

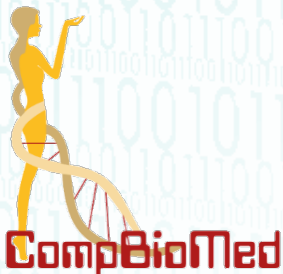
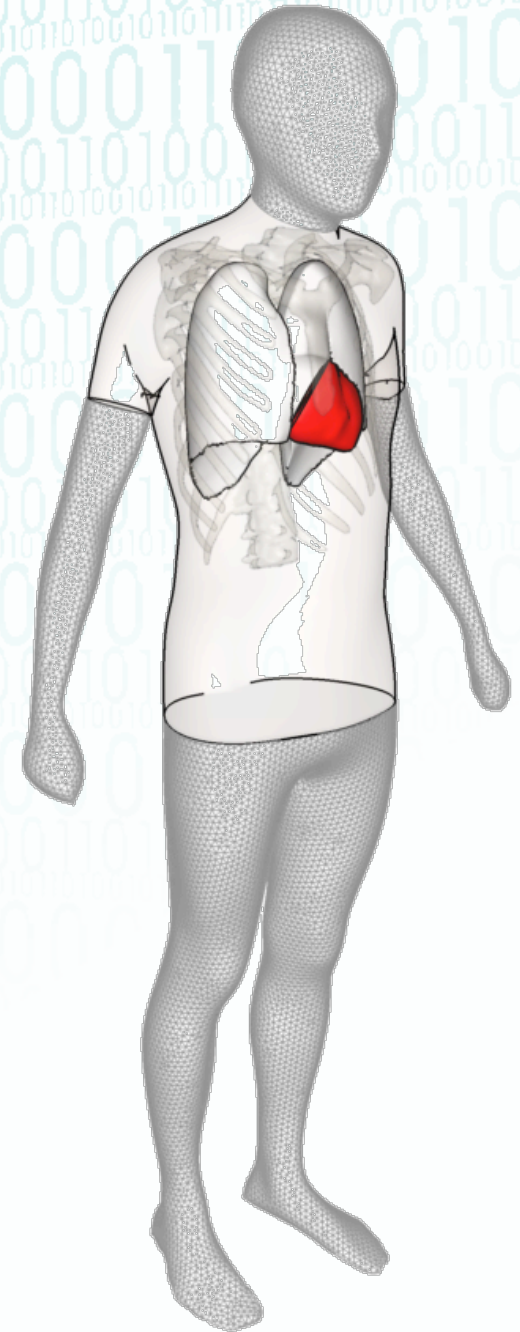
# *HPC simulations of cardiac electrophysiology using patient-specific models of the human heart*

CompBioMed & VPH Institute webinar

**Francesc Levrero-Florencio & Ana Mincholé**

Computational Cardiovascular Science Group

Department of Computer Science, University of Oxford



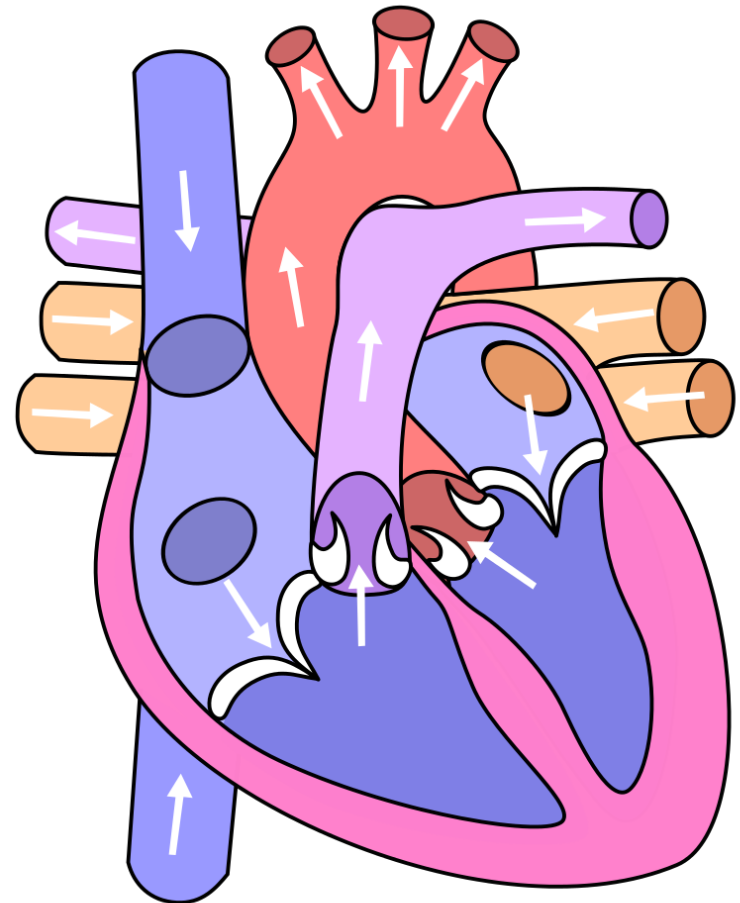
# Contents

- Introduction to the heart (electro)physiology
- Mathematical modelling
- CHASTE
- Alya
- Final Comments

# Brief introduction to (electro)physiology

# Introduction to the physiology of the heart

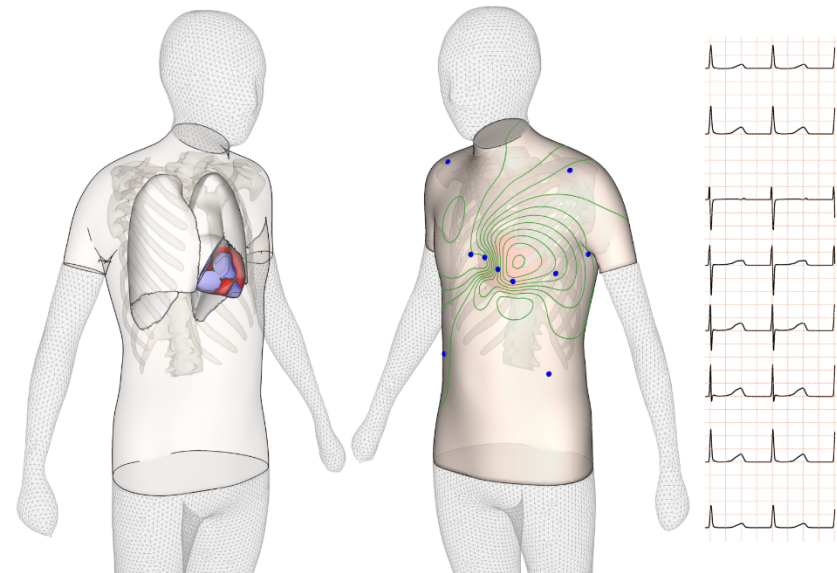
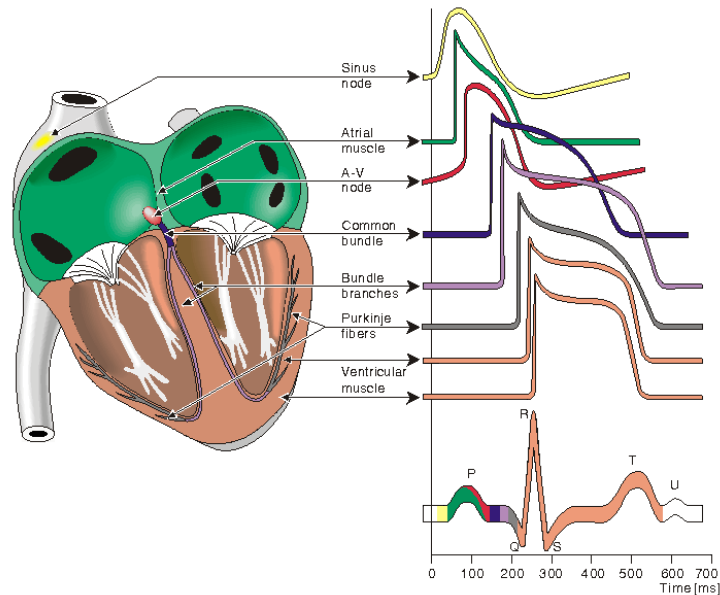
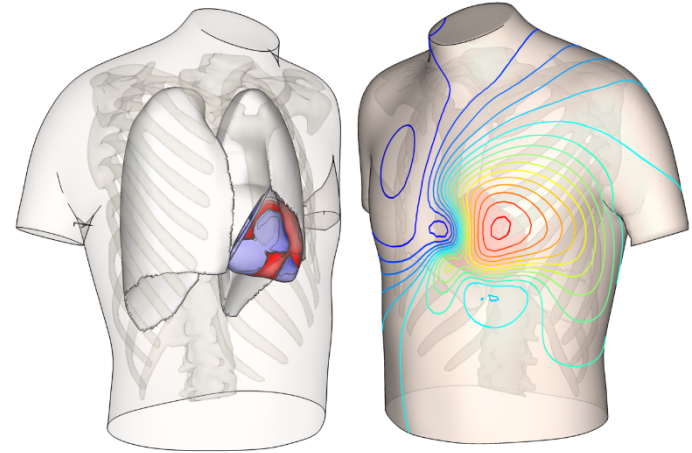
- Heart is a contracting muscle
- 4 chambers
- 2 systems: pulmonary and systemic



{<https://commons.wikimedia.org>}

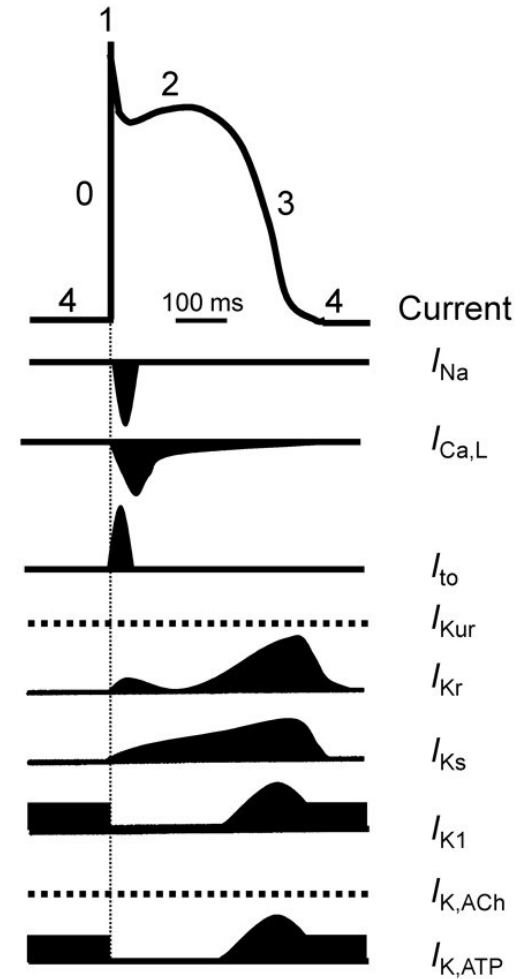
# Electrophysiology of the heart

- SA node – Atria - AV node - brief delay - His bundle branches down the septum
- Purkinje fibers allow propagation throughout the endocardium
- Cell to cell propagation through gap junctions.



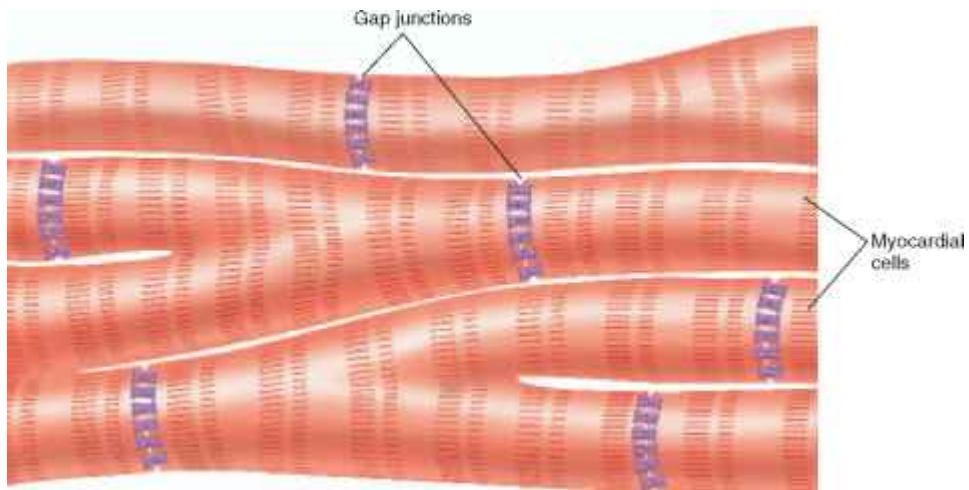
# Cell electrophysiology

- Four phases of the action potential:
  - ✓ Upstroke (0/1)
  - ✓ Plateau (2)
  - ✓ Repolarisation (3)
  - ✓ Resting potential (4)
- Calcium and Sodium currents are involved during the upstroke and plateau phases
- Potassium-based currents are involved in the repolarisation



{Ravens U and Cerbai E. Europace 2008}

# Tissue electrophysiology



{Essentials of Human Physiology 2010}



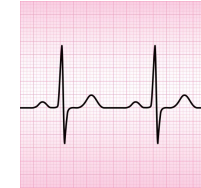
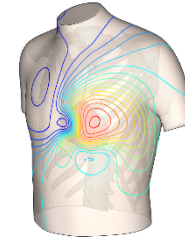
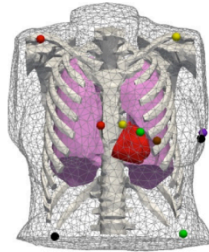
Wellcome Images

# Cardiac modelling

MULTISCALE INTEGRATION

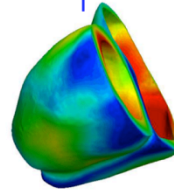
## Body

Body surface potentials  
ECG recordings  
mHealth data



## Organ

Clinical EP studies  
Multi-modality clinical MRI



### Propagation model

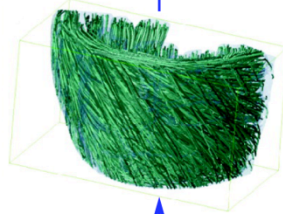
$$\nabla \cdot \bar{\sigma}_i \nabla \Phi_i = \beta \cdot I_m - I_{st,j}$$

$$\nabla \cdot \bar{\sigma}_e \nabla \Phi_e = -\beta \cdot I_m$$

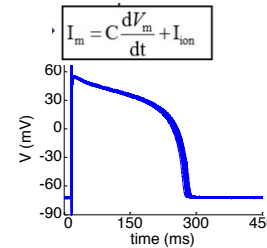


## Tissue

High-resolution MRI  
Histology  
Optical mapping  
Cell cultures

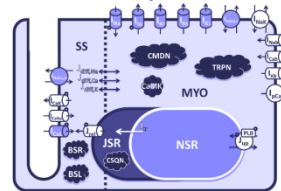


### AP cell model



## Single cell

Microelectrode recordings  
Protein/mRNA expression  
Optical mapping



### Ionic currents

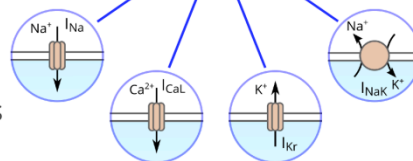
$$I_{Na} = g_{Na_{max}} * m^3 * h^j * (V_m - E_{Na})$$

$$\frac{dx}{dt} = \alpha_x(1-x) - \beta_x x$$

$$\alpha_x = \alpha_x(V_m); \beta_x = \beta_x(V_m)$$

## Ionic current

Voltage/patch clamp  
Isolated cells/hiPSC-CMs



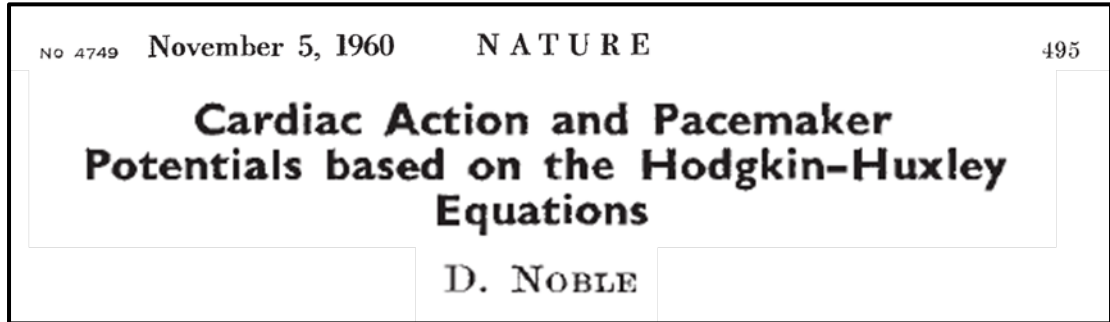
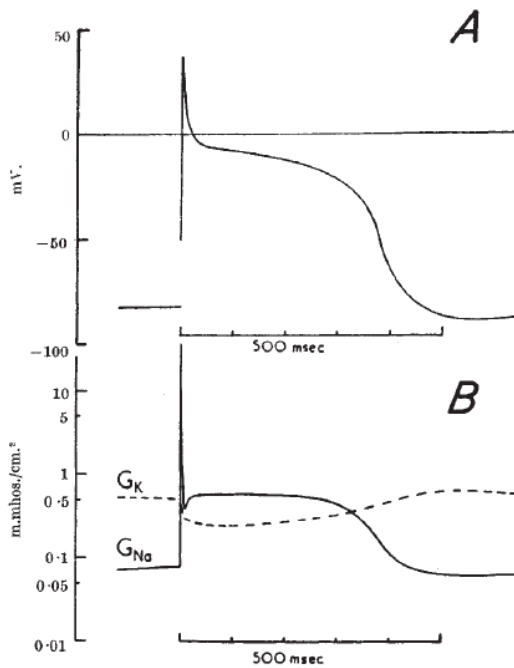


# Mathematical modelling

# First cardiac AP model



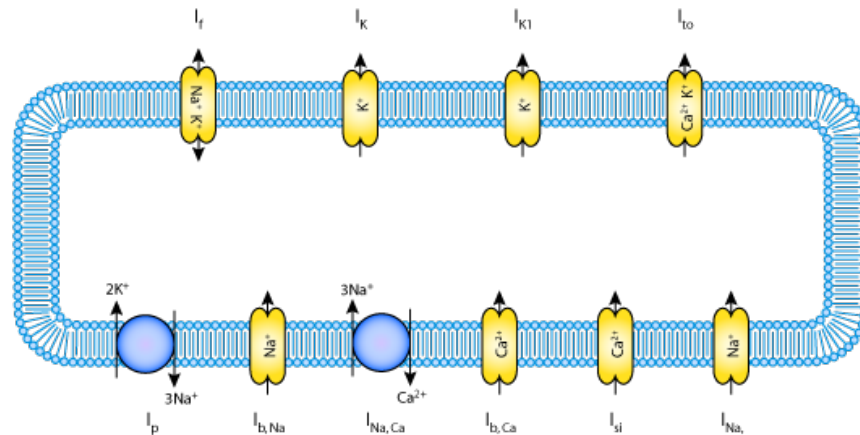
**Denis Noble**  
(1936-present)



*Phil. Trans. R. Soc. Lond. B* 307, 353-398 (1985) [ 353 ]  
Printed in Great Britain

## A MODEL OF CARDIAC ELECTRICAL ACTIVITY INCORPORATING IONIC PUMPS AND CONCENTRATION CHANGES

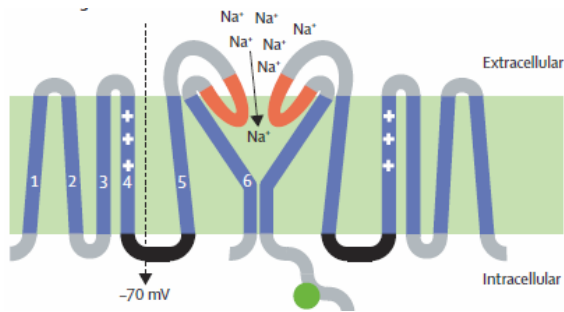
BY D. DiFRANCESCO<sup>1</sup> AND D. NOBLE, F.R.S.<sup>2</sup>



# ... to current human AP models

- ▶ Based on experimental data from >150 human hearts.
- ▶ Still largely based on Hodgkin-Huxley model and its formulation of voltage-gated ion channel behaviour.

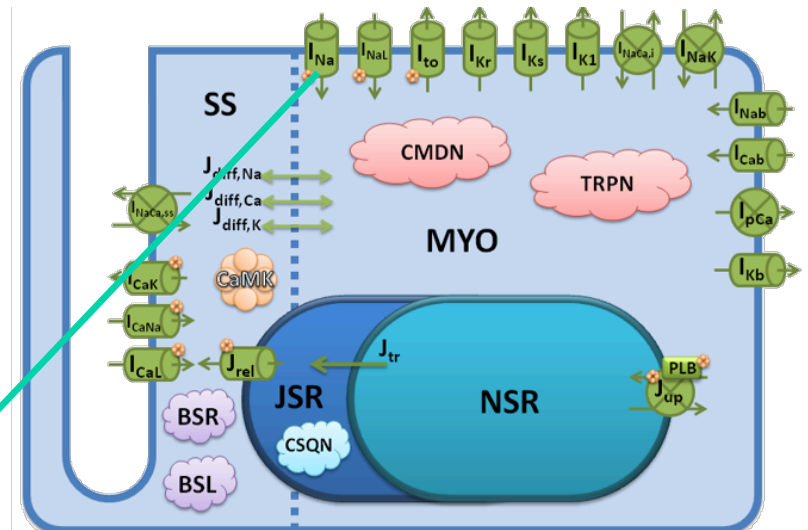
$$C_m \frac{dV_m}{dt} = I_{stim} - I_{currents}$$



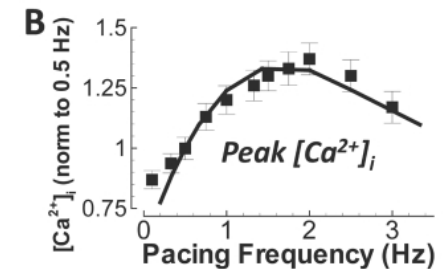
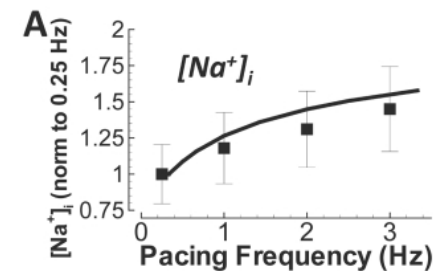
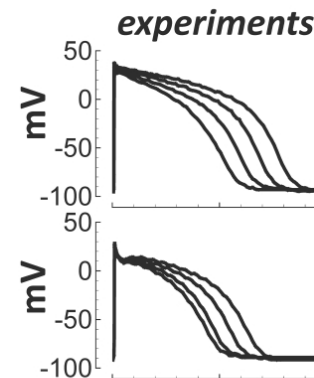
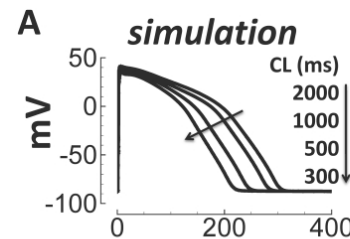
Sodium channel schematic

Sodium channel equation

$$I_{Na} = g_{Na} m^3 h (V_m - E_{Na})$$



{O'Hara et al. PLOS Comput Biol. 2011}



# Monodomain and bidomain models

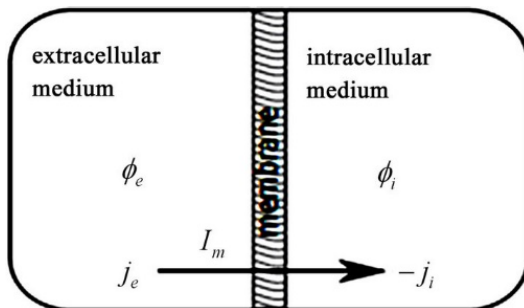
## Monodomain model

$$\nabla \cdot (\mathbf{D}_0 \nabla V) = \chi \left( C \frac{dV}{dt} + I_{ion} - I_{app} \right)$$

## Bidomain model

$$\nabla \cdot (\mathbf{D}_{int} \nabla V) + \nabla \cdot (\mathbf{D}_{int} \nabla U) = \chi \left( C \frac{dV}{dt} + I_{ion} - I_{app} \right)$$

$$\nabla \cdot [(\mathbf{D}_{int} + \mathbf{D}_{ext}) \nabla U] + \nabla \cdot (\mathbf{D}_{int} \nabla V) = 0$$



{Adebisi et al. J Biomed Sci Eng. 2013}

## Conductivity Tensors



➤ The ratio of the intracellular and extracellular conductivity tensors;

Bidomain:	Monodomain:	$\mathbf{D}_i = \begin{pmatrix} D_{  }^i & 0 & 0 \\ 0 & D_{\perp}^i & 0 \\ 0 & 0 & D_{\perp}^i \end{pmatrix}$
$\frac{D_{  }^i}{D_{  }^e} \neq \frac{D_{\perp}^i}{D_{\perp}^e}$	$\frac{D_{  }^i}{D_{  }^e} = \frac{D_{\perp}^i}{D_{\perp}^e} = \alpha$	$\mathbf{D}_e = \begin{pmatrix} D_{  }^e & 0 & 0 \\ 0 & D_{\perp}^e & 0 \\ 0 & 0 & D_{\perp}^e \end{pmatrix}$

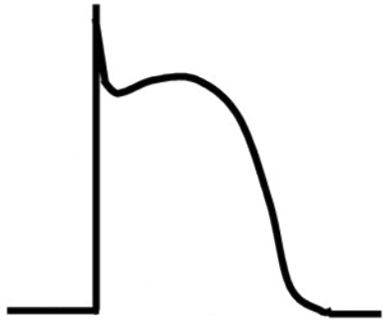
Cardiac tissue is more accurately described as a three-dimensional anisotropic *bidomain*, especially under conditions of applied external current such as in defibrillation studies.<sup>[1-2]</sup>

[1] B. J. Roth and J. P. Wikswo, IEEE Transactions on Biomedical Engineering **41**, 232-240 (1994)

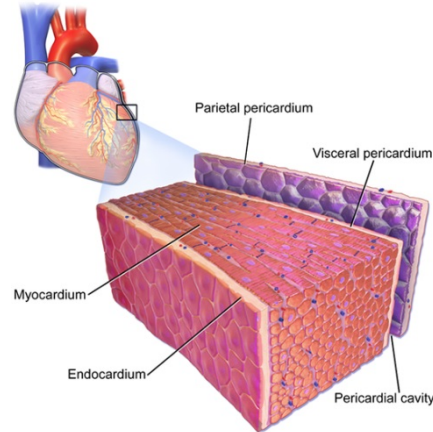
[2] J. P. Wikswo, et al., Biophysical Journal **69**, 2195-2210 (1995)

# Integrative physiology through modelling

Cellular  
Electrophysiology



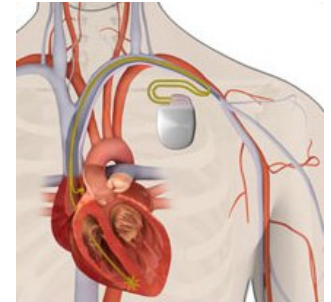
Tissue Properties



Membrane  
Kinetics



Electrical  
Stimulation



$$C_m \frac{\partial V_m}{\partial t}$$

=

$$\frac{1}{S} \nabla \cdot \sigma \nabla V_m$$

-

$$I_{ion}$$

+

$$I_{stim}$$

Propagation of the electrical impulse

# Considered simulation software



- ✓ 0D, 1D, 2D and 3D
- ✓ Petsc and Metis
- ✓ C++
- ✓ FE, BE, RK, CVODE
- ✓ Multi-scale simulation
- ✓ Highly scalable



- ✓ 0D, 1D, 2D and 3D
- ✓ In-house and Metis
- ✓ Modern Fortran
- ✓ FE
- ✓ Multiphysics
- ✓ Highly scalable (up to 100k cores)



DEPARTMENT OF  
**COMPUTER  
SCIENCE**



**Barcelona  
Supercomputing  
Center**

*Centro Nacional de Supercomputación*

# Example of electrophysiology with Chaste



**Chaste**

- **Cardiac Chaste functionalities:**
  - ✓ Monodomain and bidomain models
  - ✓ Automatic implementation of cellular action potential models from the CellML repository
  - ✓ Automatic generation of mathematical model for fibre orientation
  - ✓ Checkpoint of simulations midway through run and restart with altered parameters
  - ✓ Post-processing of simulation results to calculate electrophysiological properties such as action potential duration, conduction velocity, etc.

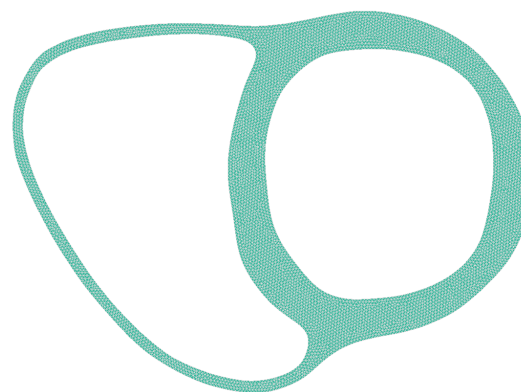
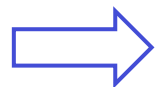


# 2D electrical propagation using Chaste

- 2D Mesh geometry



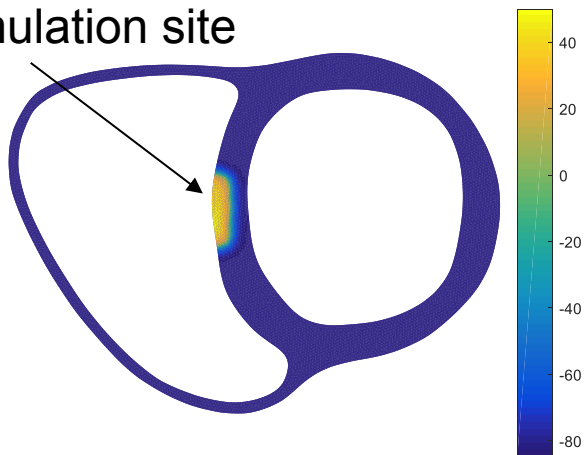
Courtesy of Dr. Rina Ariga



Courtesy of Dr. Ernesto Zacur

- Control 2D Electrical propagation

Stimulation site



- 2D tissue from MRI
- O'Hara Rudy 2011 epicardial model
- Bidomain model
- Isotropic conductivities
- Regular stimulus of 600 ms

```

<<ctxtest/TestSuite.h>
#include "PetscSetupAndFinalize.hpp"

#include "BidomainProblem.hpp"
#include "RegularStimulus.hpp"
#include "TetrahedralMesh.hpp"

#include "ORd2011epi_fkatzCvodeOpt.hpp"

class MyCellFactory : public AbstractCardiacCellFactory<2>
{
private:
    boost::shared_ptr<RegularStimulus> mpStimulus;

public:
    MyCellFactory(float period)
        : AbstractCardiacCellFactory<2>(),
          mpStimulus(new RegularStimulus(-120000.0, 1, period, 0))
    {
    }

    AbstractCvodeCell* CreateCardiacCellForTissueNode(Node<2>* pNode)
    {
        CellORd2011epi_fkatzFromCellMLCvodeOpt* p_cell;

        double x = pNode->rGetLocation()[0];
        double y = pNode->rGetLocation()[1];

        if ( (x < -0.32) && (x < -0.27) && (y > -0.1) && (y < 0.1) )
        {
            p_cell = new CellORd2011epi_fkatzFromCellMLCvodeOpt(mpSolver, mpStimulus);
        }
        else
        {
            p_cell = new CellORd2011epi_fkatzFromCellMLCvodeOpt(mpSolver, mpZeroStimulus);
        }
        return p_cell;
    }
};

class TestConductivity : public CxxTest::TestSuite
{
public:
    void TestConductivity2d() throw(Exception)
    {
        // MESH
        std::string filepath = "projects/anamin/test/data/2Dmesh";
        DistributedTetrahedralMesh<2,2> mesh;
        TrianglesMeshReader<2,2> mesh_reader(filepath);
        mesh.ConstructFromMeshReader(mesh_reader);

        // NUMERICS
        double odeT, pdeT, Print_Time;
        odeT = 0.025;
        pdeT = 0.05;
        Print_Time=1; // ms

        HeartConfig::Instance()->SetOdePdeAndPrintingTimeSteps(odeT, pdeT, Print_Time);

        // BIDOMAIN PARAMETERS
        HeartConfig::Instance()->SetSurfaceAreaToVolumeRatio(1400); // 1/cm
        HeartConfig::Instance()->SetCapacitance(1.0); // uF/cm^2
        HeartConfig::Instance()->SetIntracellularConductivities(Create_c_vector(1.2, 1.2));
        HeartConfig::Instance()->SetExtracellularConductivities(Create_c_vector(4, 4));

        // SIMULATION TIME
        HeartConfig::Instance()->SetSimulationDuration(1300); //ms

        //POSTPROCESSING OPTIONS
        // APD map
        std::vector<std::pair<double,double> > apd_map;
        apd_map.push_back(std::pair<double, double>(90.0, 0.0)); // APD90, 0 mV threshold
        HeartConfig::Instance()->SetApdMaps(apd_map);

        // Activation time map
        std::vector<double> upstroke_time_map;
        upstroke_time_map.push_back(0.0); // 0 mV threshold
        HeartConfig::Instance()->SetUpstrokeTimeMaps(upstroke_time_map);

        // Max Upstroke Velocity map
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        // OUTPUT OPTIONS
        HeartConfig::Instance()->SetOutputDirectory("./OUT");
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        HeartConfig::Instance()->SetOutputUsingOriginalNodeOrdering(true);
        HeartConfig::Instance()->SetVisualizeWithMeshalyzer(false);
        HeartConfig::Instance()->SetVisualizeWithParallelVtk(true);

        // PARAMETERS
        double period = 600.0;

        MyCellFactory cell_factory(period);

        // PROBLEM
        BidomainProblem<2> bidomain_problem( &cell_factory );

        bidomain_problem.SetMesh(&mesh);
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        //POSTPROCESSING OPTIONS
        // APD map
        std::vector<std::pair<double,double>> apd_map;
        apd_map.push_back(std::pair<double, double>(90.0, 0.0)); // APD90, 0 mV threshold
        HeartConfig::Instance()->SetApdMaps(apd_map);

        // Activation time map
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        bidomain_problem.SetMesh(&mesh);
        bidomain_problem.SetWriteInfo();
        bidomain_problem.Initialise();
        bidomain_problem.Solve();
    }
};

```

```

#include <cxxtest/TestSuite.h>
#include "PetscSetupAndFinalize.hpp"

#include "BidomainProblem.hpp"
#include "RegularStimulus.hpp"
#include "TetrahedralMesh.hpp"

#include "ORd2011epi_fkatzCvodeOpt.hpp"

```

```

#include <cxtest/TestSuite.h>
#include "PetscSetupAndFinalize.hpp"

#include "BidomainProblem.hpp"
#include "RegularStimulus.hpp"
#include "TetrahedralMesh.hpp"

#include "ORd2011epi_fkatpCvodeOpt.hpp"

class MyCellFactory : public AbstractCardiacCellFactory
{
private:
    boost::shared_ptr<RegularStimulus> mpStimulus;

public:
    MyCellFactory(float period)
        : AbstractCardiacCellFactory<>(),
          mpStimulus(new RegularStimulus(-120000.0, 1, period, 0))
    {
    }

    AbstractCvodeCell* CreateCardiacCellForTissueNode(Node<2>* pNode)
    {
        CellORd2011epi_fkatpFromCellMLCvodeOpt* p_cell;

        double x = pNode->rGetLocation()[0];
        double y = pNode->rGetLocation()[1];

        if ( (x>-0.32) && (x<-0.27) && (y>-0.1) && (y<0.1) )
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpStimulus);
        }
        else
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpZeroStimulus);
        }

        return p_cell;
    }
};

```

```

class MyCellFactory : public AbstractCardiacCellFactory<2>
{
private:
    boost::shared_ptr<RegularStimulus> mpStimulus;

public:
    MyCellFactory(float period)
        : AbstractCardiacCellFactory<2>(),
          mpStimulus(new RegularStimulus(-120000.0, 1, period, 0))
    {
    }

    AbstractCvodeCell* CreateCardiacCellForTissueNode(Node<2>* pNode)
    {
        CellORd2011epi_fkatpFromCellMLCvodeOpt* p_cell;

        double x = pNode->rGetLocation()[0];
        double y = pNode->rGetLocation()[1];

        if ( (x>-0.32) && (x<-0.27) && (y>-0.1) && (y<0.1) )
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpStimulus);
        }
        else
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpZeroStimulus);
        }

        return p_cell;
    }
};

```

```

class TestConductivity : public CxxTest::TestSuite
{
public:
    void TestConductivity2d() throw(Exception)
    {
        // MESH
        std::string filepath = "projects/anamin/TetrahedralMesh2d";
        DistributedTetrahedralMesh<2,2> mesh;
        TrianglesMeshReader<2,2> mesh_reader(filepath);
        mesh.ConstructFromMeshReader(mesh_reader);

        // NUMERICS
        double odeT, pdeT, Print_Time;
        odeT = 0.025;
        pdeT = 0.05;
        Print_Time = 1; // ms

        HeartConfig::Instance()->SetOdePdeAndPrintTime(odeT, pdeT, Print_Time);

        // BIDOMAIN PARAMETERS
        HeartConfig::Instance()->SetSurfaceAreaToVolumeRatio(1);
        HeartConfig::Instance()->SetCapacitance(1);
        HeartConfig::Instance()->SetIntracellularExtracellularVolumeRatio(1);
        HeartConfig::Instance()->SetExtracellularVolume(1);

        // SIMULATION TIME
        HeartConfig::Instance()->SetSimulationDuration(600);

        // POSTPROCESSING OPTIONS
        // APD map
        std::vector<std::pair<double, double>> apd_map;
        apd_map.push_back(std::pair<double, double>(0, 0));
        HeartConfig::Instance()->SetApdMaps(apd_map);

        // Activation time map
        std::vector<double> upstroke_time_map;
        upstroke_time_map.push_back(0.0);
        HeartConfig::Instance()->SetUpstrokeTimeMap(upstroke_time_map);

        // Max Upstroke Velocity map
        std::vector<double> maxupstroke_vel_map;
        maxupstroke_vel_map.push_back(0.0);
        HeartConfig::Instance()->SetMaxUpstrokeVelocityMap(maxupstroke_vel_map);

        // OUTPUT OPTIONS
        HeartConfig::Instance()->SetOutputDirectory("results");
        HeartConfig::Instance()->SetOutputFilename("results");
        HeartConfig::Instance()->SetOutputUsingOriginalNodeOrdering(true);
        HeartConfig::Instance()->SetVisualizeWithMeshalyzer(false);
        HeartConfig::Instance()->SetVisualizeWithParallelVtk(true);

        // PARAMETERS
        double period = 600.0;

        MyCellFactory cell_factory(period);

        // PROBLEM
        BidomainProblem<2> bidomain_problem(&cell_factory);

        bidomain_problem.SetMesh(&mesh);
        bidomain_problem.SetWriteInfo();
        bidomain_problem.Initialise();
        bidomain_problem.Solve();
    }
};

```

```

<<ctest/TestSuite.h>
#include "PetscSetupAndFinalize.hpp"

#include "BidomainProblem.hpp"
#include "RegularStimulus.hpp"
#include "TetrahedralMesh.hpp"

#include "ORd2011epi_fkatzCvodeOpt.hpp"

class MyCellFactory : public AbstractCardiacCellFactory
{
private:
    boost::shared_ptr<RegularStimulus> mpStimulus;

public:
    MyCellFactory(float period)
        : AbstractCardiacCellFactory<>(),
          mpStimulus(new RegularStimulus(-120))
    {
    }

    AbstractCvodeCell* CreateCardiacCellForTetrahedralMesh(
        CellORd2011epi_fkatzFromCellMLCvodeOpt* cell)
    {
        double x = pNode->rGetLocation()[0];
        double y = pNode->rGetLocation()[1];

        if ( (x < -0.32) && (x < -0.27) && (y > 0) )
        {
            p_cell = new CellORd2011epi_fkatz;
        }
        else
        {
            p_cell = new CellORd2011epi_fkatz;
        }
        return p_cell;
    }
};

class TestConductivity : public CxxTest::TestSuite
{
public:
    void TestConductivity2d() throw(Exception)
    {
        // MESH
        std::string filepath = "projects/anamin/test/data/2Dmesh";
        DistributedTetrahedralMesh<2,2> mesh;
        TrianglesMeshReader<2,2> mesh_reader(filepath);
        mesh.ConstructFromMeshReader(mesh_reader);

        // NUMERICS
        double odeT, pdeT, Print_Time;
        odeT = 0.025;
        pdeT = 0.05;
        Print_Time=1; // ms

        HeartConfig::Instance()->SetOdePdeAndPrintingTimeSteps(odeT, pdeT, Print_Time);

        // BIDOMAIN PARAMETERS
        HeartConfig::Instance()->SetSurfaceAreaToVolumeRatio(1400); // 1/cm
        HeartConfig::Instance()->SetCapacitance(1.0); // uF/cm^2
        HeartConfig::Instance()->SetIntracellularConductivities(Create_c_vector(0.1, 0.1));
        HeartConfig::Instance()->SetExtracellularConductivities(Create_c_vector(0.4, 0.4));

        // SIMULATION TIME
        HeartConfig::Instance()->SetSimulationDuration(1200); //ms

        //POSTPROCESSING OPTIONS
        // APD map
        std::vector<std::pair<double,double>> apd_map;
        apd_map.push_back(std::pair<double,double>(90.0, 0.0));
        HeartConfig::Instance()->SetApdMaps(apd_map);

        // Activation time map
        std::vector<double> upstroke_time_map;
        upstroke_time_map.push_back(0.0);
        HeartConfig::Instance()->SetUpstrokeTimeMap(upstroke_time_map);

        // Max Upstroke Velocity map
        std::vector<double> maxupstroke_vel_map;
        maxupstroke_vel_map.push_back(0.0);
        HeartConfig::Instance()->SetMaxUpstrokeVelocityMap(maxupstroke_vel_map);

        // OUTPUT OPTIONS
        HeartConfig::Instance()->SetOutputDirectory("./OUT");
        HeartConfig::Instance()->SetOutputFilenamePrefix("results");
        HeartConfig::Instance()->SetOutputUsingOriginalNodeOrdering(true);
        HeartConfig::Instance()->SetVisualizeWithMeshalyzer(false);
        HeartConfig::Instance()->SetVisualizeWithParallelVtk(true);

        // PARAMETERS
        double period = 600.0;
        MyCellFactory cell_factory(period);

        // PROBLEM
        BidomainProblem<> bidomain_problem( &cell_factory );

        bidomain_problem.SetMesh(&mesh);
        bidomain_problem.SetWriteInfo();
        bidomain_problem.Initialise();
        bidomain_problem.Solve();
    }
};

```

```

<cxtest/TestSuite.h>
#include "PetscSetupAndFinalize.hpp"

#include "BidomainProblem.hpp"
#include "RegularStimulus.hpp"
#include "TetrahedralMesh.hpp"

#include "ORd2011epi_fkatzvodeOpt.hpp"

class MyCellFactory : public AbstractCardiacCellFactory
{
private:
    boost::shared_ptr<RegularStimulus> mpStimulus;

public:
    MyCellFactory(float period)
        : AbstractCardiacCellFactory<>(),
          mpStimulus(new RegularStimulus(-120000.0,
        {
        }

    AbstractCvodeCell* CreateCardiacCellForTissueNode(
    {
        CellORd2011epi_fkatzvodeOpt* p_cvode
        {
            double x = pNode->rGetLocation()[0];
            double y = pNode->rGetLocation()[1];

            if ( (x<-0.32) && (x<-0.27) && (y>-0.1) &&
            {
                p_cell = new CellORd2011epi_fkatzvodeOpt* p_cvode
            }
            else
            {
                p_cell = new CellORd2011epi_fkatzvodeOpt* p_cvode
            }
            return p_cell;
        }
    }
};

```

```

class TestConductivity : public CxxTest::TestSuite
{
public:
    void TestConductivity2d() throw(Exception)
    {
        // MESH
        std::string filepath = "projects/anamin/tes
        DistributedTetrahedralMesh<2,2> mesh;
        TrianglesMeshReader<2,2> mesh_reader(filepa
        mesh.ConstructFromMeshReader(mesh_reader);

        // NUMERICS
        double odeT, pdeT, Print_Time;
        odeT = 0.025;
        pdeT = 0.05;
        Print_Time=1; // ms

        HeartConfig::Instance()->SetOdePdeAndPrintI

        // BIDOMAIN PARAMETERS
        HeartConfig::Instance()->SetSurfaceAreaToVo
        HeartConfig::Instance()->SetCapacitance(1.0
        HeartConfig::Instance()->SetIntracellularCo
        HeartConfig::Instance()->SetExtracellularCo

        // SIMULATION TIME
        HeartConfig::Instance()->SetSimulationDurat

        //POSTPROCESSING OPTIONS
        // APD map
        std::vector<std::pair<double,double> > apd
        apd_map.push_back(std::pair<double, double>
        HeartConfig::Instance()->SetApdMaps(apd_map)

        // Activation time map
        std::vector<double> upstroke_time_map;
        upstroke_time_map.push_back(0.0);
        HeartConfig::Instance()->SetUpstrokeTimeMap

        // Max Upstroke Velocity map
        std::vector<double> maxupstroke_vel_map;
        maxupstroke_vel_map.push_back(0.0);
        HeartConfig::Instance()->SetMaxUpstrokeVelo

        // OUTPUT OPTIONS
        HeartConfig::Instance()->SetOutputDirectory
        HeartConfig::Instance()->SetOutputFilenameP
        HeartConfig::Instance()->SetOutputUsingOrig
        HeartConfig::Instance()->SetVisualizeWithVtk
        HeartConfig::Instance()->SetVisualizeWithParallelVtk(true);
    }

    // PARAMETERS
    double period = 600.0;
    MyCellFactory cell_factory(period);

    // PROBLEM
    BidomainProblem<2> bidomain_problem( &cell_factory );

    bidomain_problem.SetMesh(&mesh);
    bidomain_problem.SetWriteInfo();
    bidomain_problem.Initialise();
    bidomain_problem.Solve();
}
};

```

**//POSTPROCESSING OPTIONS**

**// APD map**

```

std::vector<std::pair<double,double> > apd_map;
apd_map.push_back(std::pair<double, double>(90.0, 0.0)); // APD90, 0 mV threshold
HeartConfig::Instance()->SetApdMaps(apd_map);

```

**// Activation time map**

```

std::vector<double> upstroke_time_map;
upstroke_time_map.push_back(0.0); // 0 mV threshold
HeartConfig::Instance()->SetUpstrokeTimeMaps(upstroke_time_map);

```

**// Max Upstroke Velocity map**

```

std::vector<double> maxupstroke_vel_map;
maxupstroke_vel_map.push_back(0.0);
HeartConfig::Instance()->SetMaxUpstrokeVelocityMaps(maxupstroke_vel_map);

```

**// OUTPUT OPTIONS**

```

HeartConfig::Instance()->SetOutputDirectory("./OUT");
HeartConfig::Instance()->SetOutputFilenamePrefix("results");
HeartConfig::Instance()->SetOutputUsingOriginalNodeOrdering(true);
HeartConfig::Instance()->SetVisualizeWithMeshalyzer(false);
HeartConfig::Instance()->SetVisualizeWithParallelVtk(true);

```

**// PARAMETERS**

```

double period = 600.0;
MyCellFactory cell_factory(period);

```

**// PROBLEM**

```

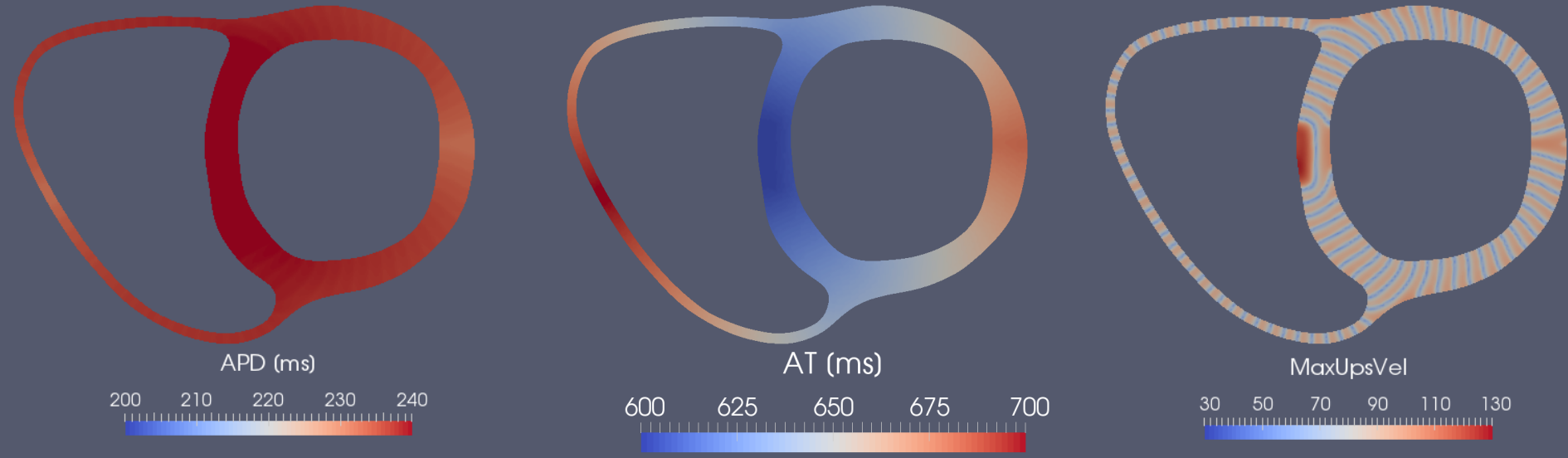
BidomainProblem<2> bidomain_problem( &cell_factory );

```

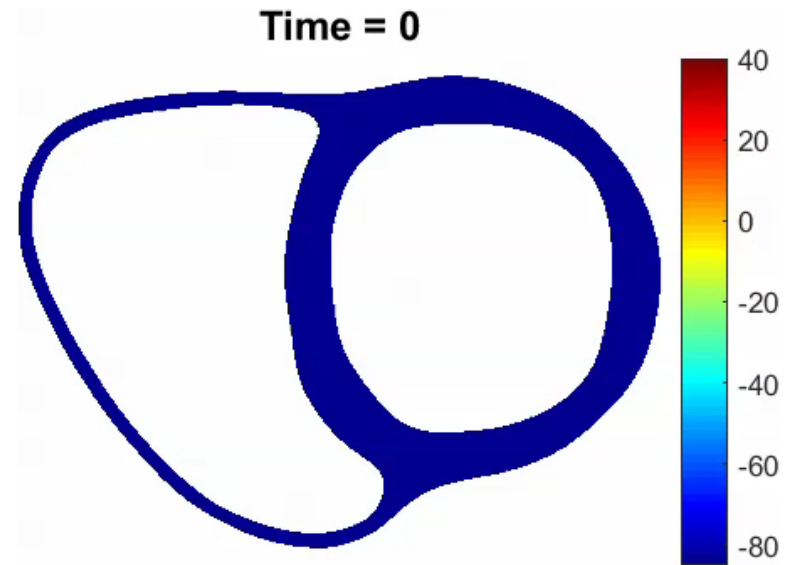
```

bidomain_problem.SetMesh(&mesh);
bidomain_problem.SetWriteInfo();
bidomain_problem.Initialise();
bidomain_problem.Solve();

```



- ✓ Similar APD values in all the geometry
- ✓ Activation time map starting from the septum
- ✓ Similar maximum upstroke velocities





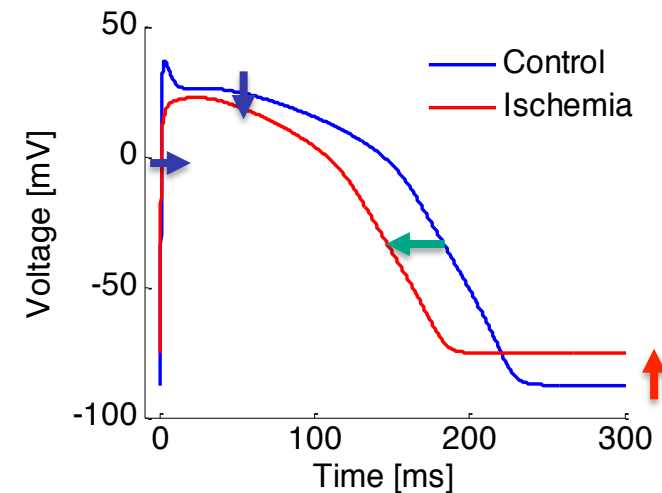
# Chaste example 2

- 2D propagation with ischemia



Ischemia in the anterior heart region:

- **Hyperkalaemia:**  $[K]_o = 8.5 \text{ mM}$
- **Hypoxia:**  $f_{katp} = 0.04$
- **Acidosis:**  $\downarrow 25\% g_{Na}$  and  $g_{CaL}$



```

class MyCellFactory : public AbstractCardiacCellFactory<2>
{
private:
    boost::shared_ptr<RegularStimulus> mpStimulus;
    double k;
    double fkatp;
    double sf_na_ca;

public:
    MyCellFactory(float Input_Ko, float Input_fkatp, float SF_Na_Ca, float period)
        : AbstractCardiacCellFactory<2>(),
          mpStimulus(new RegularStimulus(-120000.0, 1, period, 0))
    {
        k = Input_Ko;
        fkatp = Input_fkatp;
        sf_na_ca = SF_Na_Ca;
    }

    AbstractCvodeCell* CreateCardiacCellForTissueNode(Node<2>* pNode)
    {
        CellORd2011epi_fkatpFromCellMLCvodeOpt* p_cell;

        double x = pNode->rGetLocation()[0];
        double y = pNode->rGetLocation()[1];

        if ( (x>-0.32) && (x<-0.27) && (y>-0.1) && (y<0.1) )
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpStimulus);
        }
        else if ((x>-0.4) && (x<0.05) && (y>0.25) && (y<0.44))
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpZeroStimulus);
            p_cell->SetParameter("extracellular_potassium_concentration",k);
            double gna = p_cell->GetParameter("membrane_fast_sodium_current_conductance");
            double gca = p_cell->GetParameter("membrane_L_type_calcium_current_conductance");
            p_cell->SetParameter("membrane_fast_sodium_current_conductance",gna*sf_na_ca);
            p_cell->SetParameter("membrane_L_type_calcium_current_conductance",gca*sf_na_ca);
            p_cell->SetParameter("membrane_atp_dependent_potassium_current_conductance",fkatp);
        }
        else
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpZeroStimulus);
        }
    }
}

```

```

//POSTPROCESSING OPTIONS
// APD map
std::vector<std::pair<double,double> > apd_map;
apd_map.push_back(std::pair<double, double>(90.0, 0.0)); // APD90, 0 mV threshold
HeartConfig::Instance()->SetApdMaps(apd_map);

// Activation time map
std::vector<double> upstroke_time_map;
upstroke_time_map.push_back(0.0); // 0 mV threshold
HeartConfig::Instance()->SetUpstrokeTimeMaps(upstroke_time_map);

// Max Upstroke Velocity map
std::vector<double> maxupstroke_vel_map;
maxupstroke_vel_map.push_back(0.0);
HeartConfig::Instance()->SetMaxUpstrokeVelocityMaps(maxupstroke_vel_map);

// OUTPUT OPTIONS
HeartConfig::Instance()->SetOutputDirectory("./OUT");
HeartConfig::Instance()->SetOutputFilenamePrefix("results");
HeartConfig::Instance()->SetOutputUsingOriginalNodeOrdering(true);
HeartConfig::Instance()->SetVisualizeWithMeshalyzer(false);
HeartConfig::Instance()->SetVisualizeWithParallelVtk(true);

// PARAMETERS
double Input_Ko           = 8.5;
double Input_fkatp        = 0.05;
double SF_Na_Ca           = 0.75;

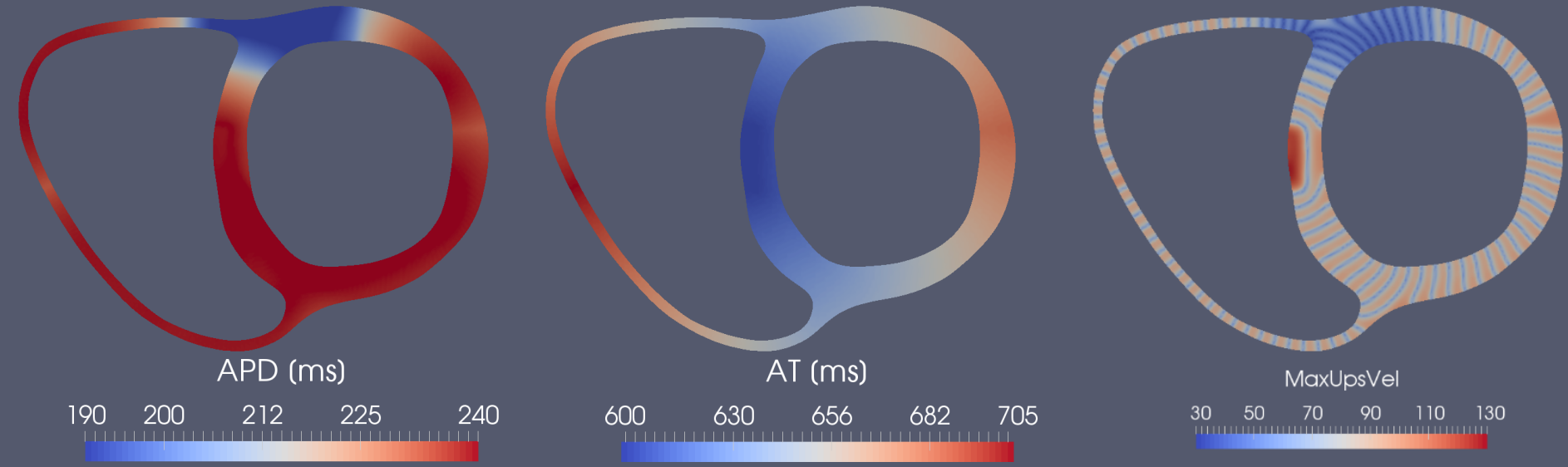
double period              = 600.0;

MyCellFactory cell_factory(Input_Ko, Input_fkatp, SF_Na_Ca, period);

// PROBLEM
BidomainProblem<2> bidomain_problem( &cell_factory );

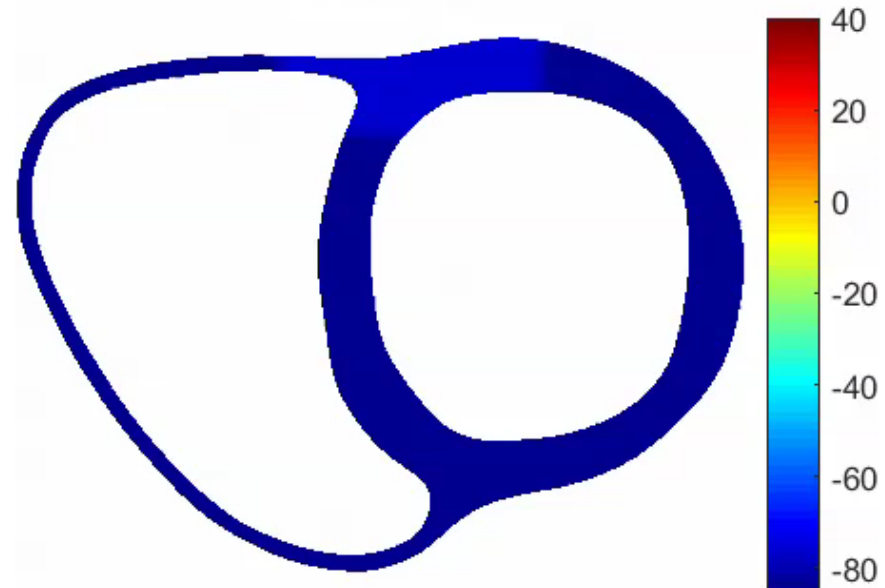
bidomain_problem.SetMesh(&mesh);
bidomain_problem.SetWriteInfo();
bidomain_problem.Initialise();
bidomain_problem.Solve();
}

```



- ✓ Shorter APD values in the ischemic region
- ✓ Activation time map starting from the septum
- ✓ Maximum upstroke velocities slower in the ischemic region

**Time = 0**



# Chaste example 3

- 2D geometry with ischemia and ectopy



Adding the effect of an ectopic beat in the border zone region

```

#include <cxxtest/TestSuite.h>
#include "PetscSetupAndFinalize.hpp"

#include "BidomainProblem.hpp"
#include "RegularStimulus.hpp"
#include "SimpleStimulus.hpp"
#include "TetrahedralMesh.hpp"

#include "ORd2011epi_fkatpCvodeOpt.hpp"

class MyCellFactory : public AbstractCardiacCellFactory<2> // <3> here
{
private:
    boost::shared_ptr<RegularStimulus> mpStimulus;
    boost::shared_ptr<SimpleStimulus> mpStimulus2;
    double k;
    double fkatp;
    double sf_na_ca;

public:
    MyCellFactory(float Input_Ko, float Input_fkatp, float SF_Na_Ca, float period, float ectopic_time)
        : AbstractCardiacCellFactory<2>(),
          mpStimulus(new RegularStimulus(-120000.0, 1, period, 0)),
          mpStimulus2(new SimpleStimulus(-120000.0, 2.0, ectopic_time))
    {
        k = Input_Ko;
        fkatp = Input_fkatp;
        sf_na_ca = SF_Na_Ca;
    }
    AbstractCvodeCell* CreateCardiacCellForTissueNode(Node<2>* pNode)
    {
        CellORd2011epi_fkatpFromCellMLCvodeOpt* p_cell;

        double x = pNode->rGetLocation()[0];
        double y = pNode->rGetLocation()[1];

        if ( (x>-0.32) && (x<-0.27) && (y>-0.1) && (y<0.1) )
        {
            p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpStimulus);
        }
    }
}

```

```

public:
MyCellFactory(float Input_Ko, float Input_fkatp, float SF_Na_Ca, float period, float ectopic_time)
: AbstractCardiacCellFactory<2>(),
  mpStimulus(new RegularStimulus(-120000.0, 1, period, 0)),
  mpStimulus2(new SimpleStimulus(-120000.0, 2.0, ectopic_time))
{
  k = Input_Ko;
  fkatp = Input_fkatp;
  sf_na_ca = SF_Na_Ca;
}
AbstractCvodeCell* CreateCardiacCellForTissueNode(Node<2>* pNode)
{
  CellORd2011epi_fkatpFromCellMLCvodeOpt* p_cell;

  double x = pNode->rGetLocation()[0];
  double y = pNode->rGetLocation()[1];

  if ( (x>-0.32) && (x<-0.27) && (y>-0.1) && (y<0.1) )
  {
    p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpStimulus);
  }

  else if ((x>-0.4) && (x<0.05) && (y>0.25) && (y<0.44))
  {
    p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpZeroStimulus);
    p_cell->SetParameter("extracellular_potassium_concentration",k);
    double gna = p_cell->GetParameter("membrane_fast_sodium_current_conductance");
    double gca = p_cell->GetParameter("membrane_L_type_calcium_current_conductance");
    p_cell->SetParameter("membrane_fast_sodium_current_conductance",gna*sf_na_ca);
    p_cell->SetParameter("membrane_L_type_calcium_current_conductance",gca*sf_na_ca);
    p_cell->SetParameter("membrane_atp_dependent_potassium_current_conductance",fkatp);
  }
  else
  {
    p_cell = new CellORd2011epi_fkatpFromCellMLCvodeOpt(mpSolver, mpZeroStimulus);
  }

  //Definition Ectopic region
  ChastePoint<2> stim_centre (-0.43,0.39);
  ChastePoint<2> stim_radius (0.05,0.05);
  ChasteEllipsoid<2> stim_region (stim_centre,stim_radius);

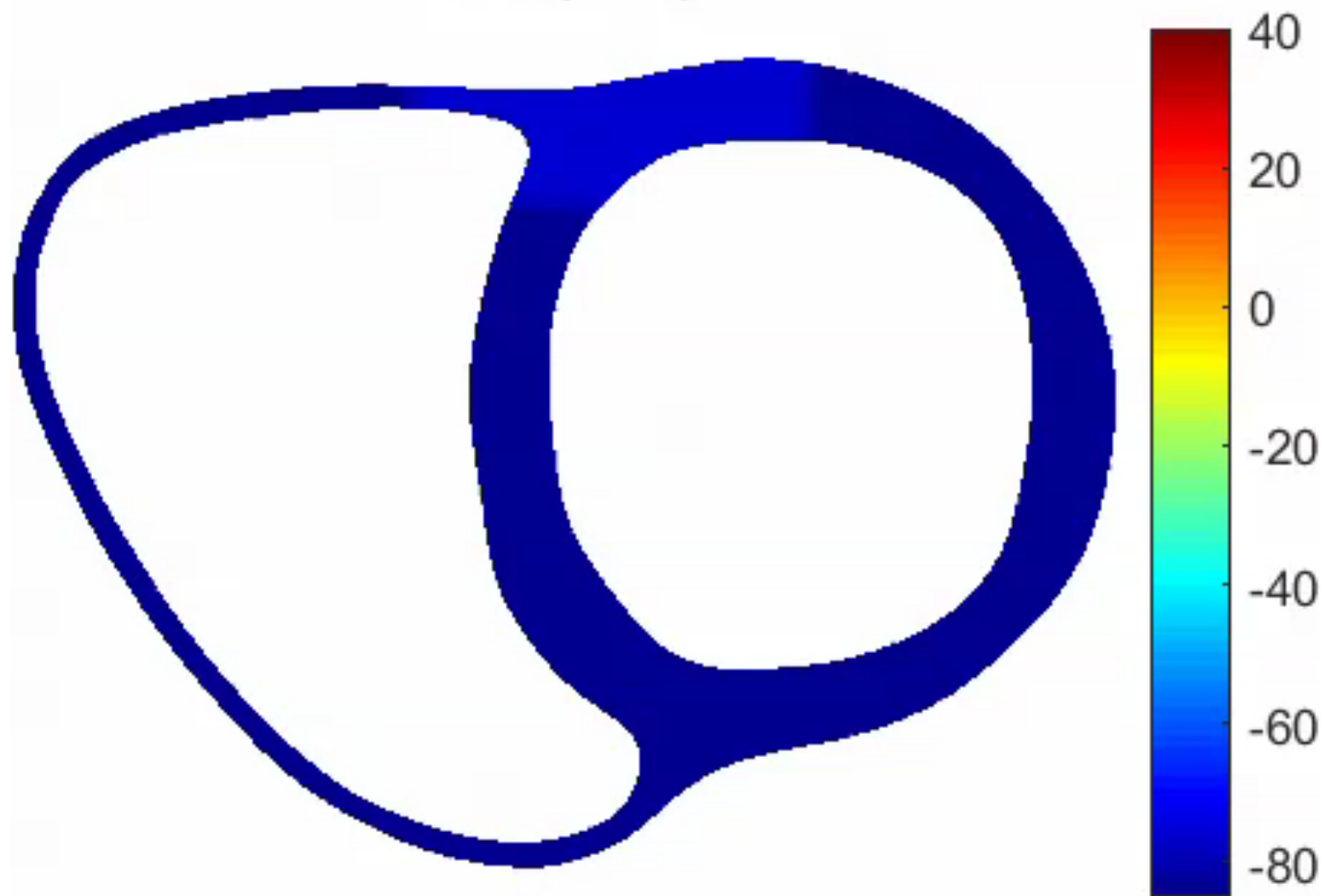
  // Modification of cell properties if it is in the ectopic region
  bool cell_is_in_stim_region = stim_region.DoesContain(this->GetMesh()->GetNode(pNode->GetIndex())->rGetLocation());

  if(cell_is_in_stim_region)
  {
    p_cell->SetStimulusFunction(mpStimulus2);
  }

  return p_cell;
}

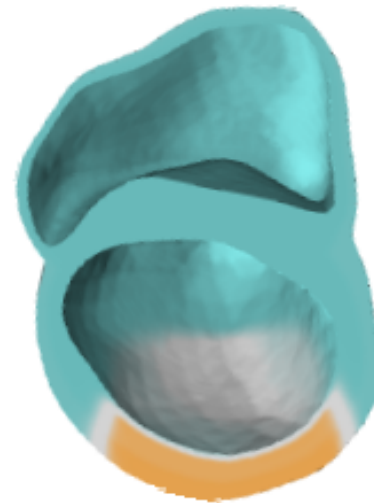
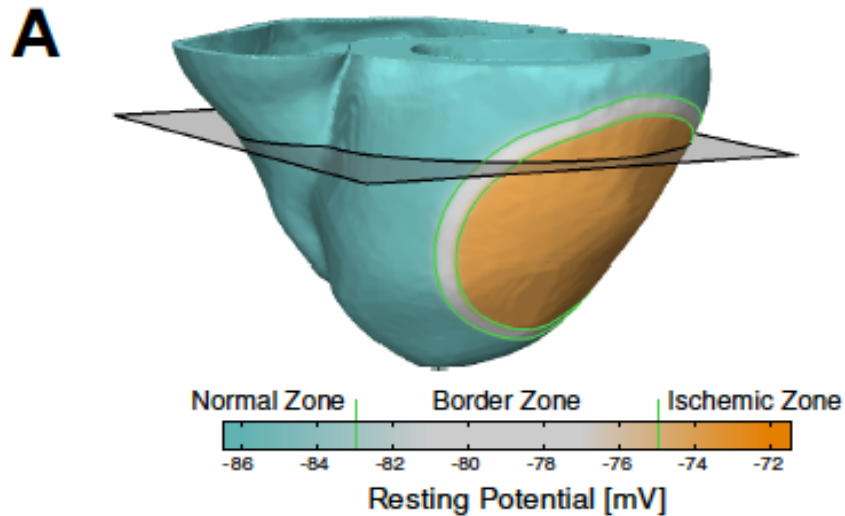
```

**Time = 0**

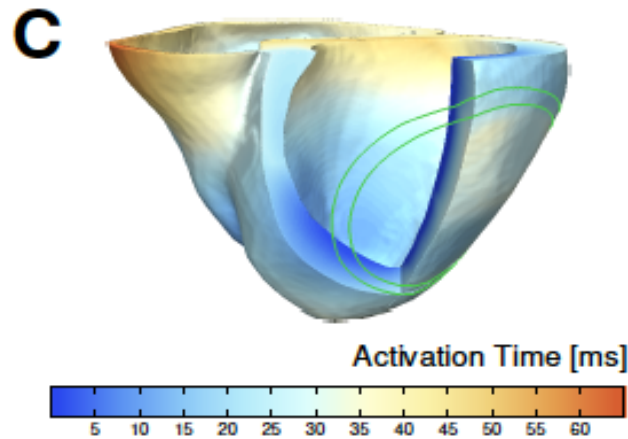
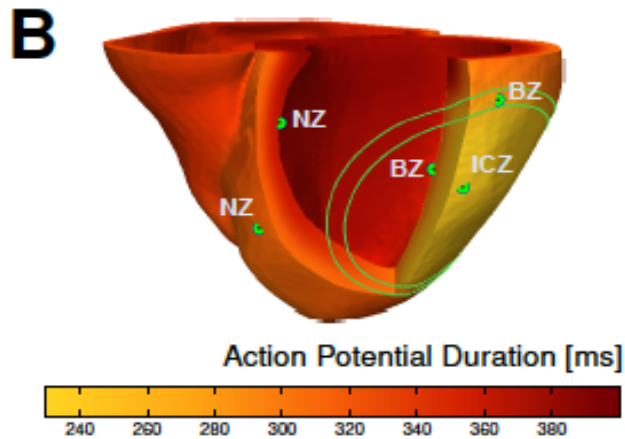




# 3D simulations in Chaste

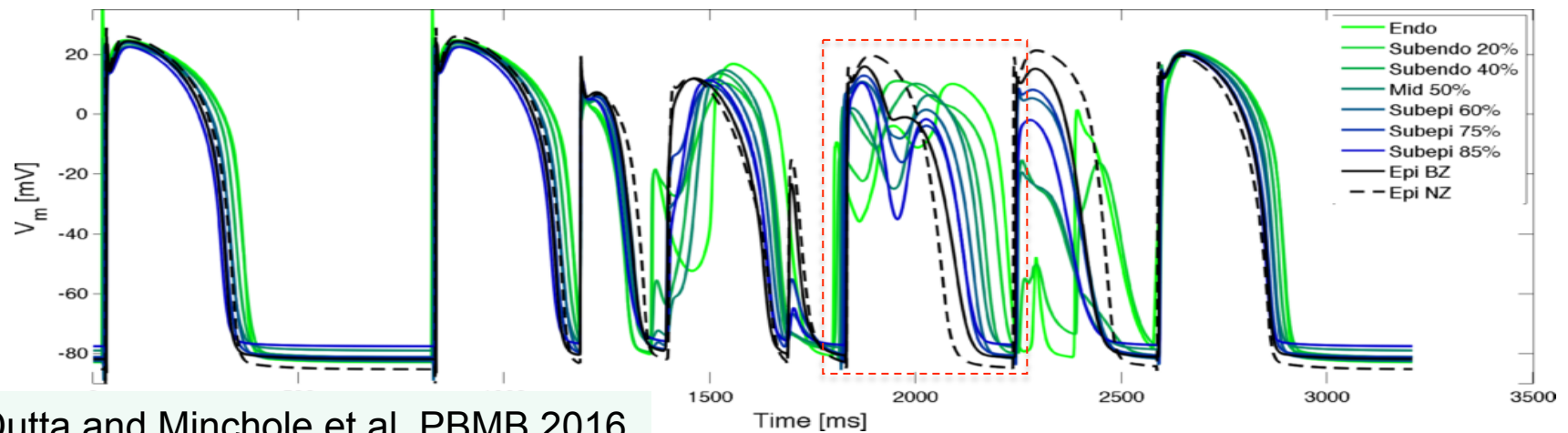
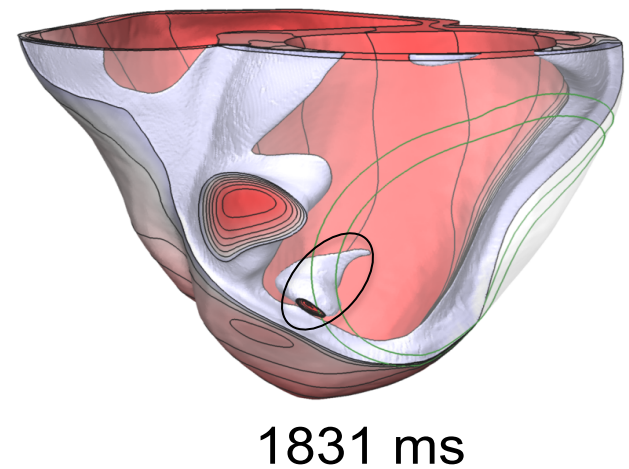
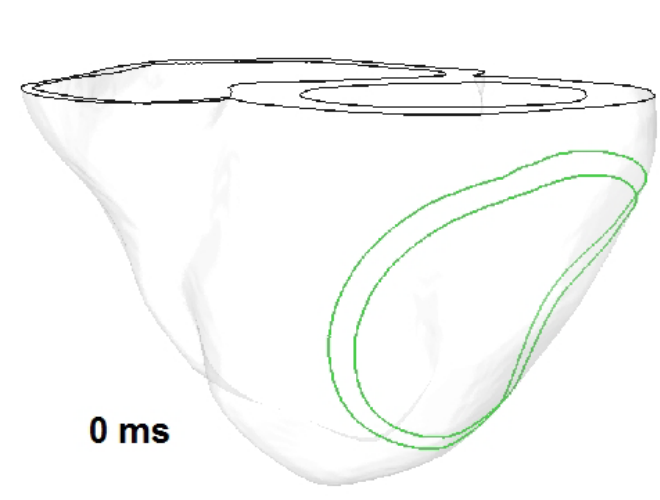


- Inclusion of border zones: endocardial and around the ischemic area
- Transmural heterogeneities



# 3D simulations in Chaste

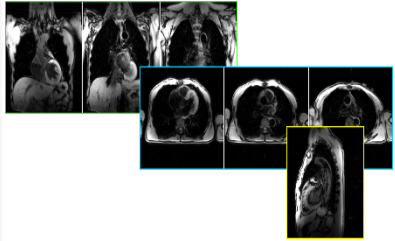
Simulated effect of IKr block in acute ischaemia



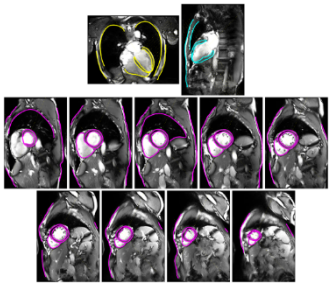
# Personalization of anatomical models

## MRI

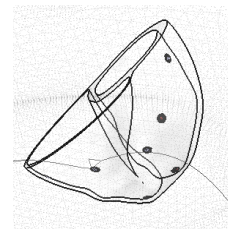
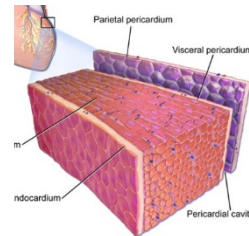
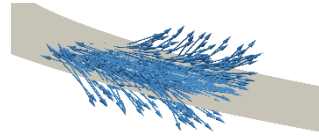
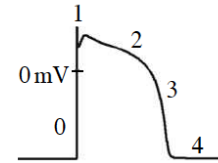
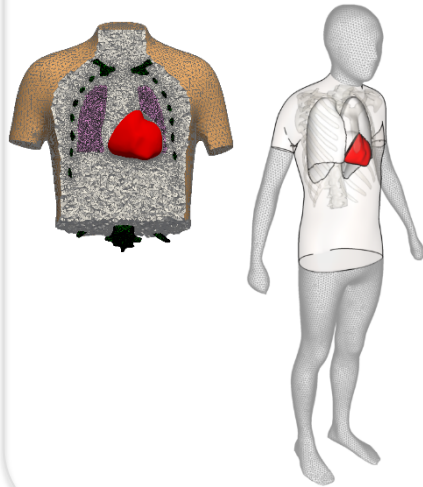
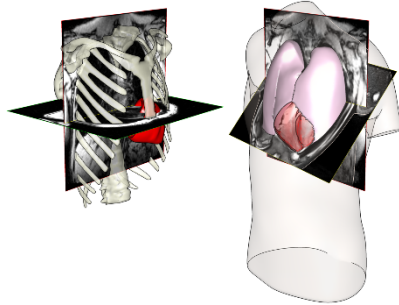
### LOCALIZERS



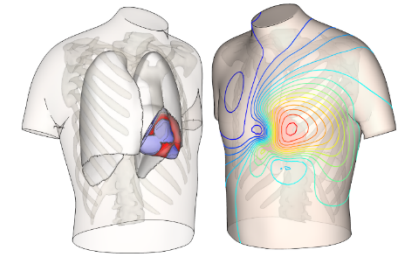
### CINE



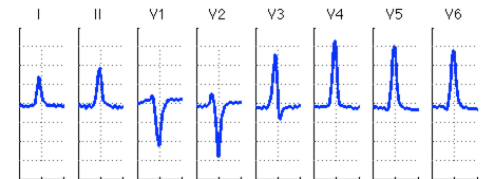
## Meshes Parametrization



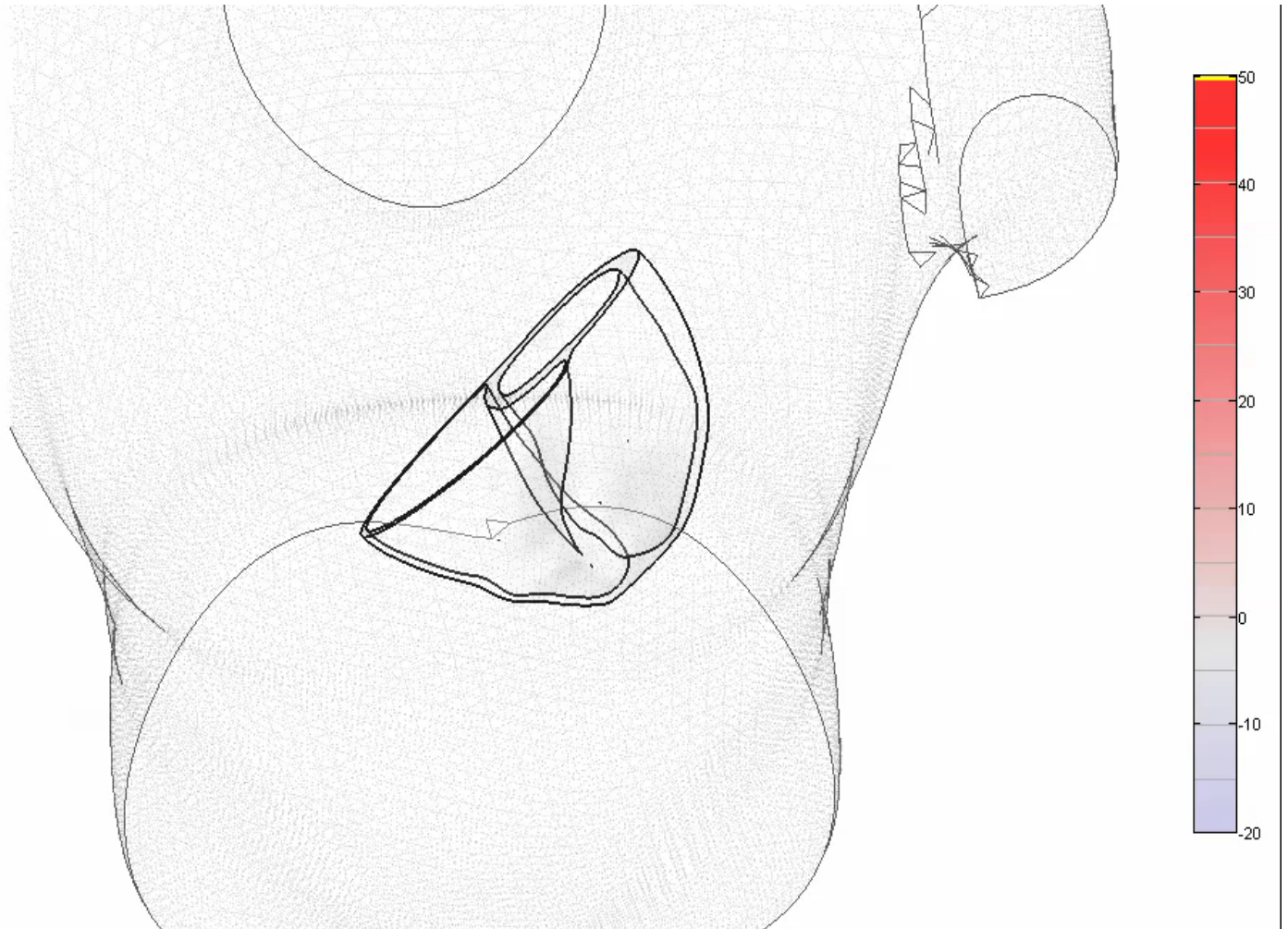
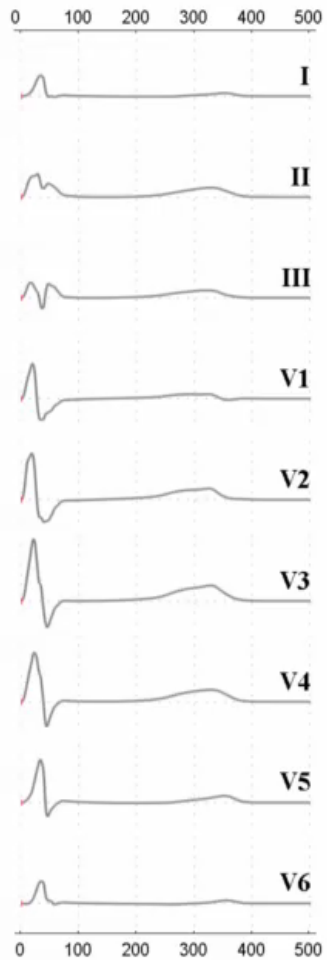
### ELECTRO-PHYSIOLOGY



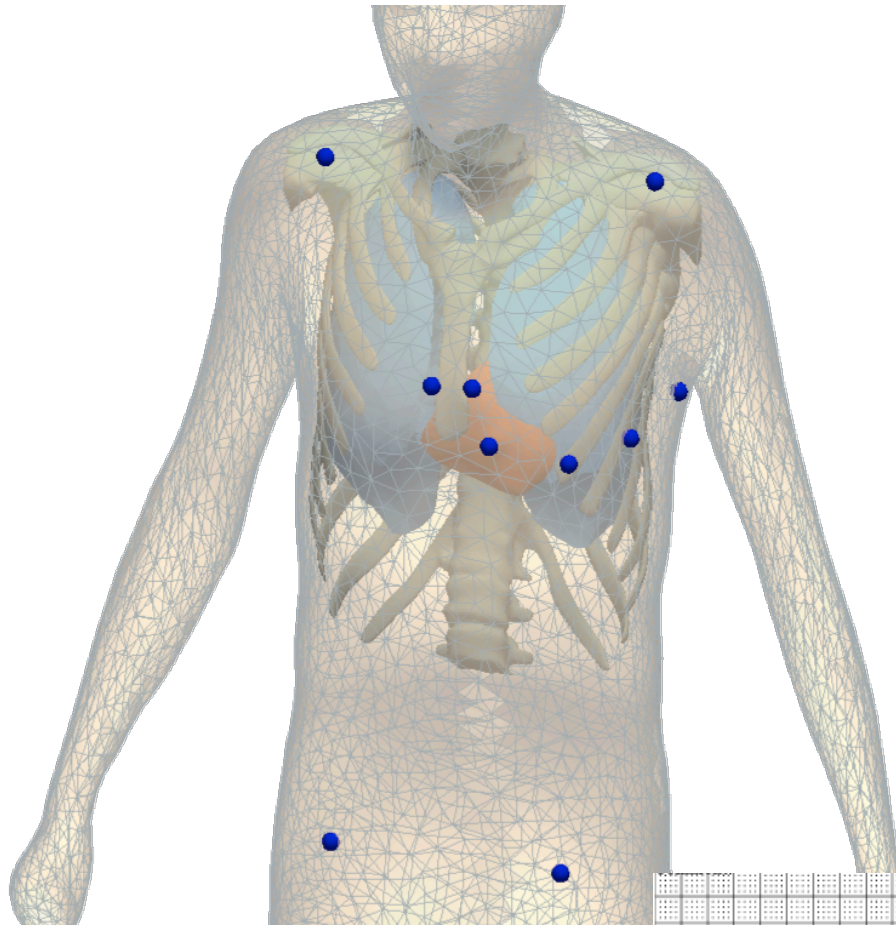
### ECG



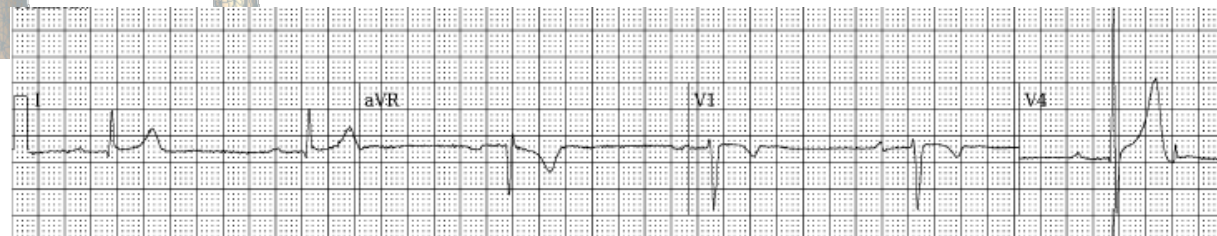
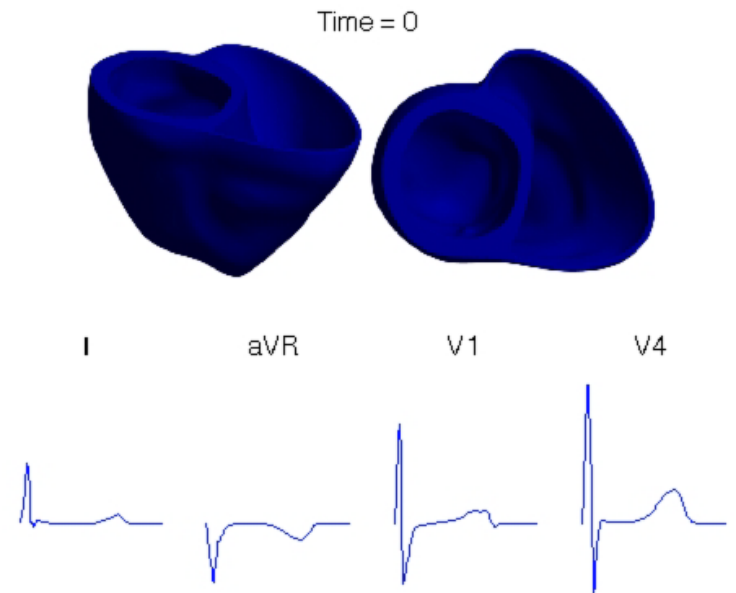
# Subject DTI003



# Computer Simulations to explain Cardiac phenotypes



Linking structure and function  
HPC cardiac simulation

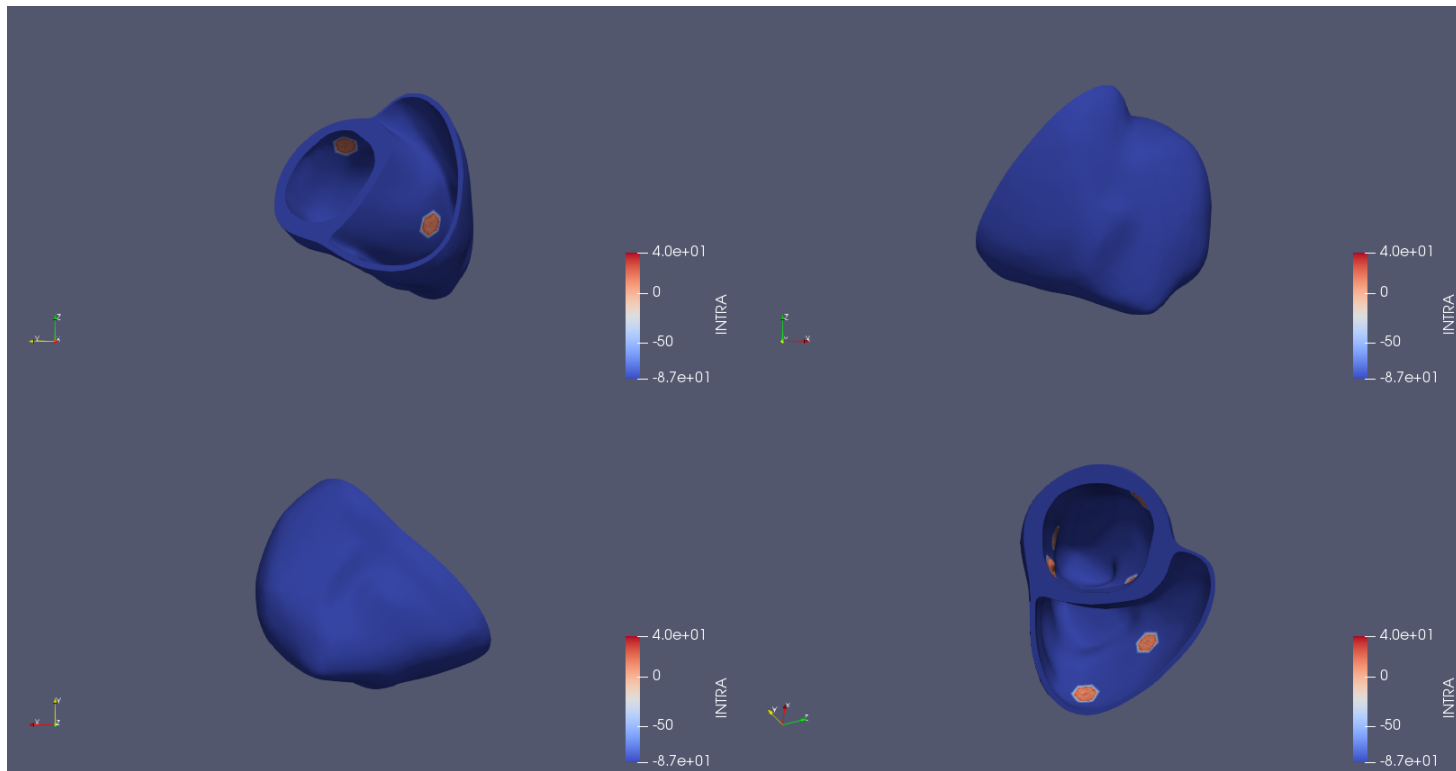


# Example of electrophysiology with Alya

- Alya functionalities:
  - ✓ Monodomain and bidomain models
  - ✓ A few cellular action potential models are implemented (FHN, ORd, TT)
  - ✓ Bidirectional electro-mechanical coupling is already implemented and fully functional
  - ✓ Excitation-contraction coupling models and models of stretch-activated ion channels are implemented
  - ✓ High efficiency and very high scalability
  - ✓ Other physics are also available (combustion, incompressible and compressible CFD...)

# Alya example 1

- Biventricular model (5.2M nodes and 32M elements)
- Run on MareNostrum IV (BSC) on 2,000 cores in 15 min
- This example comprises of 4 input files in ASCII format: .dat, .dom.dat, .ker.dat and .exm.dat
- Electrical propagation only





# Electro-mechanical modelling

## Equations of motion

$$\nabla \cdot \mathbf{P} + \rho_0 \mathbf{b} = \rho_0 \ddot{\mathbf{u}}$$

## Hyperelastic model

$$\psi(\mathbf{C}) = \frac{a}{2b} \exp[b(l_1 - 3)] + \sum_{i=f,s} \left\{ \exp[b_i(l_{4i} - 1)^2] - 1 \right\} + \frac{a_{fs}}{2b_{fs}} [\exp(b_{fs} l_{8fs}^2) - 1]$$

## Active contraction model

$$\frac{ds}{dt} = m_s(\mathbf{c}, \mathbf{s}, \lambda, \dot{\lambda})$$

## Active tension generation

$$T_{act} = h(\lambda) \frac{T_{ref}}{r_s} [(\zeta_s + 1)S + \zeta_w W]$$

## Mechano-electric coupling

$$\nabla \cdot (J\mathbf{F}^{-1} \mathbf{D}_0 \mathbf{F}^{-T} \nabla V) = \chi \left( C \frac{dV}{dt} + I_{ion} - I_{app} \right)$$

## Electro-mechanic coupling

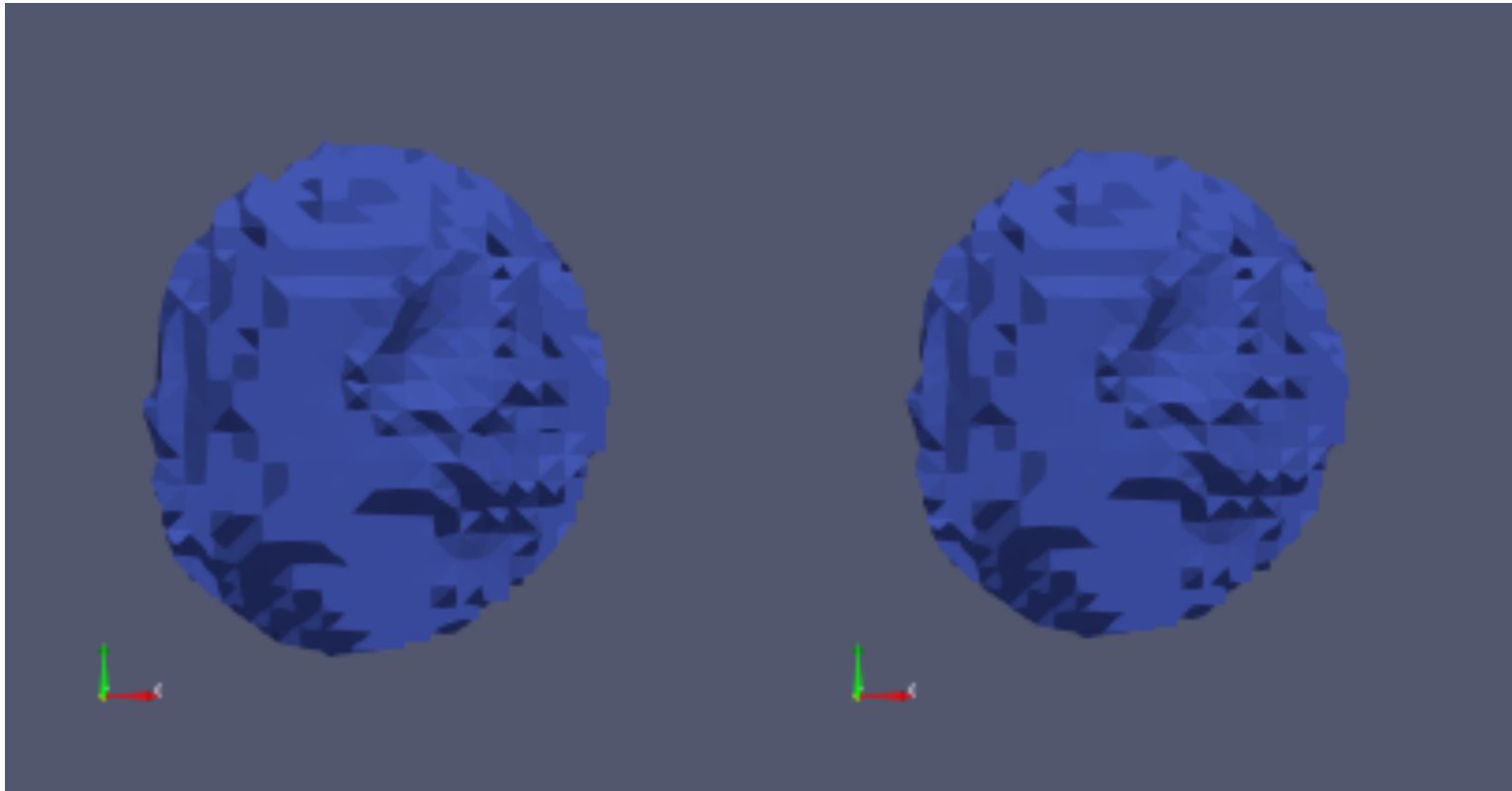
$$\frac{ds}{dt} = m_s(\mathbf{c}, \mathbf{s}, \lambda, \dot{\lambda})$$
$$T_{act} = h(\lambda) \frac{T_{ref}}{r_s} [(\zeta_s + 1)S + \zeta_w W]$$

## Stretch-activated ion channels

$$I_{inw}(V, \mathbf{F}) = f(\sqrt{\lambda} - 1)(V - E)$$

# Alya example 2

- ✓ Small blob of tissue
- ✓ Run locally with 4 cores
- ✓ Electromechanical simulation, Hunter(left) vs Land(right)



# Alya .dat and .dom.dat

```
$-----  
RUN DATA  
  ALYA: heart_test  
END_RUN_DATA  
$-----  
PROBLEM_DATA  
  MAXIMUM NUMBER GLOBAL: 1  
  NUMBER OF STEPS:      1000000  
  TIME INTERVAL:        0.000, 0.400  
$-----  
  EXMEDI:                ON  
  END_EXMEDI  
  $SOLIDZ PROBLEM:      ON  
  $END SOLIDZ  
  PARALL SERVICE:      ON  
  OUTPUT_FILE:         NO  
  POSTPROCESS:         MASTER  
  PARTITION:           FACES  
  END_PARALL  
$-----  
END_PROBLEM_DATA  
$-----
```

```
$-----  
DIMENSIONS  
  NODAL POINTS =      5196552  
  ELEMENTS =          31702122  
  SPACE_DIMENSIONS = 3  
  NODES =             4  
  BOUNDARIES =        920090  
END_DIMENSIONS  
$-----  
STRATEGY  
  INTEGRATION RULE =      OPEN  
  DOMAIN INTEGRATION_POINTS = 1  
END_STRATEGY  
$-----  
GEOMETRY, GID, WALL_DISTANCE = 0.0, ROUGHNESS = 0.0  
  FIELDS, NUMBER = 3  
  FIELD = 1, DIMENSION = 3, NODES  
    INCLUDE heart_test.fiber  
  END_FIELD  
  FIELD = 2, DIMENSION = 1, NODES  
    INCLUDE heart_test.celltype  
  END_FIELD  
  INCLUDE heart_test.stimuli  
END_FIELDS  
  INCLUDE heart_test.geo  
  INCLUDE heart_test.surfaces  
END_GEOMETRY  
$-----  
BOUNDARIES, EXTRAPOLATE  
$ INCLUDE heart_test.boundaries  
END_BOUNDARIES
```

# Alya .ker.dat and .exm.dat

```
$-----  
PHYSICAL PROBLEM  
$COUPLING  
$ SOLIDZ EXMEDI  
$END COUPLING  
END_PHYSICAL_PROBLEM  
$-----  
NUMERICAL_TREATMENT  
MESH  
$ DIVISION = 1  
END MESH  
END_NUMERICAL_TREATMENT  
$-----  
OUTPUT & POST PROCESS  
ON LAST MESH  
STEPS = 50  
END_OUTPUT & POST_PROCESS  
$-----
```

```
$-----  
PHYSICAL_PROBLEM  
$-----  
PROBLEM_DEFINITION  
STARTING POTENTIAL, FIELD  
CURDENSITY: FIELD = 3  
END STARTING POTENTIAL  
$GEO COUPLING: LAGRANGIAN  
END_PROBLEM_DEFINITION  
$-----  
PROPERTIES  
MATERIAL = 1  
CONTINUUM MODEL  
IN DIFFUSION = 0.00107143 0.00024107 0.00024107 [cm2/ms]  
INITIAL MEMBRANE POTENTIAL= -87.0 [mV]  
END CONTINUUM MODEL  
CELL MODEL: OHARA  
CM CONST = 1.0  
ENDCELL MODEL  
INI CELL, HETEROGENEOUS  
INCLUDE heart_test.framework  
END INI CELL  
END_PROPERTIES  
$-----  
FIBER MODEL: FIELD = 1  
CELL_TYPE: FIELD = 2  
$-----  
END_PHYSICAL_PROBLEM  
$-----  
NUMERICAL_TREATMENT  
$CONSTANT MATRIX  
SAFETY FACTOR = 25.0  
TIMETREATMENT: IMPLICIT, CRANK  
ALGEBRAIC_SOLVER  
SOLVER: CG  
CONVERGENCE: ITERA=1000, TOLER=1.0e-6, ADAPTIVE, RATIO=0.01  
OUTPUT: CONVERGENCE  
PRECONDITIONER: DIAGONAL  
END ALGEBRAIC_SOLVER  
END_NUMERICAL_TREATMENT  
$-----  
OUTPUT & POST_PROCESS  
POSTPROCESS_INTRA  
END_OUTPUT & POST_PROCESS  
$-----
```

# Final comments

- ▶ Complex nature of biology:
  - Models as tools to augment experimental/clinical findings.
- ▶ HPC is needed for whole biventricular and torso simulations:
  - Models can several tens of millions of elements!
- ▶ Highlight of the potential of mathematical modelling for *in silico* clinical and drug trials:
  - Substitution of current protocols for human *in silico* protocols
- ▶ Highlight of the potential of mathematical modelling for explaining the underlying mechanisms of abnormalities in the ECG:
  - Potential improvement of the associated inverse problem

# Acknowledgements

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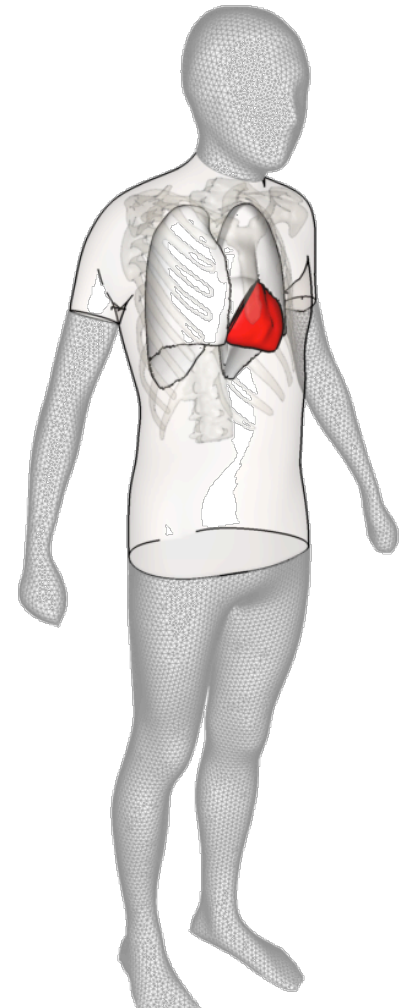
Rina Ariga, Hugh Watkins

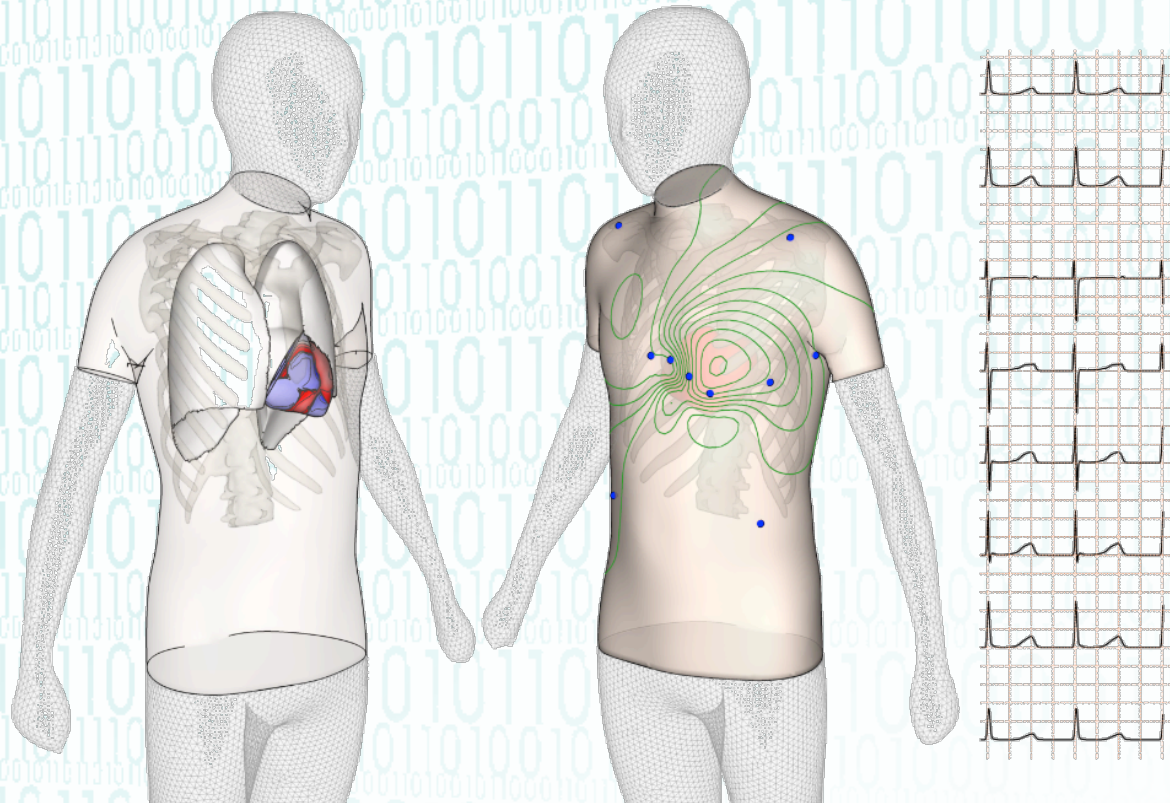
## Food and Drug Administration, USA

Sara Dutta, David Strauss

## Simula, Norway

Valentina Carapella





# Thank You!



National Centre  
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Refinement & Reduction  
of Animals in Research

**wellcome**trust



British Heart  
Foundation

