

# Computational Physiology DSLs - LibCellML and related ideas

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## Background

The ABI Software Laboratory contributes to a number of open source software projects that are focussed on the computational modelling needs of bioengineering, computational physiology, and related fields.

- **Model representation** – CellML, FieldML<sup>[1]</sup>
- **Repository**
  - Models – PMR
  - Data – PMR, CAP
- **Simulation** – OpenCMISS-Iron
- **Visualisation** – OpenCMISS-Zinc, Cmgui
- **Workflow**: MAP Client
- **Tools**: CAP Client, OpenCOR

## Challenges

Multiphysics, multiscale, population variation, agent based modelling.

Computational demands

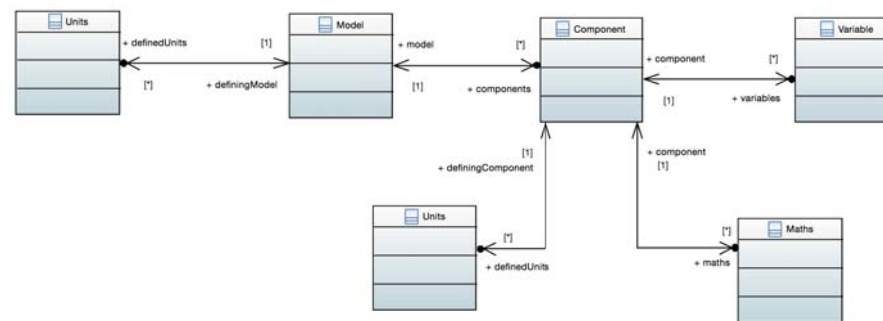
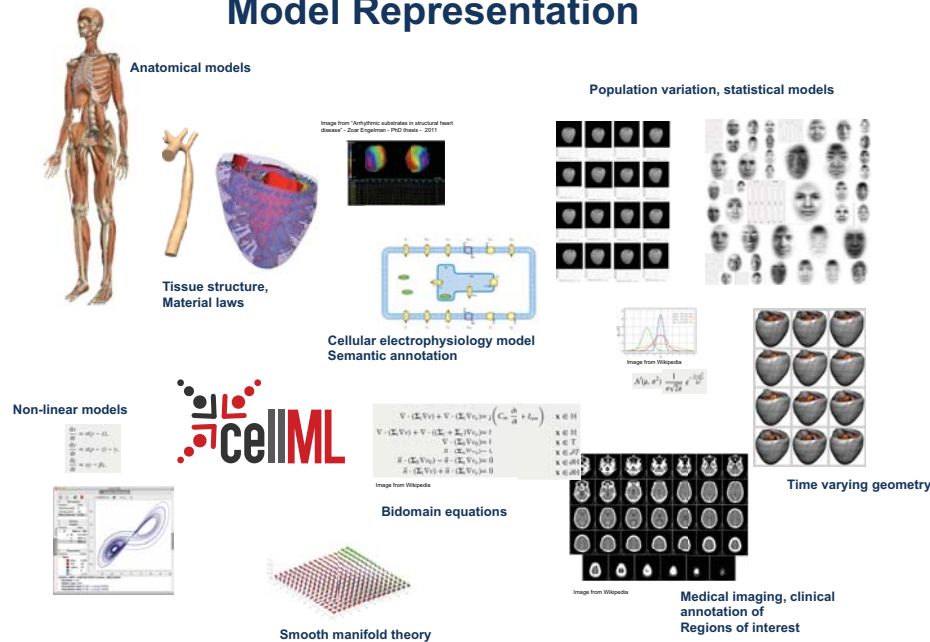
Research reproducibility

Model repurposing

## Approaches

- **System Modularity** - separation of concerns:
  - Solver vs. model: Domain Specific Language (DSL) for model, e.g. CellML, FieldML, SBML
  - Modularity within model – e.g. separate CellML components.

## Model Representation



LibCellML Domain Specific Object model

## Related work

### CellML API

- Library for processing CellML v1.0 and v1.1 models.
- Used by: OpenCOR, OpenCell, PMR, OpenCMISS-Iron, Antimony

### FieldML

- Similar goals, but focussed on time varying spatial models (geometry and advanced fields)

### Prototypes

- Haskell: DSLs, EDSLs (embedded DSLs) and model processing.

## Recent work

**LibCellML**: Library for processing CellML models.

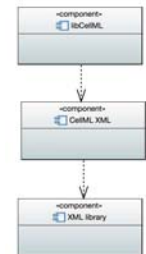
Just started.

Planned support for CellML version 1.2 (still WIP)

C++11 codebase

Open source, on GitHub<sup>[2]</sup>

XML bindings (probably Generated using Code Synthesis XSD<sup>[3]</sup>)



## References

1. Britten, R.D., et al., *FieldML, a proposed open standard for the Physiome project for mathematical model representation*. Medical & biological engineering & computing, 2013. **51**(11): p. 1191-1207.
2. <https://github.com/cellml/libcellml/>
3. <http://www.codesynthesis.com/products/xsd/>