Coupled/Nonlinear PDEs

Coupled (linear) PDEs in Chaste

We consider solving a set of linear coupled PDEs, and assume it is a case in which the use of linear basis functions for all unknowns is appropriate (for example, a set of reaction-diffusion equations).

It is not possible to write a generic 'PDE class' for all such coupled systems, so a user wishing to solve such systems in Chaste will have to write their own solver.

However, using the tools available, this requires significantly less work than coding up from scratch



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Chaste

The parameter PROBLEM_DIM

Define PROBLEM_DIM to be the size of the system of PDEs. For example, for the PDE system $\,$

$$u_t = \nabla^2 u + v$$

$$v_t = \nabla^2 v + a \nabla^2 u$$

$$w_t = \nabla^2 w + u$$

we have PROBLEM_DIM equal to 3

The PDE solver classes are written to work with general PROBLEM_DIM. Ir particular the following classes are all templated over this:

BoundaryConditionsContainer<ELEM.DIM, SPACE.DIM, PROBLEM.DIM>
AbstractFeObjectAssembler<ELEM.DIM, SPACE.DIM, PROBLEM.DIM,
CAN_ASSEMBLE_VEC, CAN_ASSEMBLE_MAT>

AbstractLinearPdeSolver<ELEM_DIM, SPACE_DIM, PROBLEM_DIM>



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Striping

With the system

$$u_t = \nabla^2 u + v$$

$$v_t = \nabla^2 v + \alpha \nabla^2 u$$

$$w_t = \nabla^2 w + u$$

and with a mesh of N nodes, and linear basis functions for each unknown, the unknown vectors will be \mathbf{U}^n , \mathbf{V}^n , \mathbf{W}^n , each of size N.

In the linear system to be set up to solve this problem, the solution vector is chosen to be *striped*, i.e. the full solution vector is given by

$$\mathcal{U}^{n} = [U_{1}^{n}, V_{1}^{n}, W_{1}^{n}, U_{2}^{n}, V_{2}^{n}, W_{2}^{n}, \dots, U_{N}^{n}, V_{N}^{n}, W_{N}^{n}]$$

This is largely for parallelisation reasons

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Weak form

Consider the static problem:

$$\nabla^{2} u + \alpha \nabla^{2} v + f = 0$$

$$\nabla^{2} v + u + w = 0$$

$$\nabla^{2} w + \beta \nabla^{2} v = 0$$

subject to u = v = w = 0 on Γ_1 and natural boundary conditions on Γ_2 .

The linear system (in block form) can be read off to be

$$\begin{bmatrix} K & \alpha K & 0 \\ -M & K & -M \\ 0 & \beta K & K \end{bmatrix} \begin{bmatrix} \mathbf{U} \\ \mathbf{V} \\ \mathbf{W} \end{bmatrix} = \begin{bmatrix} \mathbf{b} \\ 0 \\ 0 \end{bmatrix}$$

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Coupled problems in Chaste

For such coupled linear PDEs, it is reasonably straightforward to set-up a (parallel, efficient, trustworthy) solver in Chaste.

The user needs to be able to convert their set of PDEs into a linear system as above, then only needs to implement functions <code>ComputeMatrixTerm()</code> and <code>ComputeVectorTerm()</code> saying what (the elemental contributions of) the matrix and vector are (remembering the striped nature of the data structures).

For examples, see tutorial on writing PDE solvers

Consider a nonlinear elliptic problem, such as

$$\nabla \cdot (D(\mathbf{u})\nabla \mathbf{u}) + f = 0$$

with boundary conditions

$$egin{array}{lll} & u & = & 0 & & \mbox{on } \Gamma_1 \\ & D(u) oldsymbol{
abla} u \cdot \mathbf{n} & = & g & & \mbox{on } \Gamma_2 \end{array}$$

Computing the weak form as before, we obtain: find $u \in \mathcal{V}_0$ satisfying

$$\int_{\Omega} (D(u)\nabla u) \cdot \nabla v \, \mathrm{d}V - \int_{\Omega} f v \, \mathrm{d}V - \int_{\Gamma_2} g v \, \mathrm{d}S = 0 \qquad \forall v \in \mathcal{V}_0$$

Write this as: find $u \in \mathcal{V}_0$ satisfying

$$\mathcal{F}(u,v)=0 \quad \forall v \in \mathcal{V}_0$$

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The finite element problem is obtained as before: find $u_h \in \mathcal{V}_0^h$ satisfying

$$\mathcal{F}(\mathbf{u}_h, \mathbf{v}) = 0 \qquad \forall \mathbf{v} \in \mathcal{V}_0^h$$

i.e. find coefficients U_1, \ldots, U_N of $u_h = \sum U_i \phi_i$ such that

$$\mathcal{F}(u_h, \phi_i) = 0$$
 for $i = 1, \dots, N$

This is a general N-dimensional nonlinear system

An iterative approach is required to solve nonlinear systems. Let u_h^k (equivalently, $\mathbf{U}^k = [U_1^k, \dots, U_N^k]$) be the current guess. Then the vector \mathbf{F}' defined by

$$F_i^k = \mathcal{F}(u_h^k, \phi_i)$$

is known as the k-th residual vector. We require a guess satisfying

$$\|\mathbf{F}^k\| < \text{TOL}$$

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Newton's method

Suppose we want to solve the nonlinear set of N equations

$$F(U) = 0$$

Given an initial guess U^0 , Newton's method is: let $\mathbf{U}^{k+1} = \mathbf{U}^k + \delta \mathbf{U}^{k+1}$, where $\delta \mathbf{U}^{k+1}$ satisfies the linear system

$$J(\mathbf{U}^k) \, \delta \mathbf{U}^{k+1} = -\mathbf{F}(\mathbf{U}^k)$$

where $J_{ij} = \frac{\partial F_i}{\partial U_j}$.

Newton's method provides quadratic convergence when the current guess is 'close enough' to the true solution. To avoid initial divergence however, it may be necessary to use damping

$$\mathbf{U}^{k+1} = \mathbf{U}^k + \mathbf{s}^k \delta \mathbf{U}^{k+1}$$

for some s^k generally smaller than 1. (There are various ways to go about choosing s^k , the simplest is to pick one from a small list of possibilities which leads to the smallest $||\mathbf{F}||$)

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- There are other methods for solving nonlinear systems, for example solve x = f(x) using **fixed point iterations**: $x^{n+1} = f(x^n)$.
- For F(U) = 0, this is

$$\boldsymbol{\mathsf{U}}^{n+1} = \boldsymbol{\mathsf{U}}^n + \boldsymbol{\mathsf{F}}(\boldsymbol{\mathsf{U}}^n)$$

- Very loosely speaking, methods which use the Jacobian will be more effective.
- If used, the Jacobian can be either provided analytically (if so, has to be calculated on paper on paper and coded up); or estimated numerically (slow).
- Petsc has (black-box) solvers for nonlinear systems. The user has to provide functions telling Petsc how to compute the residual (and optionally, the Jacobian)
- Chaste sometimes uses the Petsc nonlinear solvers (eg AbstractNonlinearAssemblerSolverHybrid), sometimes Newton's method is coded from scratch (solid mechanics solvers).

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Solving nonlinear problems with finite elements and Newton's method

- Choose a (good!) initial guess U⁰
- Compute the initial residual $F^0 = F(U^0)$ (loop over elements, compute elemental contribution, add to full vector).
- While $\|\mathbf{F}^k\| > \text{TOL}$

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Cardiac electro-physiology

The monodomain and bidomain equations

The monodomain equations (dropping stimulus currents) is essentially the heat equation coupled to ODEs:

$$\chi \left(C \frac{\partial V}{\partial t} + I_{\text{ion}}(\mathbf{u}, V) \right) - \nabla \cdot (\sigma \nabla V) = 0$$

$$\frac{\mathrm{d}\mathbf{u}}{\mathrm{d}t} = \mathbf{f}(\mathbf{u}, V)$$

(with zero-Neumann BCs on entire boundary)

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Placing cell models

We know that the forcing term of the heat equation enters the RHS vector of the FEM discretisation as (using ψ rather than ϕ for basis functions):

$$b_j = \int_{\Omega} f \psi_j \, \mathrm{d}V$$

which here is (assuming the reaction term is treated explicitly in the time-discretisation)

$$b_j^{n+1} = \int_{\Omega} \mathit{l}_{\mathsf{ion}}(\mathbf{u}^n, \mathit{V}^n) \psi_j \, \mathrm{d}\mathit{V}$$

Therefore we require the ionic current at the quadrature points, i.e. **u** is required at the quadrature points. The natural approach is therefore to solve cell models at quad points.

However this can be computationally-expensive (and a pain to implement), instead, solve cell models at nodes and interpolate onto quadrature points.

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Cell models at nodes

Solving cell models at nodes, we write the ionic current evaluated at the nodes as $\mathbf{I} = (I_1, \dots, I_N)$. Interpolating the ionic current onto the quadrature point using linear basis functions ψ_j , we have

$$I_{\mathsf{ion}} = \sum I_k \psi_k$$

which means that

$$b_j = \int_{\Omega} I_{\text{ion}} \psi_j \, dV = \int_{\Omega} \sum_k I_k \psi_k \psi_j \, dV = \sum_k I_k \int_{\Omega} \psi_k \psi_j \, dV = \sum_k M_{jk} I_k$$

so that

$$\mathbf{b} = M\mathbf{I}$$

Monodomain discretisation

Solve

$$\chi \left(C \frac{\partial V}{\partial t} + I_{\text{ion}}(\mathbf{u}, V) \right) - \nabla \cdot (\sigma \nabla V) = 0$$

$$\frac{d\mathbf{u}}{dt} = \mathbf{f}(\mathbf{u}, V)$$

subject to initial conditions and zero-Neumann boundary conditions.

We de-couple the ODEs from the PDEs, and use a time-discretisation which treats the conductivity implicitly and the (nonlinear) reaction term explicitly, and place cell models at nodes, obtaining, for the PDE solve:

$$\left(\frac{\chi C}{\Delta t} M + K\right) \mathbf{V}^{m+1} = \frac{\chi C}{\Delta t} M \mathbf{V}^m - M \mathbf{I}^m$$



Monodomain discretisation

Solve

$$\chi \left(C \frac{\partial V}{\partial t} + I_{\text{ion}}(\mathbf{u}, V) \right) - \nabla \cdot (\sigma \nabla V) = 0$$

$$\frac{d\mathbf{u}}{dt} = \mathbf{f}(\mathbf{u}, V)$$

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We de-couple the ODEs from the PDEs, and use a time-discretisation which treats the conductivity implicitly and the (nonlinear) reaction term explicitly, and place cell models at nodes, obtaining, for the PDE solve:

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Bidomain discretisation

Solve

$$\chi \left(C \frac{\partial V}{\partial t} + I_{\text{ion}}(\mathbf{u}, V) \right) - \nabla \cdot (\sigma_i \nabla (V + \phi_e)) = 0$$

$$\nabla \cdot (\sigma_i \nabla V + (\sigma_i + \sigma_e) \nabla \phi_e) = 0$$

$$\frac{d\mathbf{u}}{dt} = \mathbf{f}(\mathbf{u}, V)$$

subject to initial conditions and zero-Neumann boundary conditions.

$$\begin{bmatrix} \frac{\chi C}{\Delta t} M + K[\sigma_i] & K[\sigma_i] \end{bmatrix}$$

$$\begin{bmatrix} \frac{\chi \mathcal{C}}{\Delta t} M + K[\sigma_i] & K[\sigma_i] & K[\sigma_i + \sigma_e] \end{bmatrix} \begin{bmatrix} \mathbf{V}^{m+1} \\ \mathbf{\Phi}_e^{m+1} \end{bmatrix} = \begin{bmatrix} \frac{\chi \mathcal{C}}{\Delta t} M \mathbf{V}^m - M \mathbf{I}^m \\ 0 \end{bmatrix}$$

$$K[\sigma]_{jk} = \int_{\Omega} \nabla \psi_k \cdot (\sigma \nabla \psi_j) \, \mathrm{d}V$$



Bidomain discretisation

Solve

$$\chi\left(\mathcal{C}\frac{\partial V}{\partial t} + I_{\text{ion}}(\mathbf{u}, V)\right) - \nabla \cdot (\sigma_i \nabla (V + \phi_e)) = 0$$

$$\nabla \cdot (\sigma_i \nabla V + (\sigma_i + \sigma_e) \nabla \phi_e) = 0$$

$$\frac{\mathrm{d}\mathbf{u}}{\mathrm{d}t} = \mathbf{f}(\mathbf{u}, V)$$

subject to initial conditions and zero-Neumann boundary conditions.

Going through the same procedure, we obtain

$$\begin{bmatrix} \frac{\chi \mathcal{C}}{\Delta t} M + K[\sigma_i] & K[\sigma_i] \\ K[\sigma_i] & K[\sigma_i + \sigma_e] \end{bmatrix} \begin{bmatrix} \mathbf{V}^{m+1} \\ \mathbf{\Phi}_e^{m+1} \end{bmatrix} = \begin{bmatrix} \frac{\chi \mathcal{C}}{\Delta t} M \mathbf{V}^m - M \mathbf{I}^m \\ 0 \end{bmatrix}$$

where

$$K[\sigma]_{jk} = \int_{\Omega} \nabla \psi_k \cdot (\sigma \nabla \psi_j) \, \mathrm{d}V$$



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Solution overview

For either the monodomain or bidomain equations

- Set up the left-hand side matrix, A say (loop over elements, etc)
- Set up the mass matrix (loop over elements, etc)
- \bullet Set up the initial conditions V^0 , also initialise cell models at each node
- While $t < t_{\rm end}$
 - Pass nodal voltages to each cell model
 - · Solve cell models at each node using choice of ODE solver
 - Compute ionic current at each node
 - Set up linear system RHS vector (matrix-vector products only, no need for assembly)
 - Solve linear system



Cell models in Chaste

```
AbstractOdeSystem
    mStateVariables
    EvaluateYDerivatives(t,y)
```

```
AbstractCardiacCell<sup>1</sup> inherits from AbstractOdeSystem
mOdeSolver

▷ of type AbstractOdeSolver

Compute(40.±1)
```

□ Use solver to solve between given times, updating internal state

Class has various other cardiac cell related functionalit

LuoRudyCellModel inherits from AbstractCardiacCell



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Cell models in Chaste

```
AbstractOdeSystem
    mStateVariables
    EvaluateYDerivatives(t,y)
```

> of type AbstractOdeSolver

Compute(t0,t1)

- ▷ Class has various other cardiac cell related functionality

LucRudyCellModel inherits from AbstractCardiacCell:



¹Slightly simplified

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Cell models in Chaste

```
AbstractOdeSystem
    mStateVariables
    EvaluateYDerivatives(t,y)
```

▷ of type AbstractOdeSolver

Compute(t0,t1)

- ▷ Class has various other cardiac cell related functionality





¹Slightly simplified

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Cardiac PDE solvers in Chaste

AbstractLinearPdeSolver:

```
PrepareForSetupLinearSystem()
```

> Empty implementation here (ie does nothing) SetupLinearSystem()

AbstractDynamicPdeSolver inherits from AbstractLinearPdeSolver :

SetTimes(t0,t1)

SetInitialCondition(initialCondition)

Solve()

▷ Calls PrepareForSetupLinearSystem(), then calls SetupLinearSystem(), then solves linear system.



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Cardiac PDE solvers in Chaste

MonodomainSolver

mMonodomainTissue

- ▷ Basically a set of AbstractCardiacCells for each node
- ▷ plus conductivity information

 ${\tt mMonodomainAssembler}$

mMassMatrixAssembler

PrepareForSetupLinearSystem()

> Overloaded to solve all the cell models

Implemented method: SetUpLinearSystem()

> Uses the above assemblers to set up the linear system

Motos

- BidomainSolver uses the same design (but uses PROBLEM_DIM=2
- There is MonodomainProblem and BidomainProblem (both inheriting from AbstractCardiacProblem). These own solvers and deal with set-up and output etc.



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