



An HPC Implementation of the Finite Element Method

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Turning Discovery Into Health

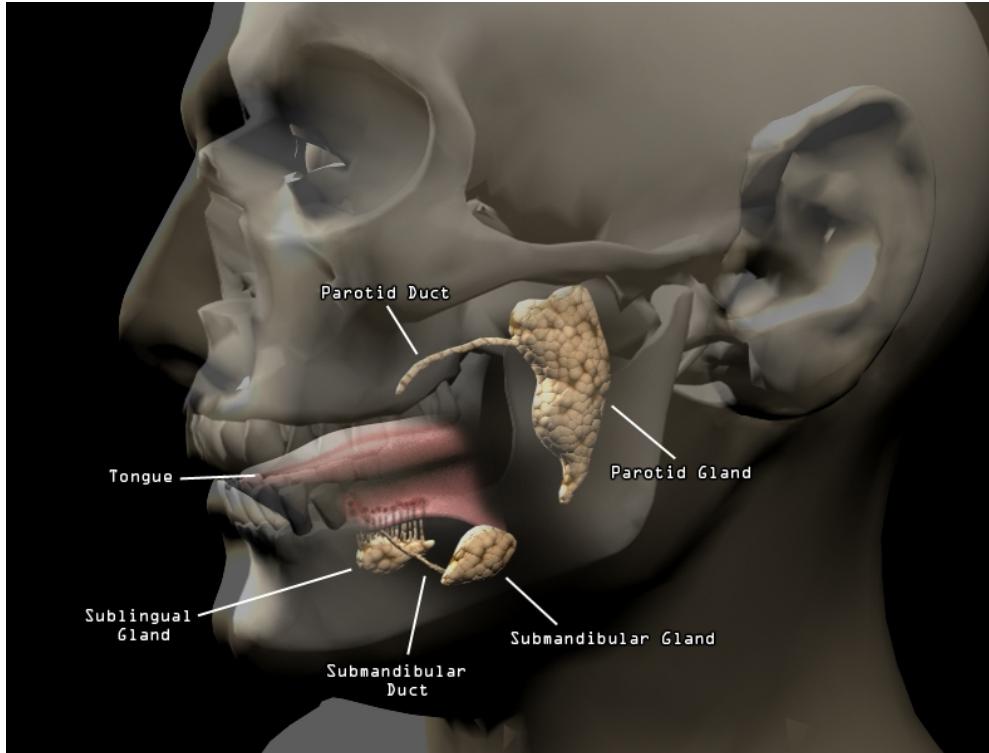
The science goal:

Physically accurate modeling, simulation and visualisation
of biological cell function.

Scalable!

- High Performance Computing
- *An Object Oriented Approach to Biological Cell Modeling*

Salivary glands

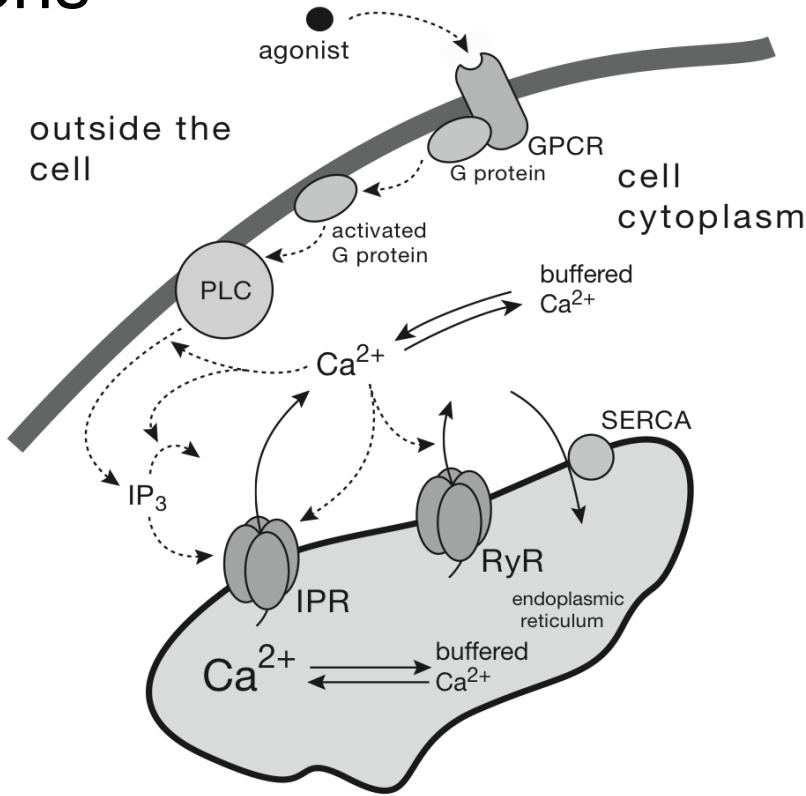


A Finite Element Method workflow overview

- Modeling equations (*How does the system work?*)
- Mesh generation (*What does the system look like?*)
- Discretisation (*How can this be implemented on a computer?*)
- Run simulation (*What does the system do?*)
- Evaluate results (*What exactly happened?*)

The Modeling Equations

Start with a conceptual model...



The Modeling Equations

Calcium and IP₃ dynamics

Partial differential equations model the cell calcium dynamics.

Reaction-Diffusion:

$$\frac{\partial c}{\partial t} = D_c \nabla^2 c + (J_{\text{IPR}} + J_{\text{leak}})(c_e - c) - J_{\text{serca}}$$

$$\frac{\partial p}{\partial t} = D_p \nabla^2 p + V_{\text{PLC}}(\vec{x}) - V_{\text{deg}} \left(\frac{c^2}{K_{3K}^2 + c^2} \right) p$$

$$\frac{\partial h}{\partial t} = \frac{h_\infty - h}{\tau}$$

$$J_{\text{serca}} = V_s \frac{c^2}{K_s^2 + c^2}$$

$$J_{\text{IPR}} = k_{\text{IPR}}(\vec{x}) P_O$$

$$P_O = \phi_c \phi_p h$$

$$\phi_c = \frac{c^3}{K_a^3 + c^3}$$

$$\phi_p = \frac{p^4}{K_p^4 + p^4}$$

$$h_\infty = \frac{K_i^2}{K_i^2 + c^2}$$

$$c_e = (c_t - c)/\gamma$$

Mesh Generation - Data Acquisition

Confocal microscopy

2D image stacks

32 slices in Z direction.

Cells & Lumen

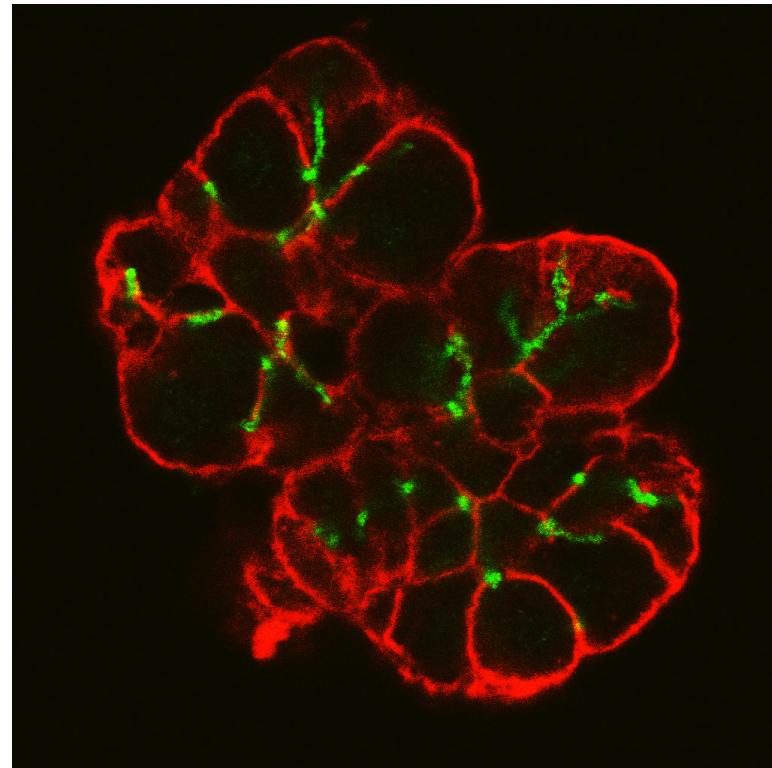
red: Na-KATPase

green: Cl Channel

Real dimensions used:

$70.7\mu\text{m}^2$, $2.2\mu\text{m}$ spacing

TIFF files: 1024x1024

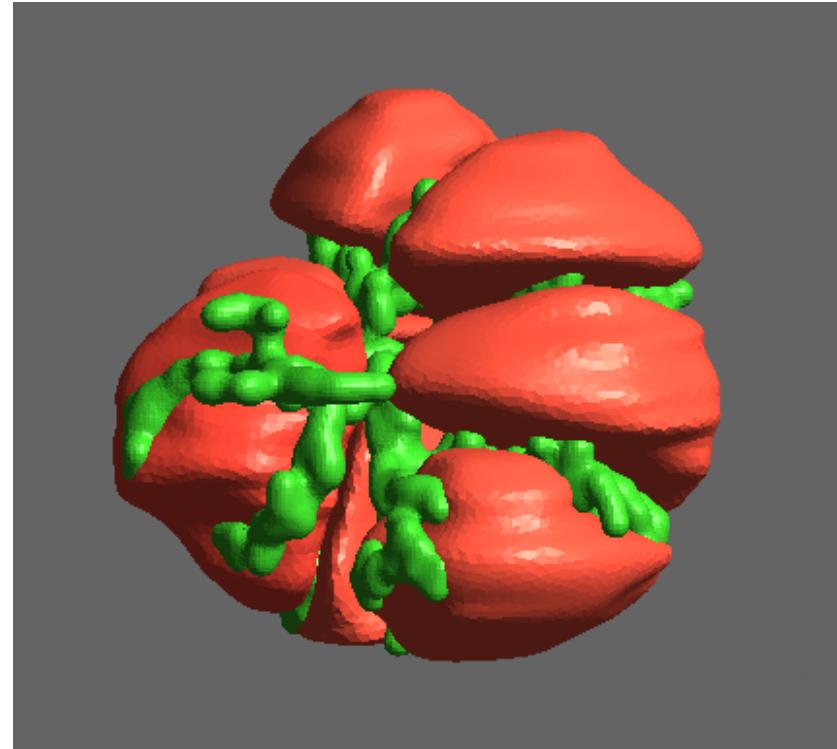


Mesh Generation - 3D Physical Model

Cells & Lumen

The cells are grouped in tight clusters.

The lumen has a tree-like branching structure.

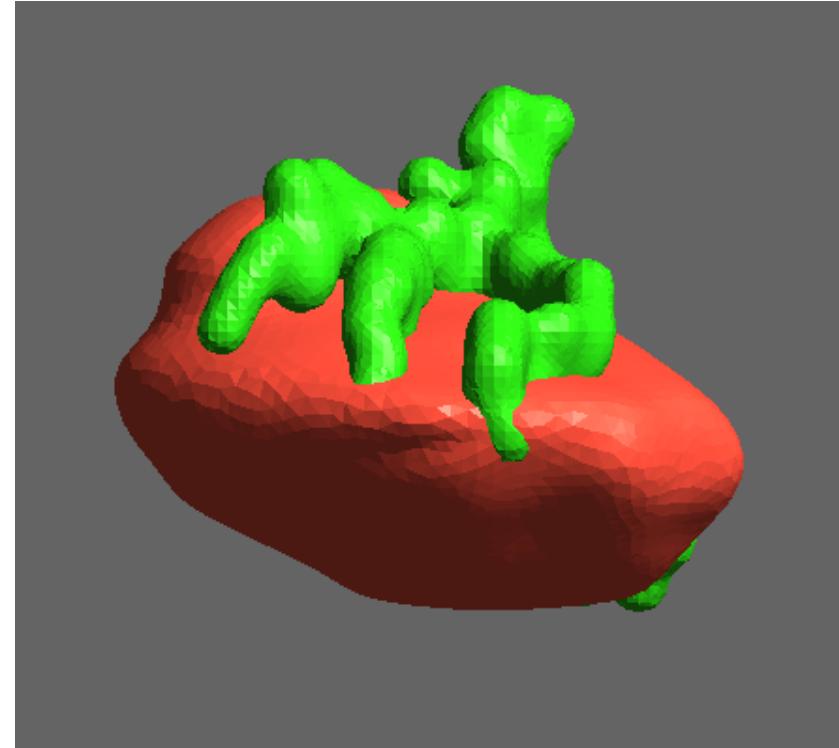


Mesh Generation - 3D Physical Model

Cells & Lumen

Each cell is held by a lumen “claw”.

The lumen has a central trunk.



Mesh Generation - for FEM

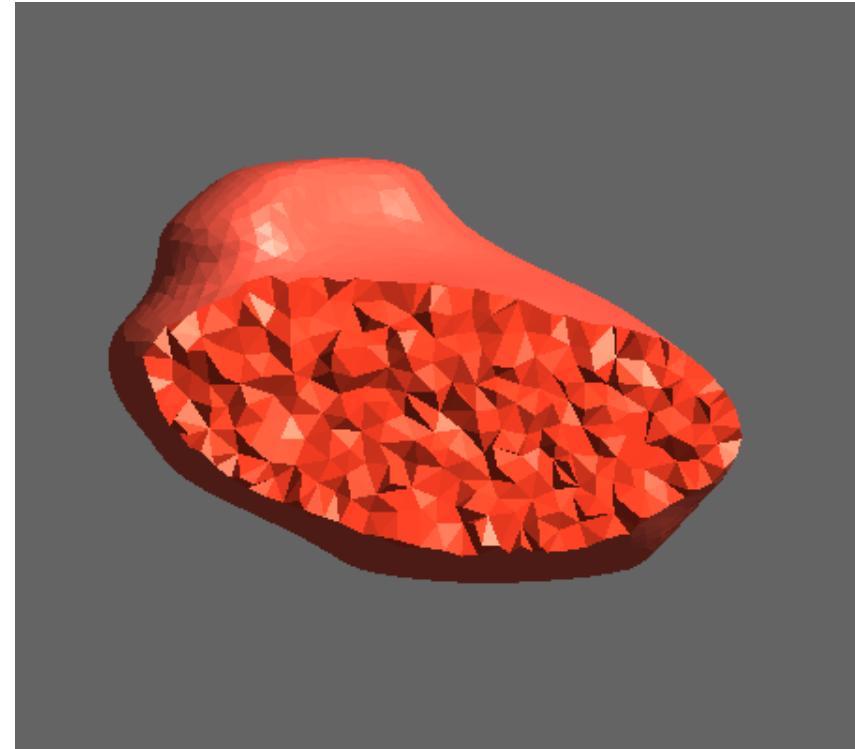
Cells

Solid volumetric meshing with tetrahedrons.

Tetrahedrons have vertices and edges.

Element = tetrahedron

Node = vertex



Discretisation

Facilitates (almost direct) translation to computer code.

- Build “mass” and “load” matrices.
- Construct (sparse) system matrix: ~ 400,000,000 elements!

NOTE: Most of the simulation run-time is spent solving the sparse system matrix.

Computer Code

Tool selection

Sparse matrix solver: *PETSc*

Portable, Extensible Toolkit for Scientific Computation

<http://www.mcs.anl.gov/petsc>

Vector and matrix library: *Eigen*

Eigen is a C++ template library for linear algebra

<http://eigen.tuxfamily.org>

Computer Code

Object oriented!

Language: C++

Once the behaviour of a cell “object” is defined, we can instantiate clusters of cells that interact with each other and their environment.

Computer Code

Object definition

```
class cGeneric3dModel {
public:
    cGeneric3dModel(cCellMesh *mesh, cPetscSolver *solver);
    virtual ~cGeneric3dModel();
    void run();
    void save_results();

    MatrixXXC Amat, mass, u; // A, mass and solution matrices

private:
    void get_parameters();
    void init_u();
    MatrixX1C make_load(long i);
    ArrayRefMass make_ref_mass();
    Array1VC getbodyreactions(tCalcs c, tCalcs ip, tCalcs h, tCalcs ipr_f, tCalcs plc_f);
    tCalcs getboundaryflux(tCalcs c);
    void make_matrices();
    void load_node_data(std::string file_name, int dindex);
    void save_matrix(std::string file_name, MatrixXXC mat);

    cCellMesh *mesh;
    cPetscSolver *solver;
    tCalcs p[PCOUNT]; // the model parameters array
    long numt; // number of time steps
    Eigen::Array<tCalcs, Eigen::Dynamic, MODELNCOUNT> node_data;
    Eigen::Array<tCalcs, Eigen::Dynamic, MODELECOUNT> element_data;
};
```

Computer Code

HPC – *How can we scale up?*

- One cell object per compute node
- Scale “out” (i.e. use more compute nodes)
 - MPI for cell interactions (light weight!)
- Scale “in” (i.e. use more cores per compute node)
 - Threads for utility functions
 - Accelerators for the main solver (i.e. GPU’s, Intel Phi)

Simulation

HPC - *Auckland Pan Cluster*

- Job submission scripting (shell scripting and python)
- Multiple parameter sweeps
- File and directory structures facilitate reproducibility!

Simulation

Job submission script (extract)

```
# create the top level results directory
csdir = os.getcwd()
path = "results/CS" + time.strftime("%y%m%d%H%M%S")
os.mkdir(path)
os.chdir(path)

# iterate through the parameters
for v1 in valsA:
    for v2 in valsB:
        pdir = ""
        for pv in parms:
            pdir += pv[0] + '-' + str(pv[1]) + '_'
        pdir += parmA + '-' + str(v1) + '_' + parmB + '-' + str(v2)
        pdir = pdir.replace('.', 'p')
        os.mkdir(pdir)
        os.chdir(pdir)

        # copy data files and execute the simulation
        os.system("cp " + csdir + "/parameters/" + model + ".dat cs.dat")
        if(os.path.isfile("ipr_REF.bin") and os.path.isfile("plc_REF.bin")):
            os.system("cp " + csdir + "/parameters/ipr_REF.bin .")
            os.system("cp " + csdir + "/parameters/plc_REF.bin .")

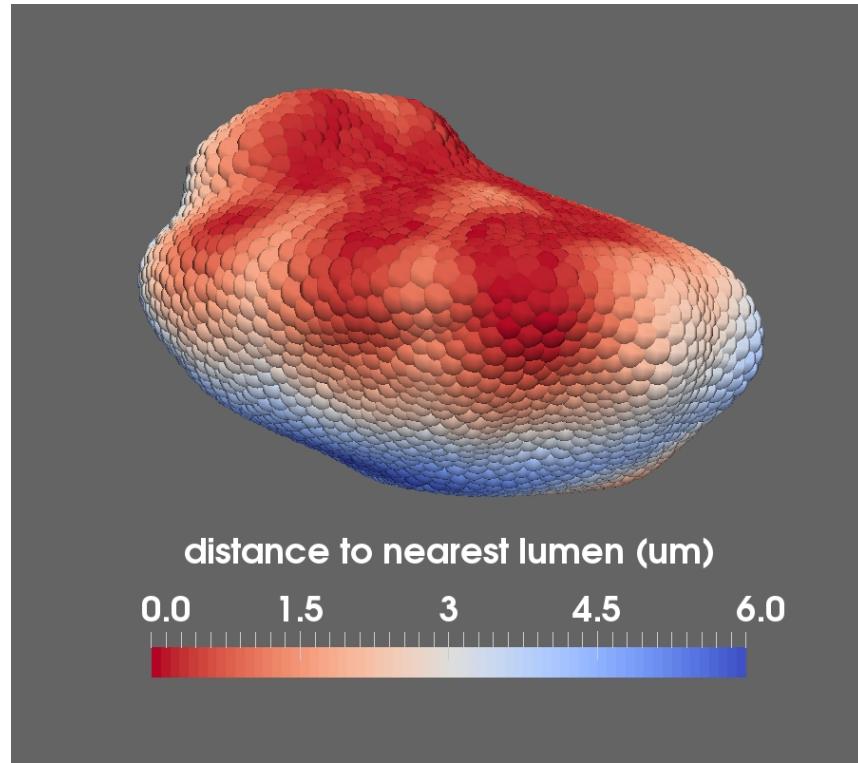
        for pv in parms: # set the fixed values
            replace_pvalue(pv[0], pv[1])
        replace_pvalue(parmA, v1) # set the swept values
        replace_pvalue(parmB, v2)
        os.system("cp " + csdir + "/meshes/" + mesh + ".msh cs.msh")
        os.system("sbatch " + csdir + "/run_sim.sl " + model + " " + mesh + " " + csdir)
        os.chdir("../")
```

Evaluate Results

Cell
Nodal view

Precomputed values.

Imprint of the lumenal
“claw”.



Evaluate Results

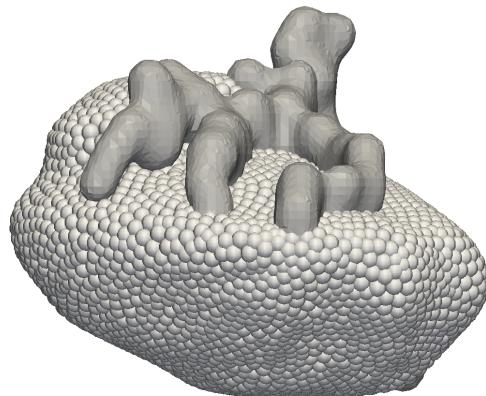
Cell
Nodal view

Static distributions.

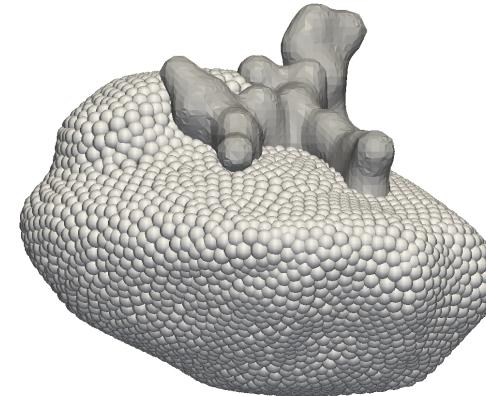
IPR in red.

PLC in blue.

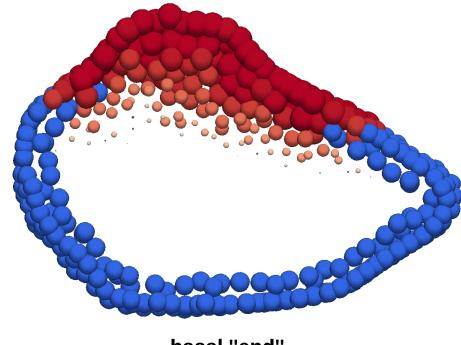
100% lumen



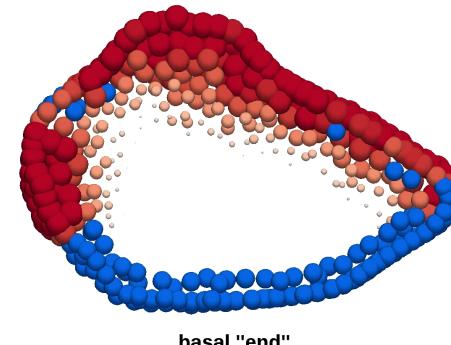
60% lumen



apical "end"



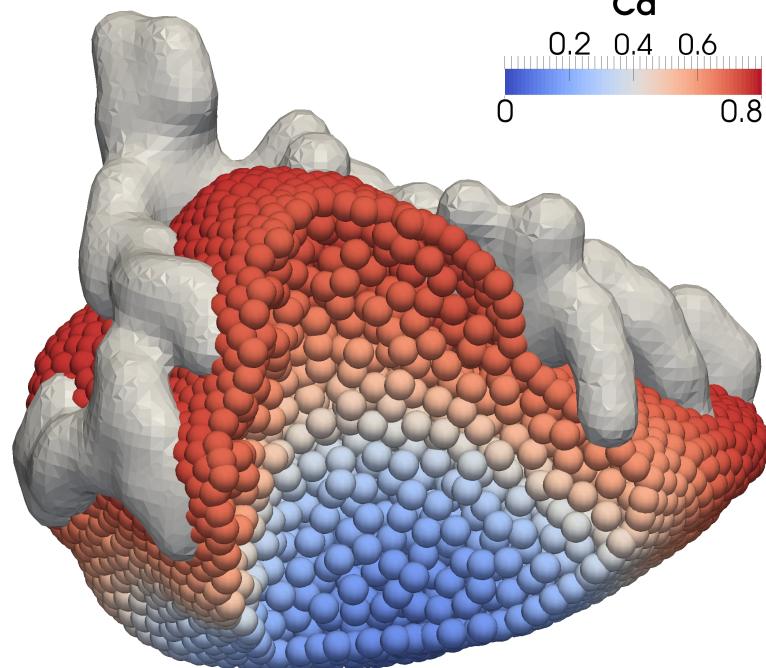
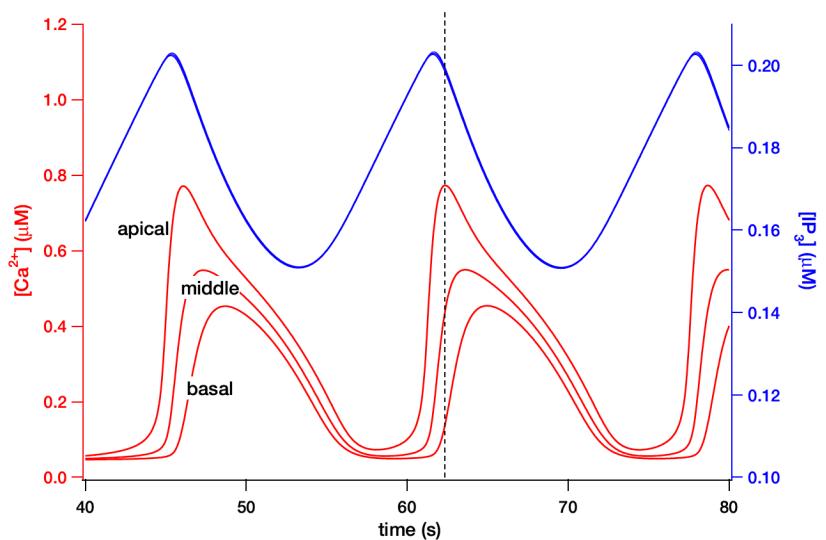
apical "end"



FEM Simulation Results

Calcium waves

Wave-fronts propagate from the apical end to the basal end the cell.



What's next?

- Run simulations with coupled cells (i.e. gap junctions)
- Include fluid flow in lumen (computational fluid dynamics)
- Higher resolution digitisations
- Stay tuned, more to come...

Questions & Answers?

Tuesday 11:15 (Queenstown)	NeSI Update	Nick Jones
Tuesday 11:45 (Remarkables)	Growing NZ's Researcher's Computing Capability	Georgina Rae & John Rugis
Tuesday 14:30 (Queenstown)	An HPC Implementation of the Finite Element Method	John Rugis
Tuesday 16:00 (Queenstown)	The NeSI National Platform Framework	Michael Uddstrom
Tuesday 16:30 (Queenstown)	Early Experiences with Cloud Bursting	Jordi Blasco
Tuesday 16:30 (Remarkables)	NeSI – NZGL Alliance	Dan Sun & Nic Mair
Tuesday 17:00 (Queenstown)	Acceleration Made Easy	Wolfgang Hayek

