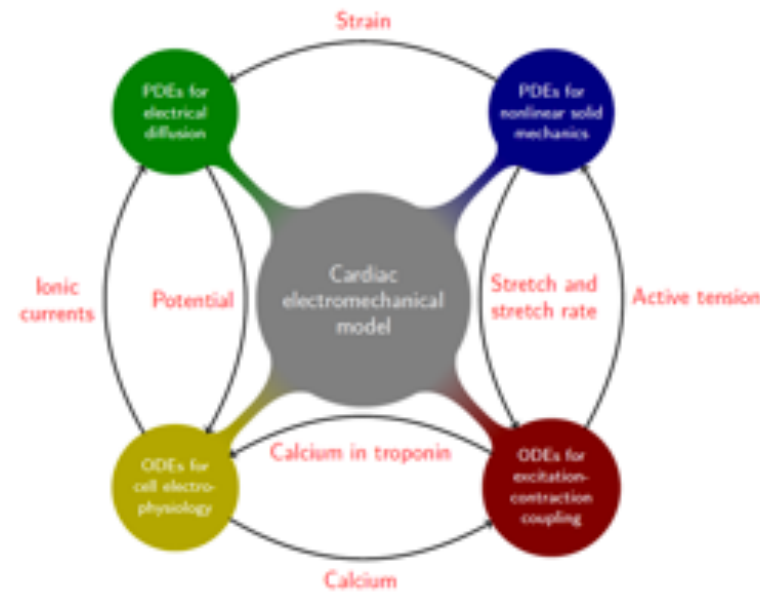
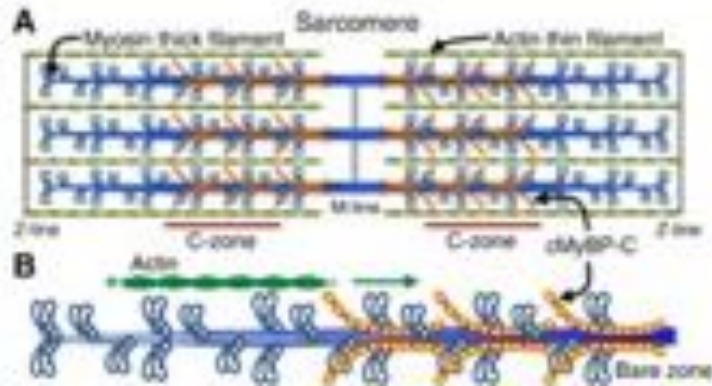
WP5 Integration of Modelling and Simulation Platform

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

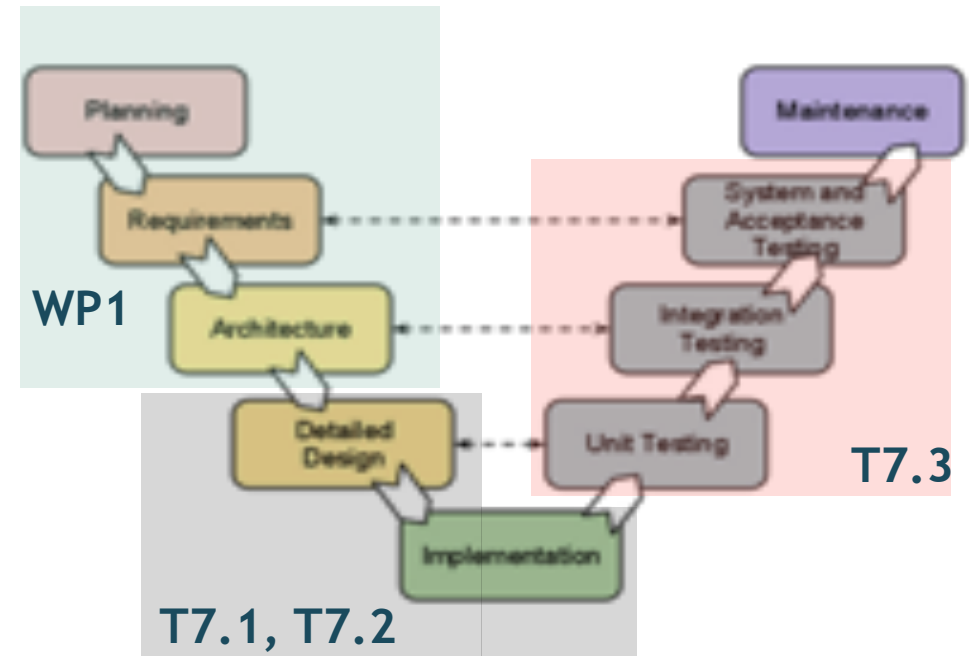
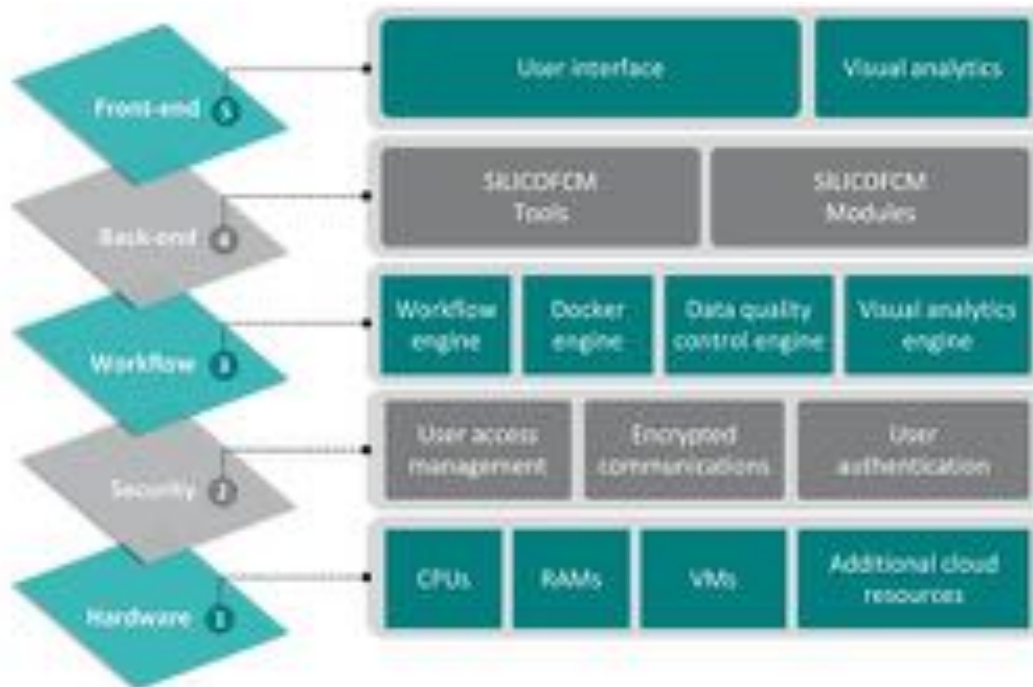
Building *in Silico* Cloud Platform, Integration, Testing and Refinement

Leader UOI

M12-M32

Main objective

- To integrate the SILICOFM 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&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

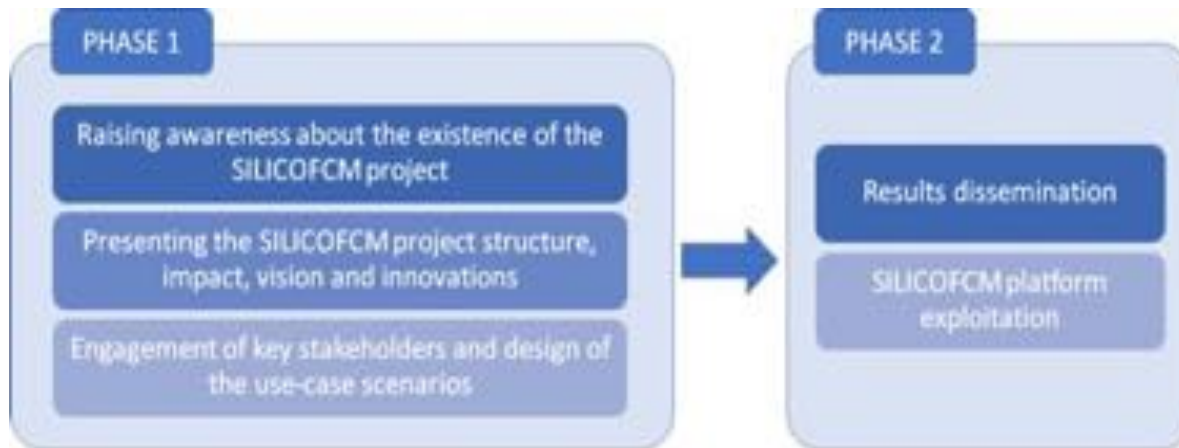
Exploitation and Dissemination

Leader BioIRC

M1-M42

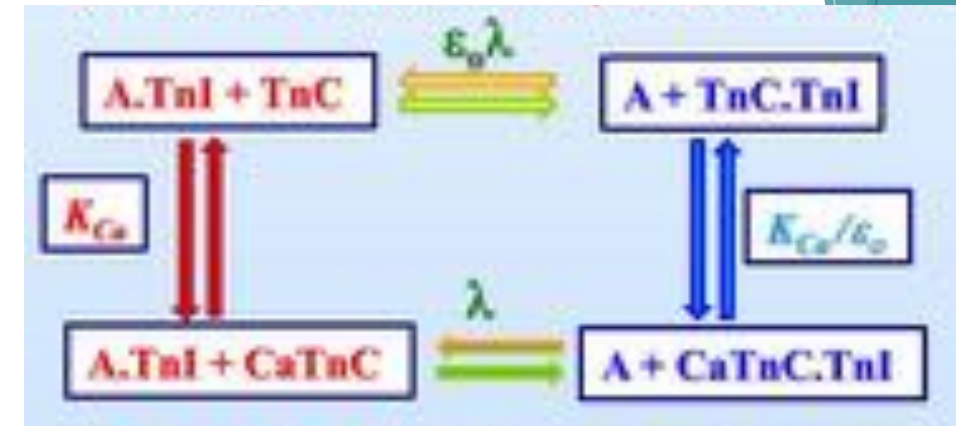
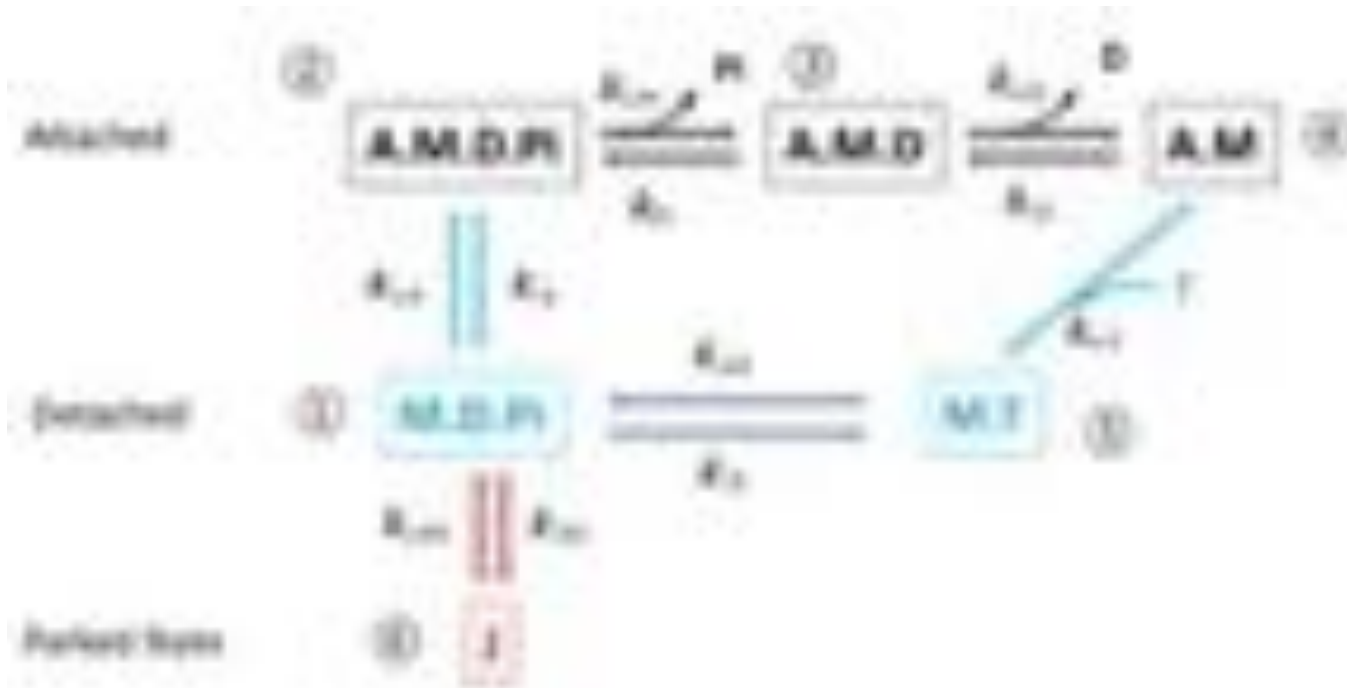
Main objective

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

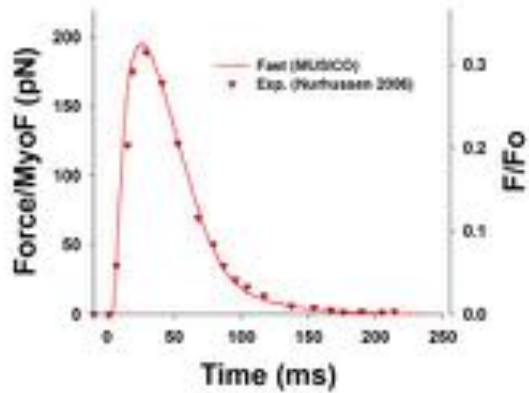


MUSICO: Six State Crossbridge Model & Calcium Kinetics with TnC

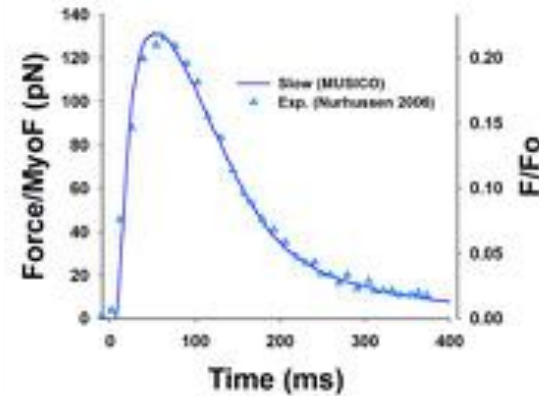
Cardiac Muscle



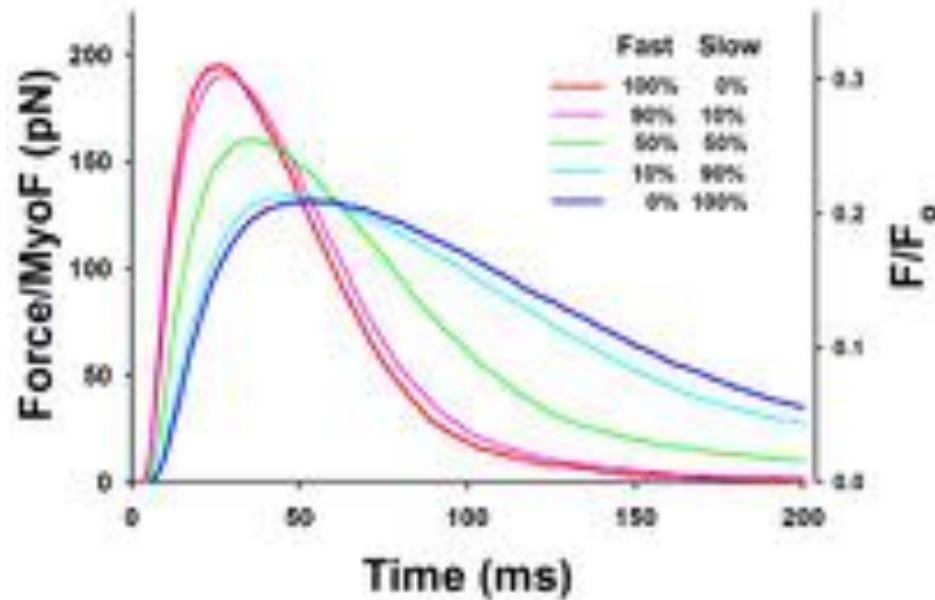
Fast Twitch Contraction



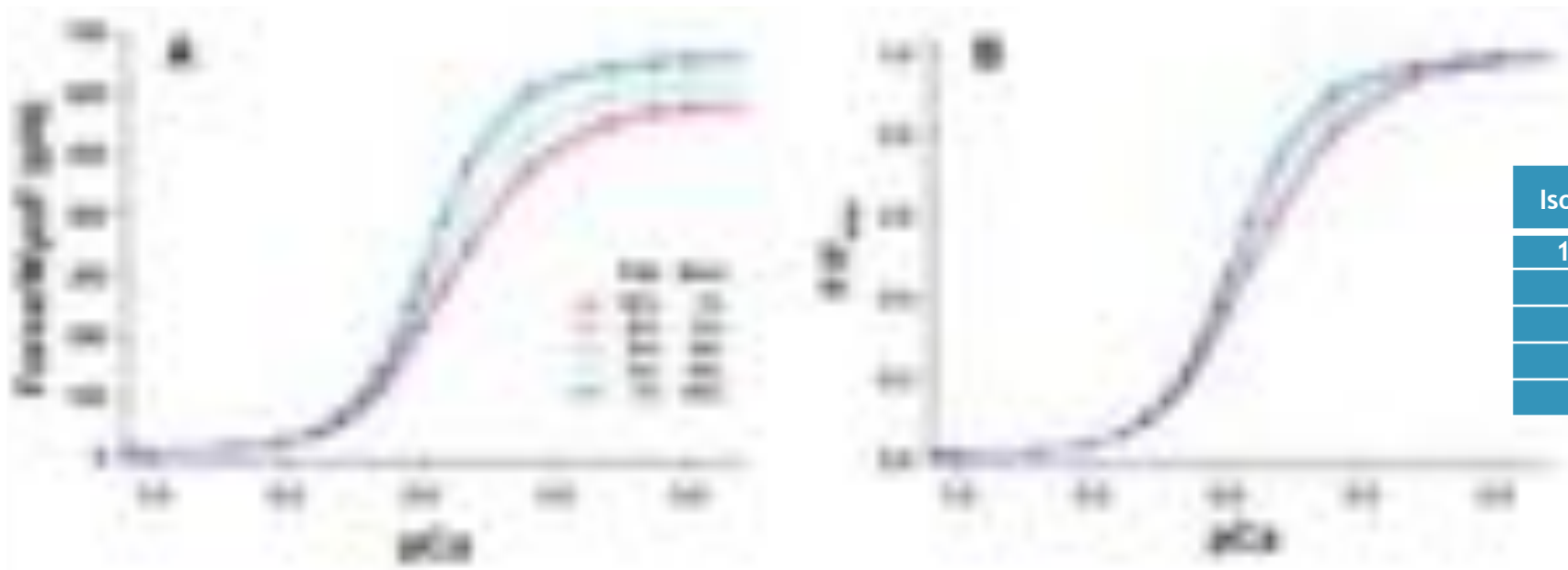
Slow Twitch Contraction



- ❖ The calcium transient:
- ❖ Peak of 1.6 μ M at 6 ms
- ❖ Decays to low calcium concentrations at about 35 ms.



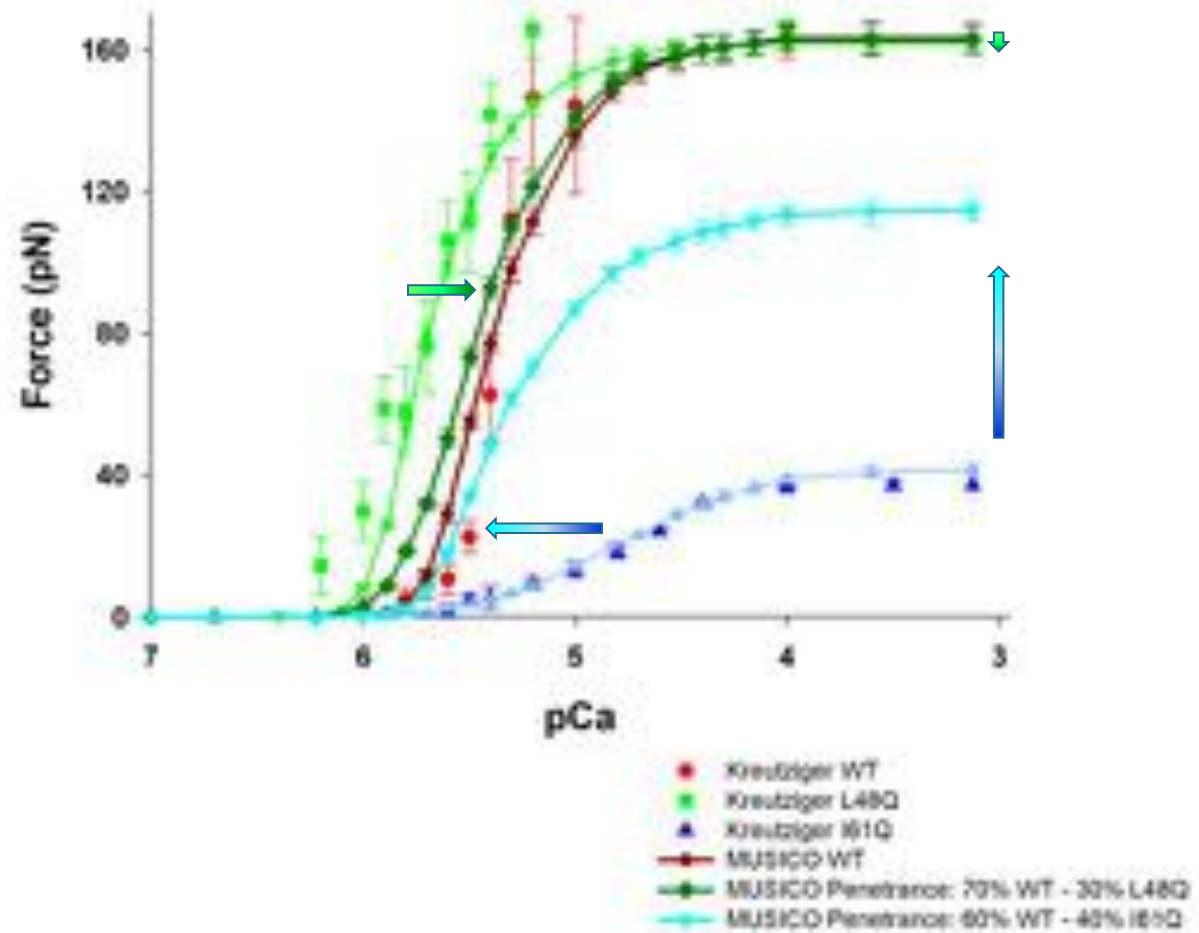
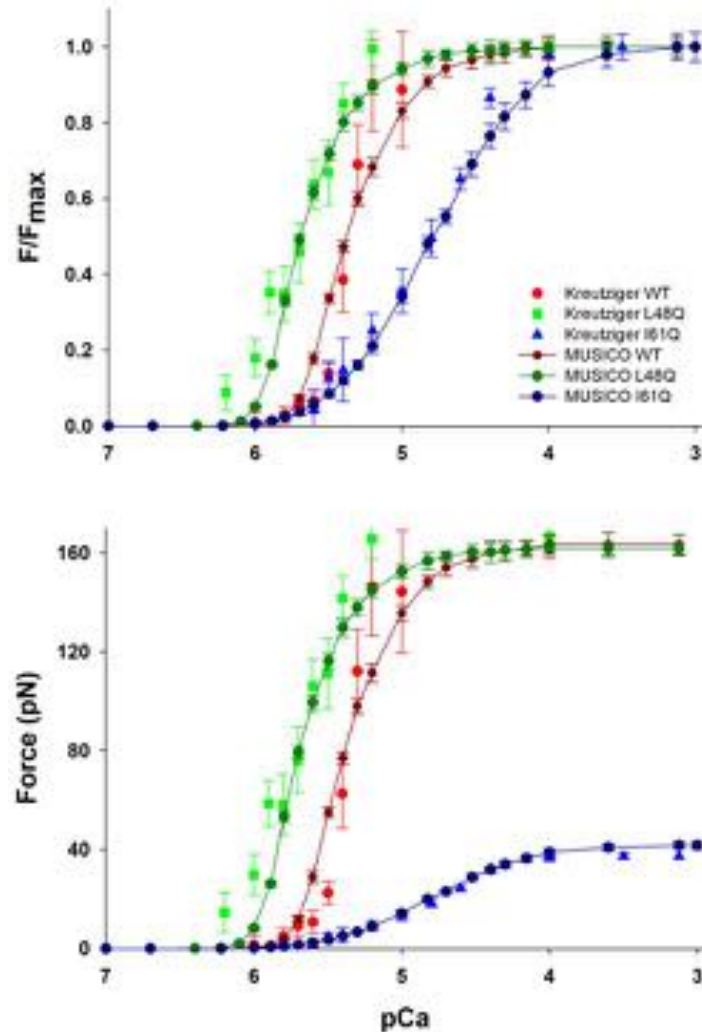
Effect on Hill Coefficient and Calcium Sensitivity



Isoform Mixture		Hill Coeff	pCa ₅₀
100% α	0% β	2.302	5.890
90% α	10% β	2.342	5.899
50% α	50% β	2.531	5.933
10% α	90% β	2.807	5.966
0% α	100% β	2.889	5.973

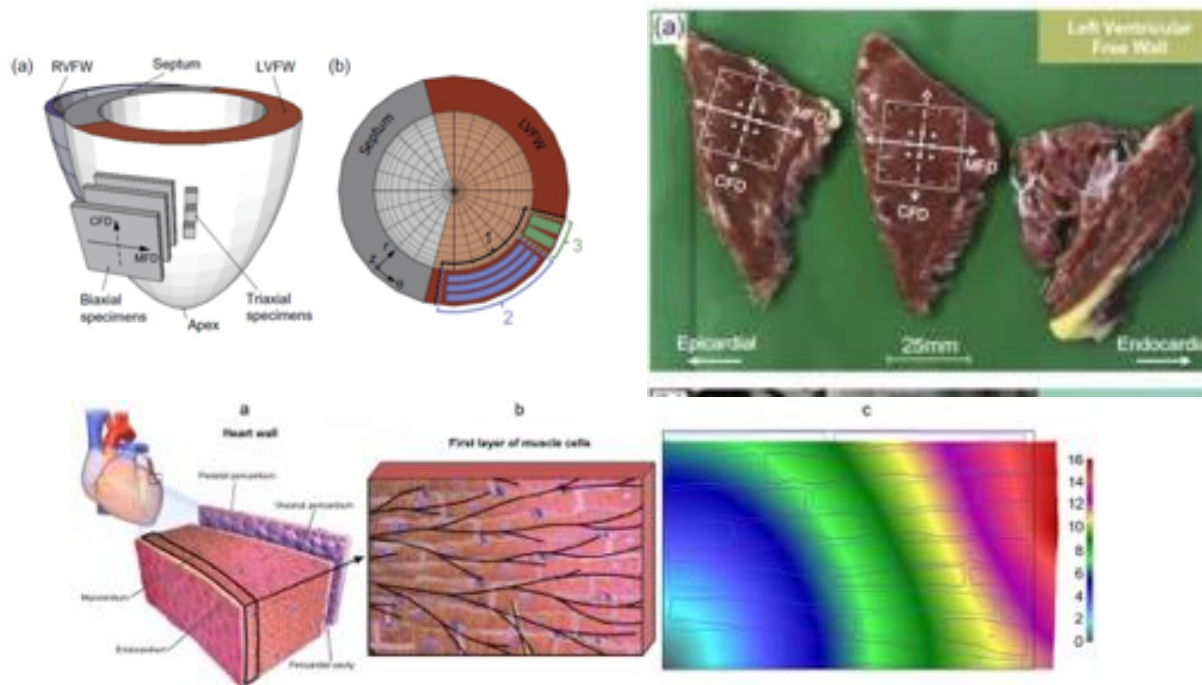
Fit to Kreutziger Experiments - Rat Trabeculae 100% Mutants

Kreutziger - Predicted Penetrance (Rat Trabeculae)

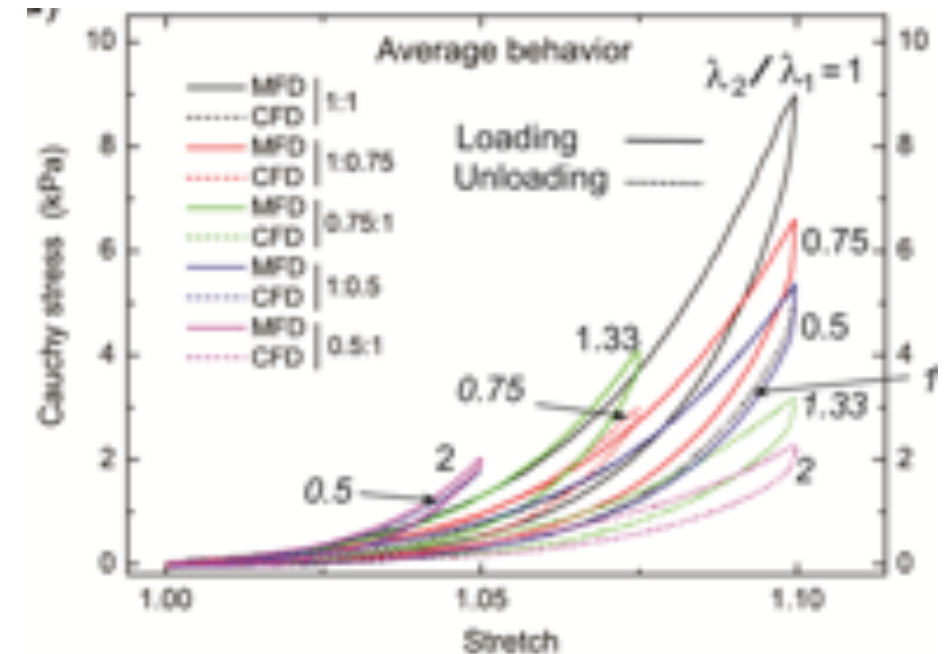


Upgrade FE biomechanical simulation PAK Solver

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



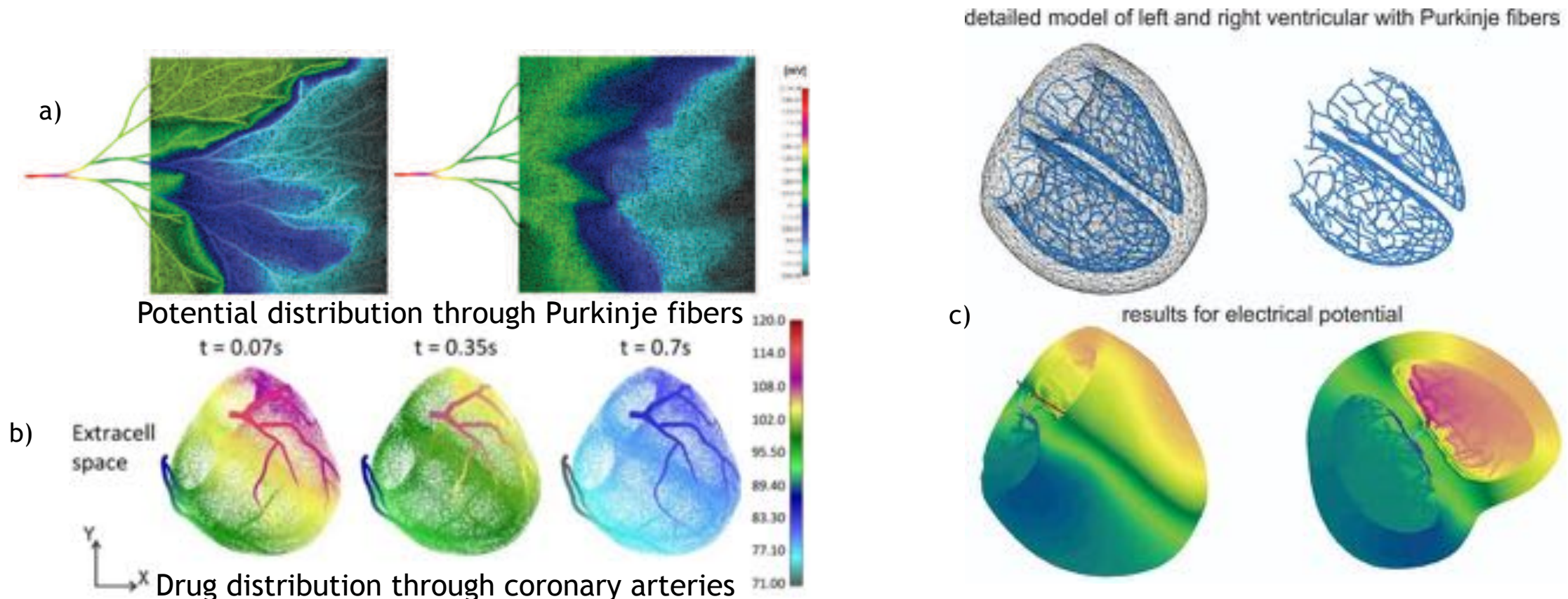
Field of displacements in heart tissue due to uniaxial straining



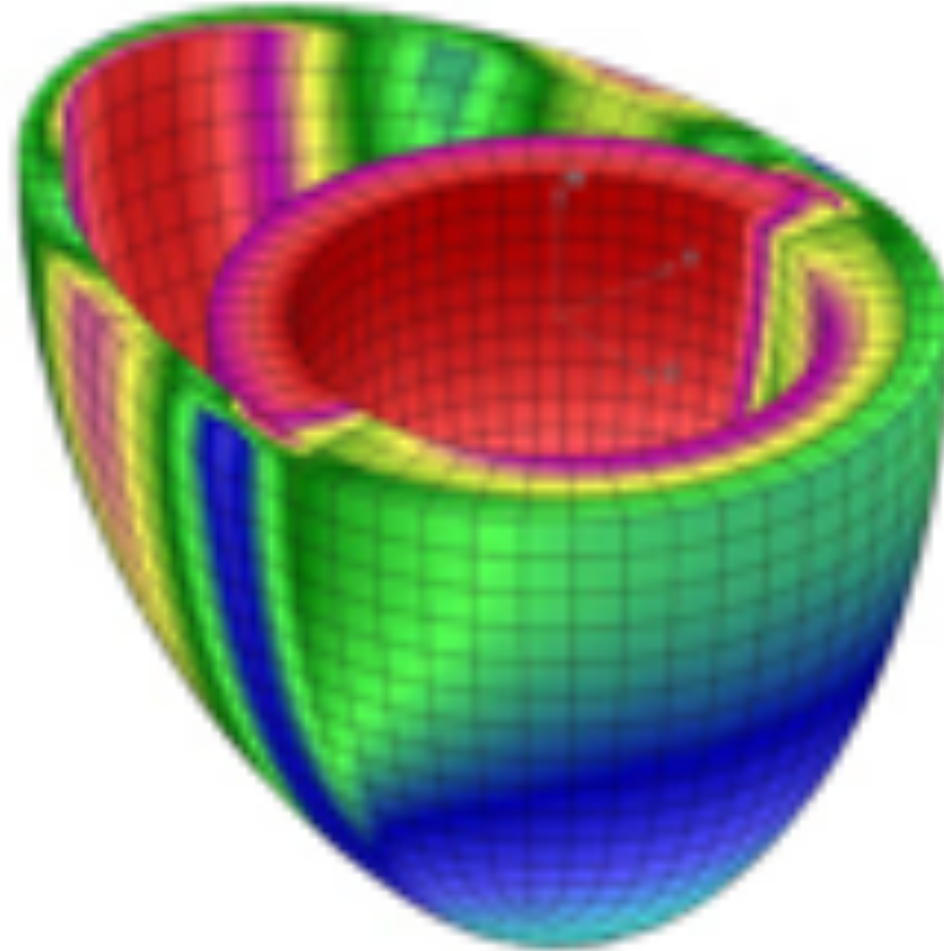
Experimental curves with hysteresis for biaxial loading of myocardium tissue

Upgrade FE biomechanical simulation PAK Solver

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

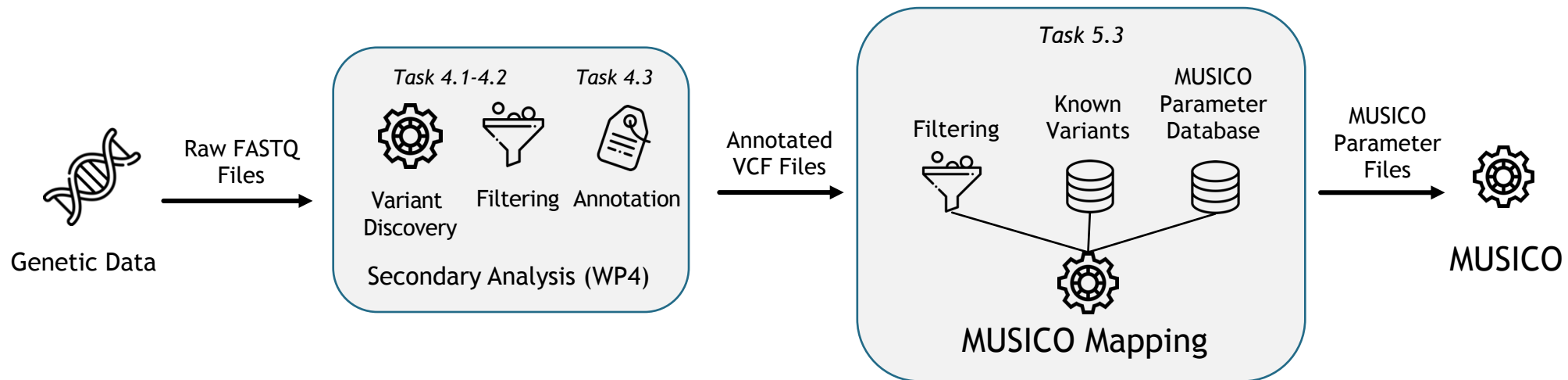


Fluid-solid interaction-PAK



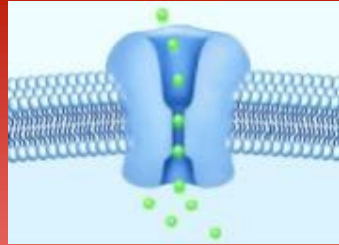
Linking bioinformatics and MUSICO

- ▶ 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.

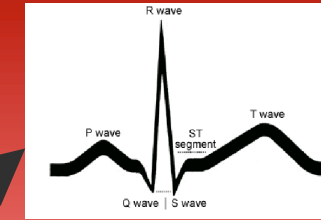


The Heart Physiology as an Electro-Mechanic System

Electrophysiology

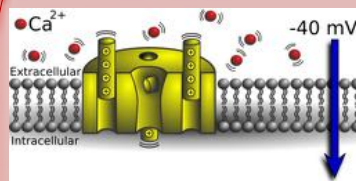


ion channels



ventricles

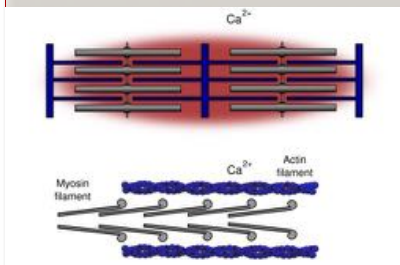
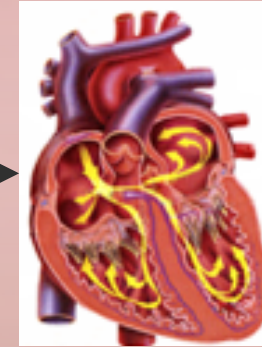
electrocardiogram



Ca²⁺ ion channels

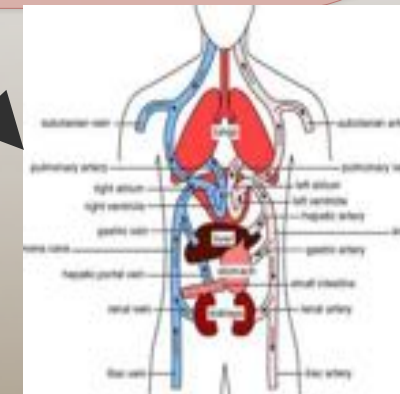
whole cell

myocardium



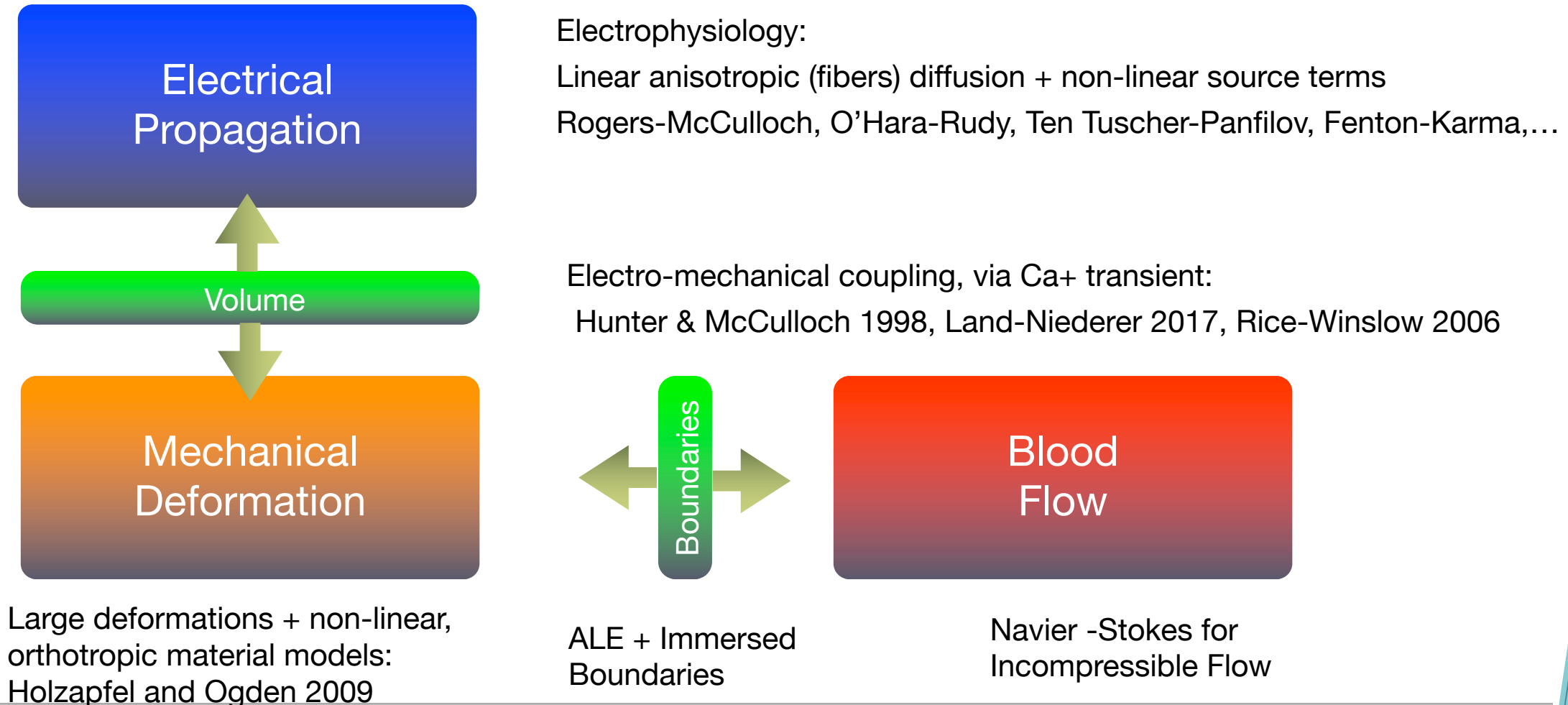
sarcomere

Biomechanics



circulation

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

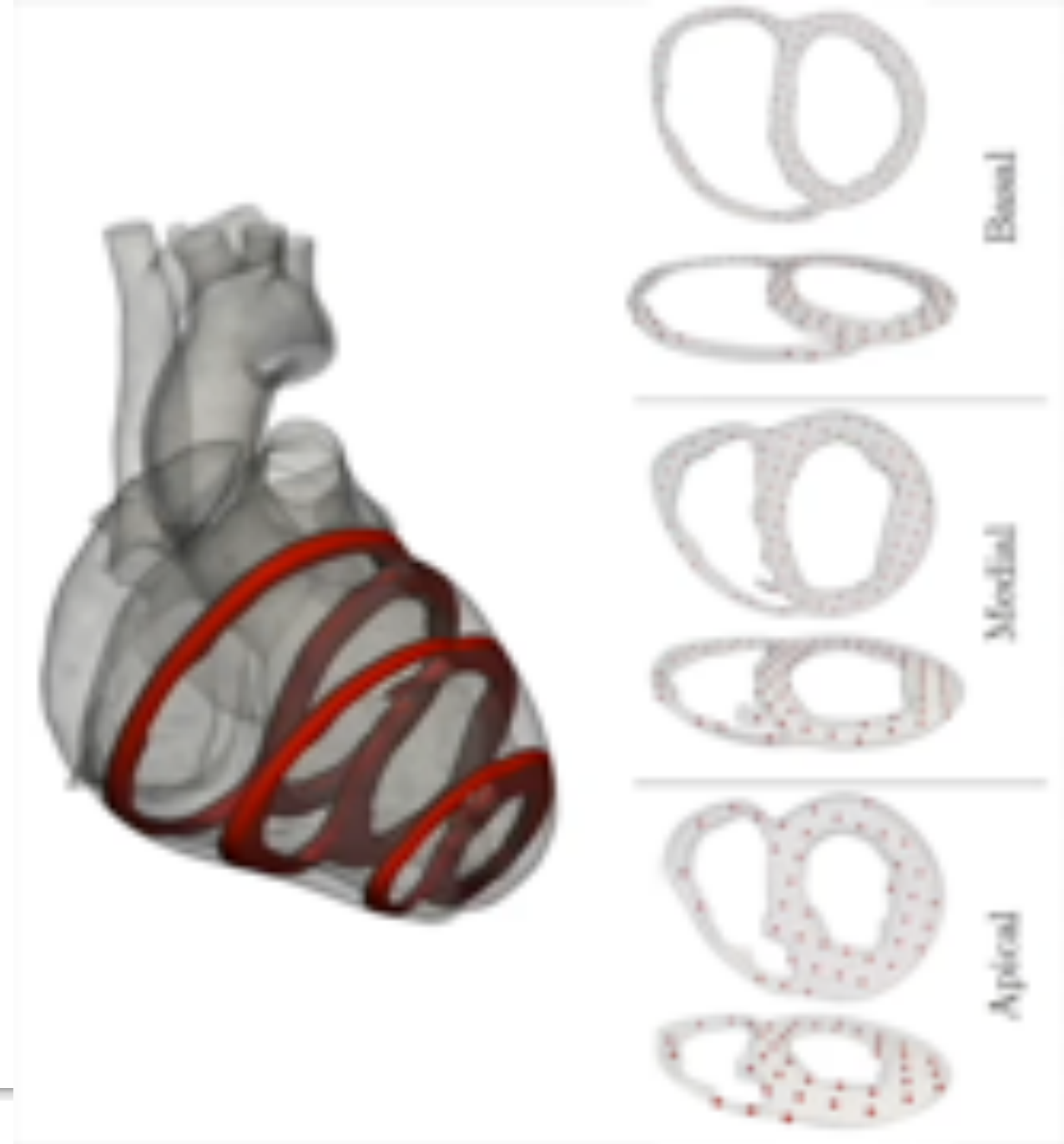
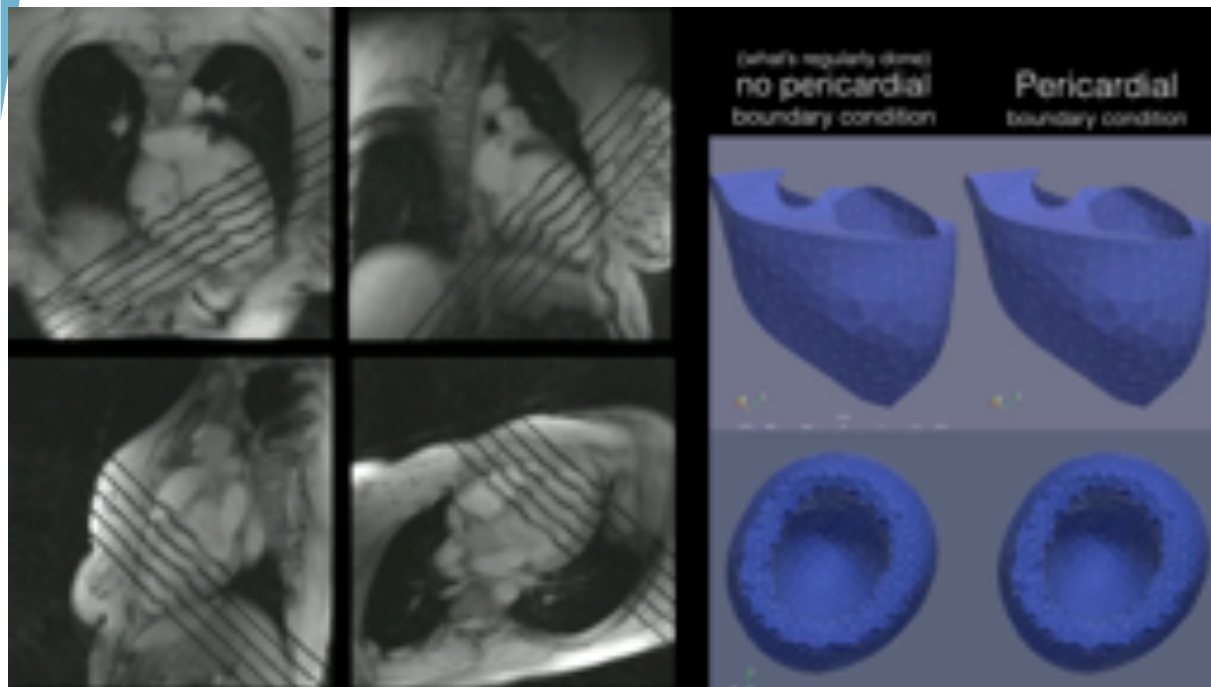


Fully Coupled Electro-Mechanic-Fluid simulation

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

Segmentation and Surface
representation

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

Biventricular Detailed
Octree
Volumetric Meshes

MAXIMUM ELEMENT SIDE LENGTH: 0.4
mm



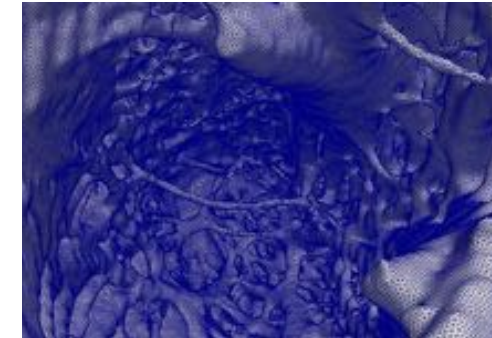
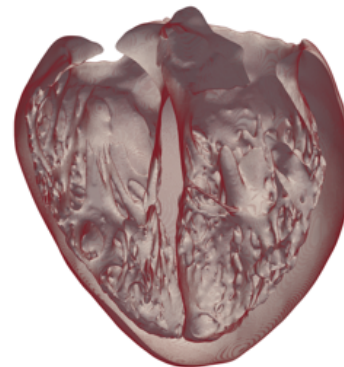
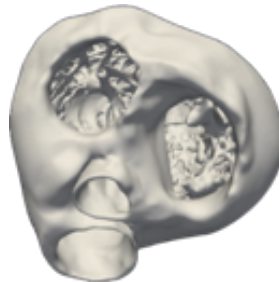
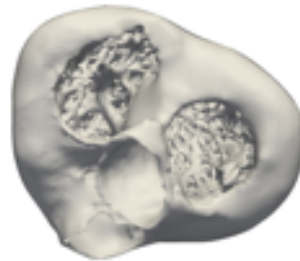
● Male Heart
● Female Heart



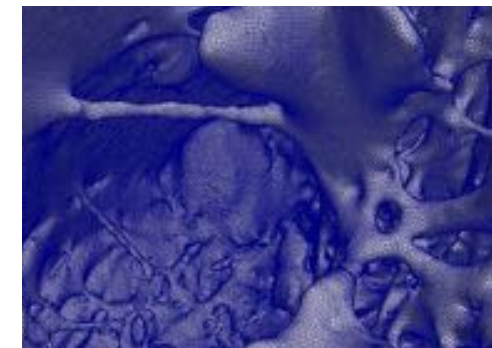
AUTODESK
MESHMIXER



ReMESH



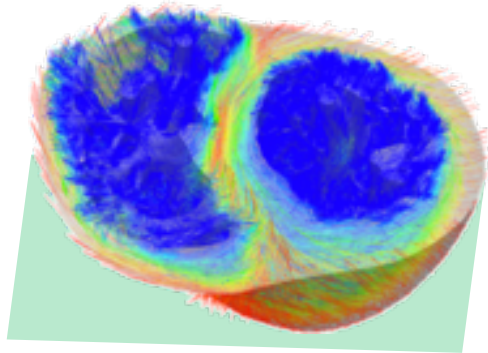
n° elements: 86.318.429
Volume: 394.2 cm³
n° points: 14.994.563



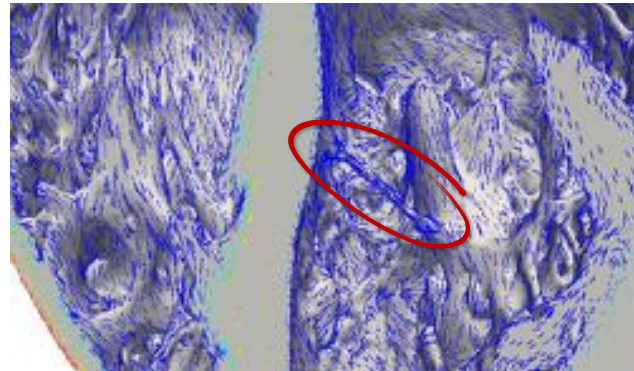
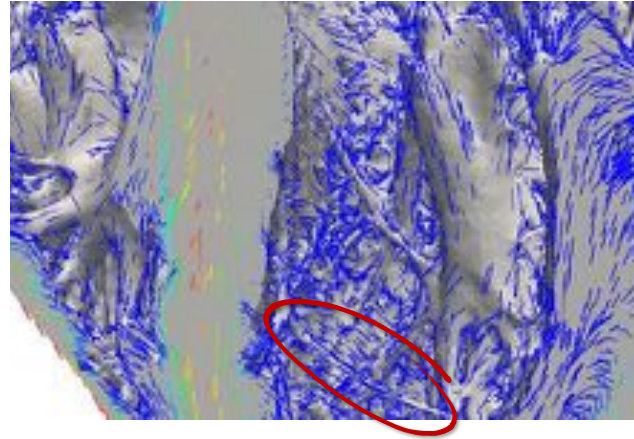
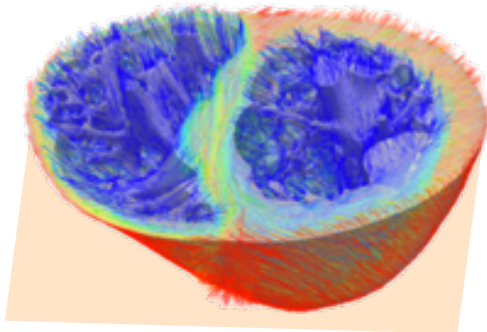
n° elements: 65.501.799
Volume: 299.2 cm³
n° points: 11.416.445

Fiber orientations

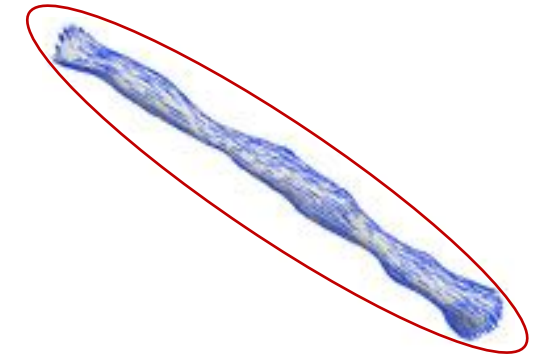
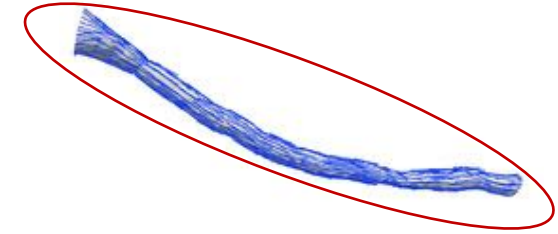
● Male Heart



● Female Heart



False Tendons



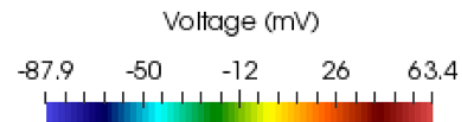
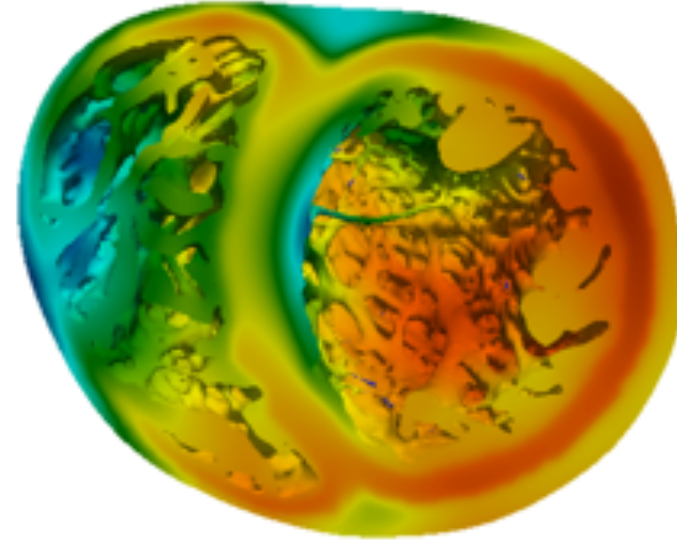
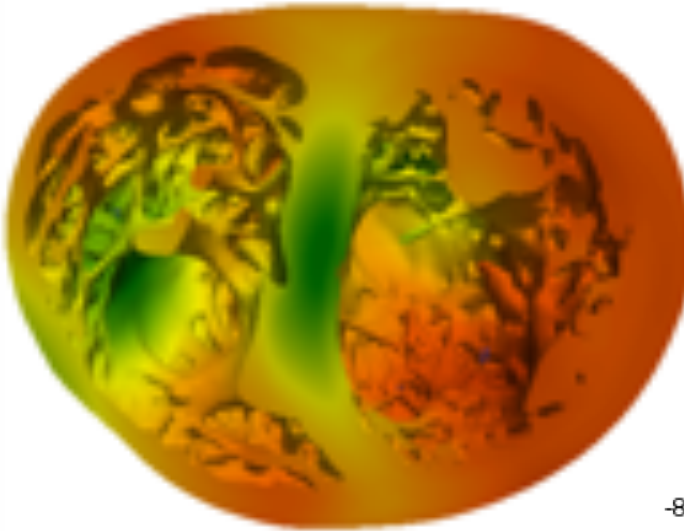
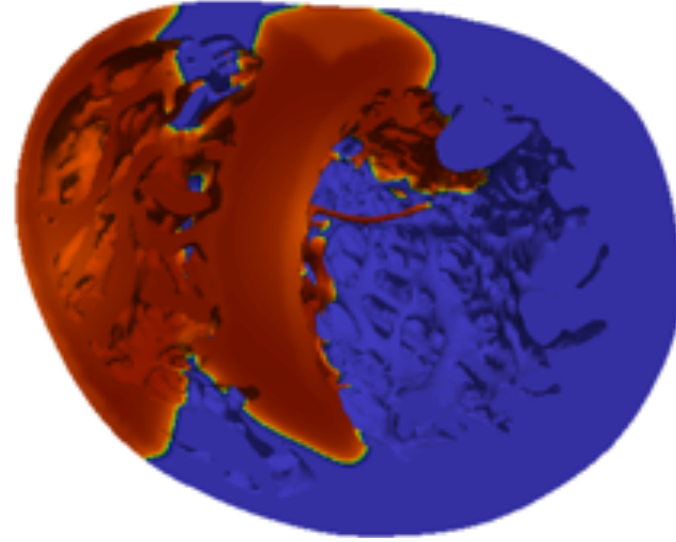
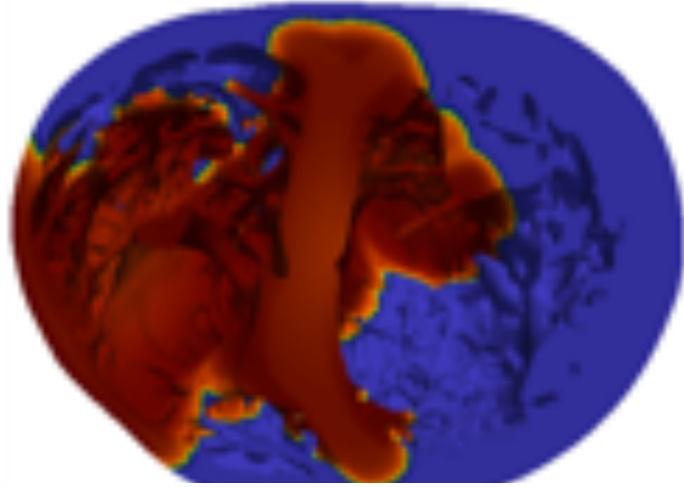
A Rule-Based Method to Model Myocardial Fiber Orientation for Simulating Ventricular Outflow Tract Arrhythmias. Doste et al. FIMH 2017

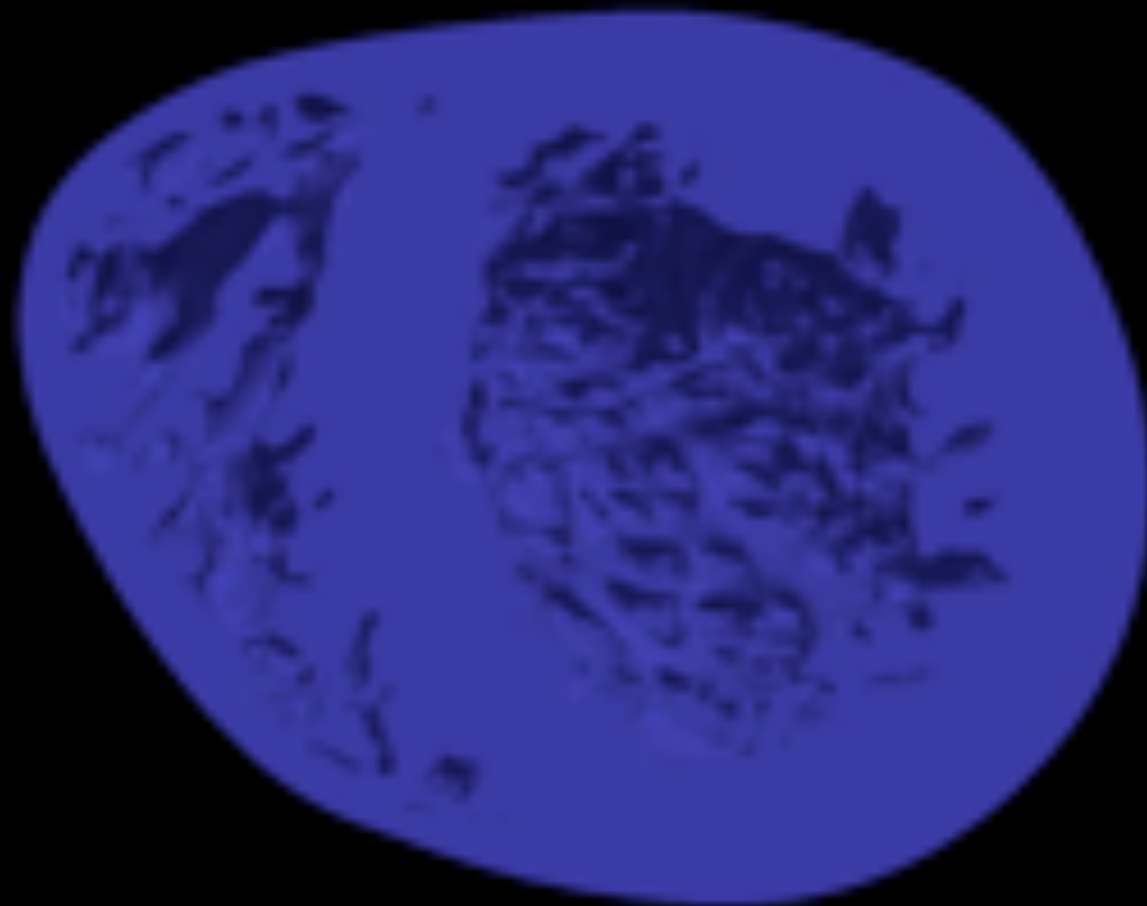
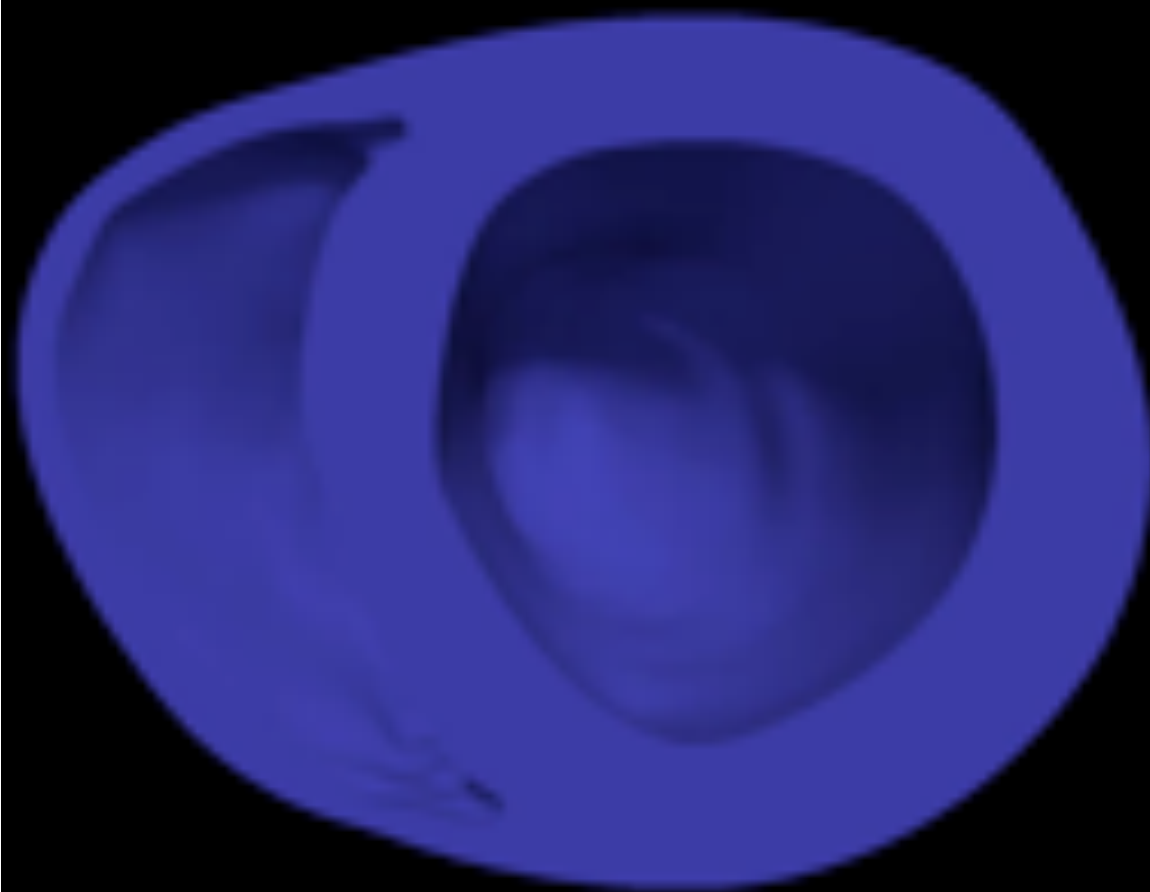


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Barcelona



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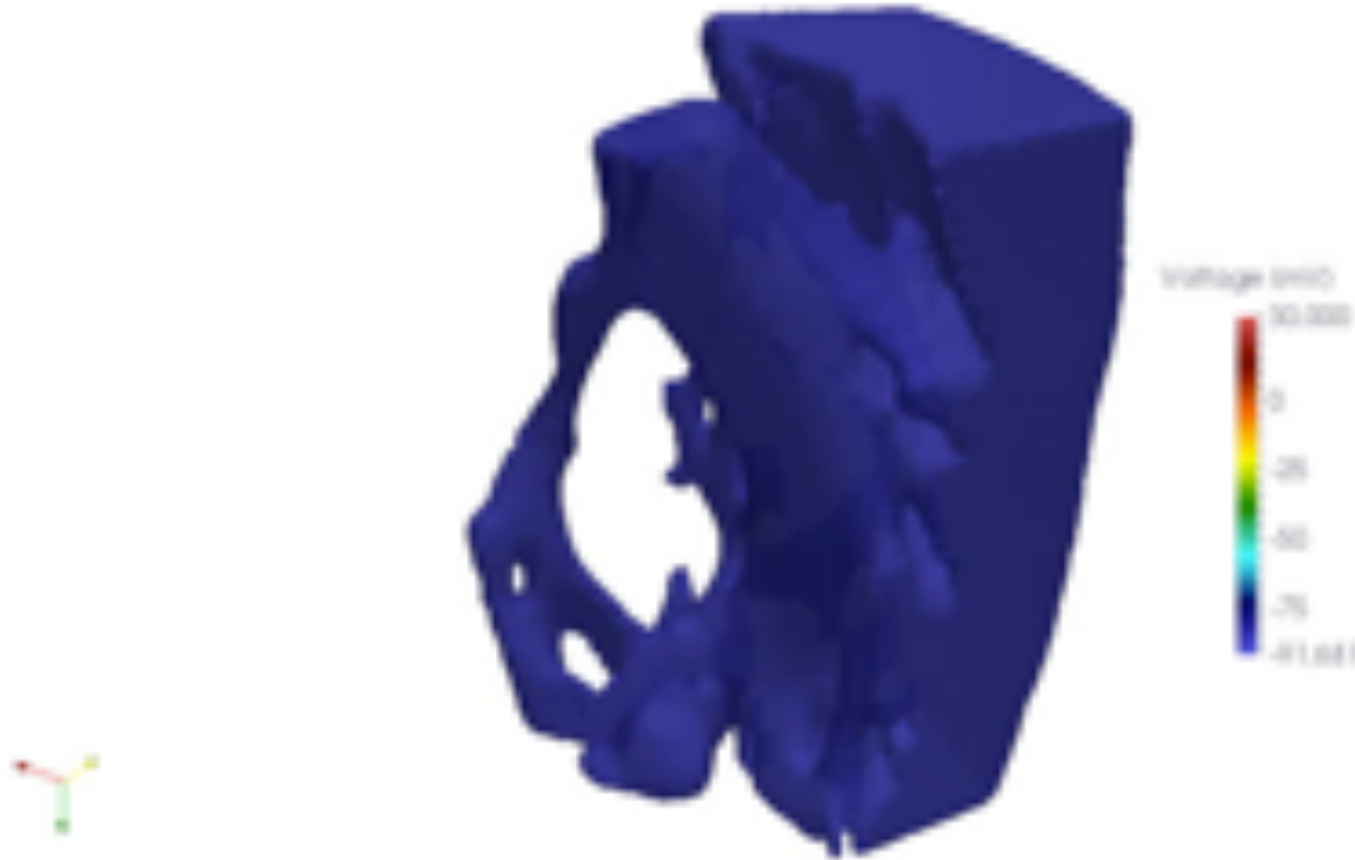




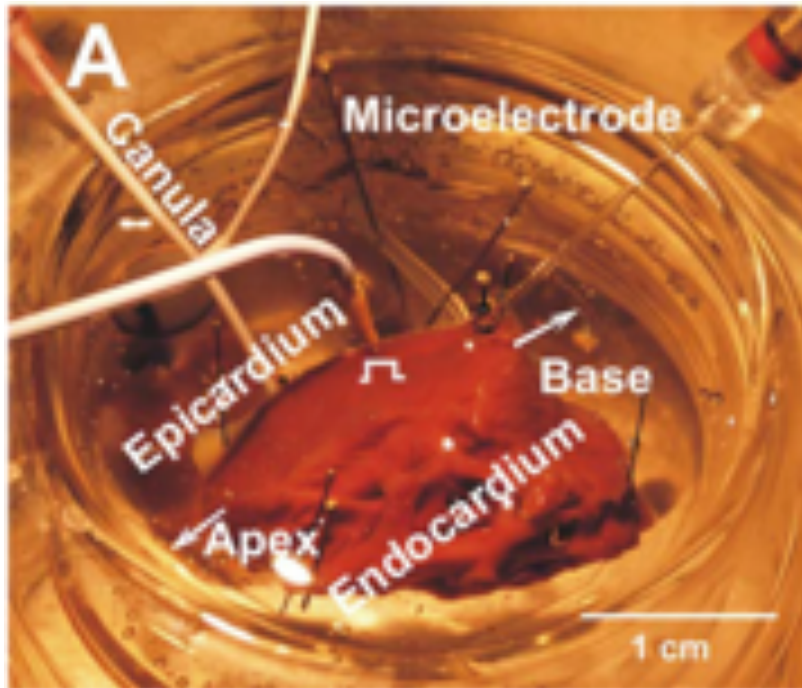
Time: 0.000000



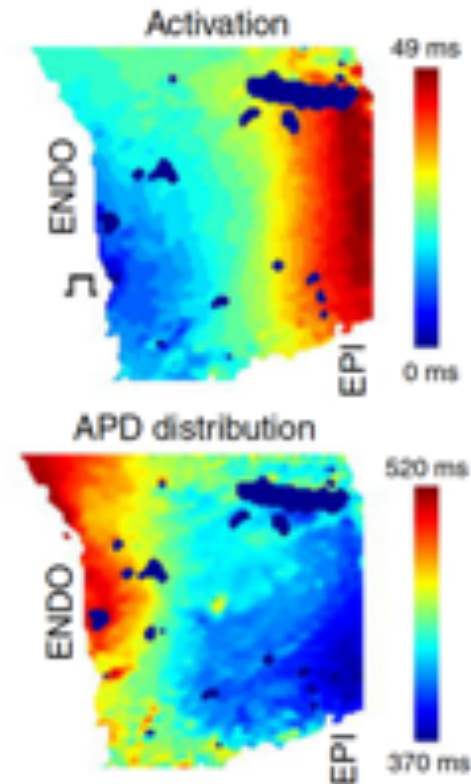
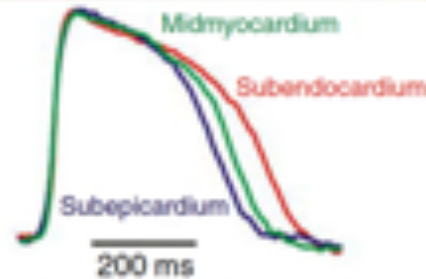
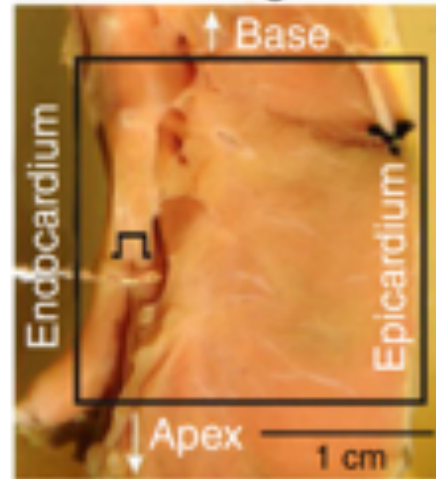
Impact of endocardial structures on electro-mechanics



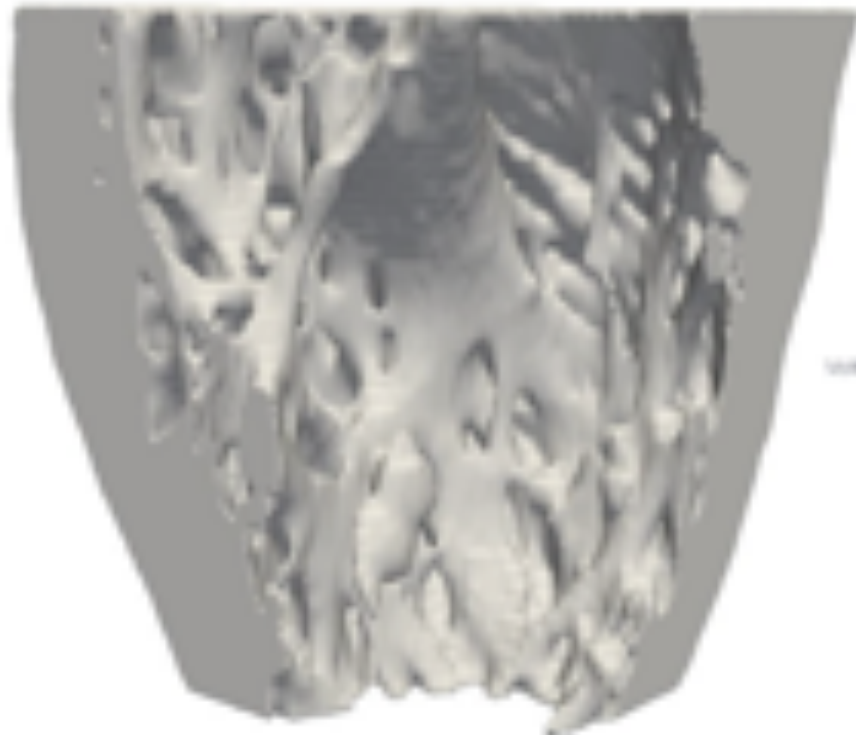
Human Ventricular Wedge Preparation



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

