

What's next for PMR2

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Te Whare Wānanga o Tāmaki Makaurau

What does PMR2 have now?

- Integrated Mercurial for storage of models.
 - Embedded workspaces.
- Uses Plone for presentation and content management.
 - Also user access control.
- Provide a presentational interface that is customizable and extensible through plugins through the Zope Component Architecture.



PMR2 was planned for

- The CellML model repository.
- A repository of other types of models, next up will most likely be FieldML support (cmgui/zinc viewer).
- Ultimately it will fulfill its role as The Physiome Model Repository.

The screenshot shows the CellML Model Repository homepage. The header includes the CellML logo, navigation tabs (Models Home, Exposures, Workspaces), a search bar, and user links (Log in, Register). The main content area is titled 'CellML Model Repository' and 'Main Model Listing'. It provides a list of processed model exposures with a note about the list's purpose. A 'Browse by category' section lists various biological and physiological categories. A 'Searching' section explains how to use the search bar. The footer contains copyright information and links to the site map, accessibility, and contact.

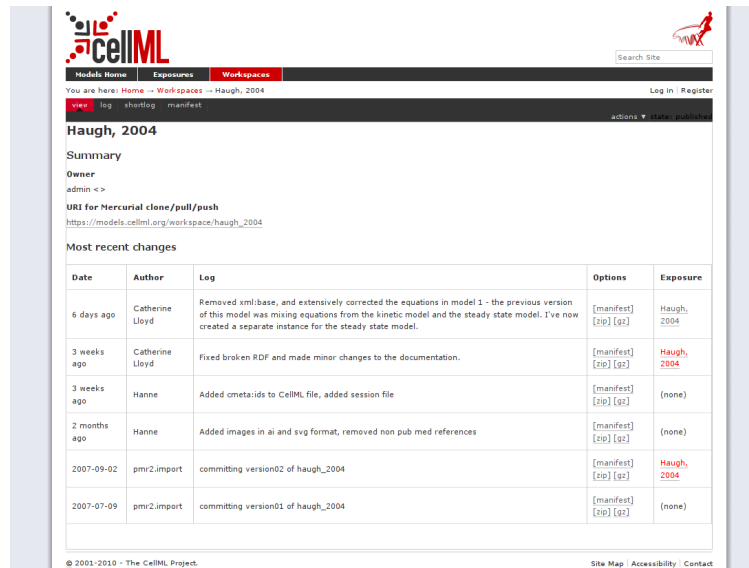
The screenshot shows the CellML Model Repository page for the 'Reconstruction of the action potential of ventricular myocardial fibres' model. The header is identical to the homepage. The main content area is titled 'Reconstruction of the action potential of ventricular myocardial fibres'. It includes a 'Model Status' section with a reference to the original paper, a 'Model Structure' section with a detailed description of the model, and a 'Model Curation' section with star ratings for various metrics. A 'Download Methods' section provides links to the complete archive and the file used to solve the model. A 'Views available' section lists the model metadata, mathematics, procedural C code, and OpenCell session. A 'Navigation' section provides a link to the model page. The footer contains copyright information and links to the site map, accessibility, and contact.

Recent new features

- User workspaces.
 - This allows user to create their own workspaces of any name.
 - Implemented; will be released in v0.3.
- Exposure File Notes.
 - Implemented as plugins for PMR2, they provide the views and its creation methods for specific file formats.
 - Implemented; released in v0.2.

User workspaces

- Every users of PMR2 can create workspaces in their user workspace folder.
- This eliminates workspace name clashes.
- Can support forking of models in the future if needed.



The screenshot shows the CellML website interface. At the top, there's a navigation bar with links for Models, Home, Exposures, and Workspaces. Below this, a breadcrumb trail indicates the current location: Home > Workspaces > Haugh, 2004. The main content area displays the workspace details for 'Haugh, 2004', including a summary, owner information (admin), and a URI for Mercurial clone/pull/push. A table titled 'Most recent changes' lists several updates, including corrections to equations, broken RDF, and added session files. The table has columns for Date, Author, Log, Options, and Exposure.

Date	Author	Log	Options	Exposure
6 days ago	Catherine Lloyd	Removed xml:base, and extensively corrected the equations in model 1 - the previous version of this model was mixing equations from the kinetic model and the steady state model. I've now created a separate instance for the steady state model.	[manifest] [zip] [gz]	Haugh, 2004
3 weeks ago	Catherine Lloyd	Fixed broken RDF and made minor changes to the documentation.	[manifest] [zip] [gz]	Haugh, 2004
3 weeks ago	Hanne	Added cmet:ids to CellML file, added session file	[manifest] [zip] [gz]	(none)
2 months ago	Hanne	Added images in ai and svg format, removed non pub med references	[manifest] [zip] [gz]	(none)
2007-09-02	pmr2.import	committing version02 of haugh_2004	[manifest] [zip] [gz]	Haugh, 2004
2007-07-09	pmr2.import	committing version01 of haugh_2004	[manifest] [zip] [gz]	(none)

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Exposure presentational plugins

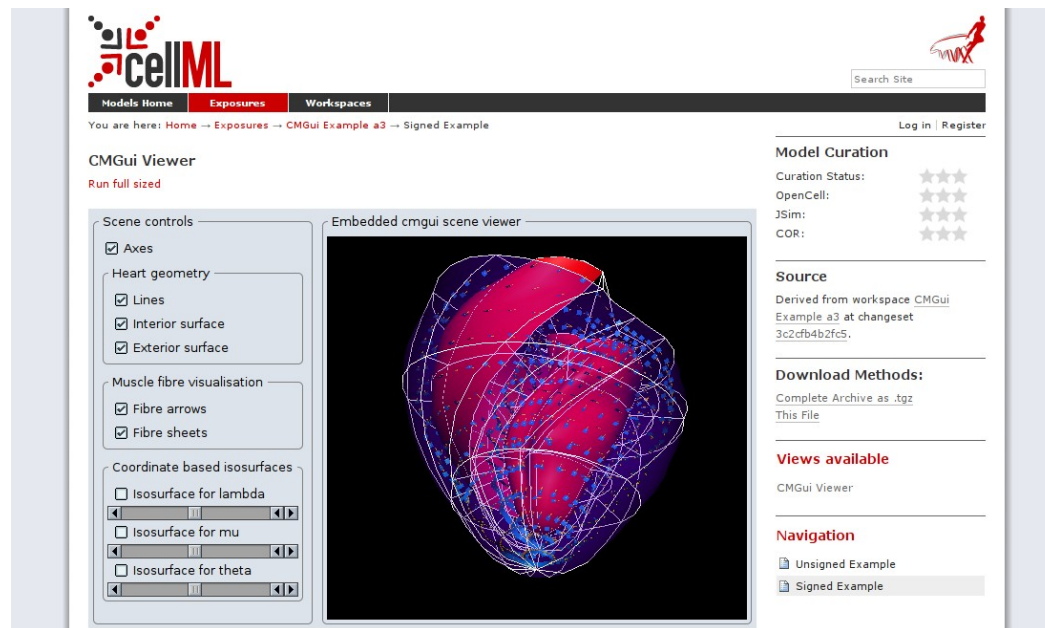
- There may be types of model (files) in the repository.
- Each of these files will then be "built", with notes (Exposure File Notes) attached to them.
- These notes provide storage for the generated data; viewers can be customized to render this data.
- Built on top of Zope's annotation module.
- Utilizes the Zope Component Architecture, which makes heavy use of the adapter pattern.
- This makes it relatively easy to add new plugins to improve the presentation of existing models or other model types.

Example Zope configuration file.

```
<configure
  xmlns="http://namespaces.zope.org/zope"
  xmlns:browser="http://namespaces.zope.org/browser"
  i18n_domain="fieldml">
  <adapter
    for="pmr2.app.content.interfaces.IExposureFile"
    factory=".CMGuiViewerNoteFactory"
    provides="fieldml.pmr2.interfaces.ICMGuiViewerNote"
    name="cmgui_viewer"
  />
  <utility
    name="cmgui_viewer"
    component=".annotator.CMGuiViewerAnnotatorFactory"
    provides="pmr2.app.annotation.interfaces.IExposureFileAnnotator"
  />
  <browser:page
    for="pmr2.app.content.ExposureFile"
    name="cmgui_viewer"
    class=".browser.CMGuiViewerView"
    permission="zope2.View"
  />
</configure>
```

Improving presentation of models

- The ZCML file in the previous slide was from the plugin that enabled this Zinc viewer view for exnode model files.
- In the future FieldML will be used instead.



Improving presentation of models

- Currently, exposures are just a rendering of simple, free formatted text with static images.

The screenshot displays the CellML website interface. At the top, the CellML logo is visible on the left, and a search bar is on the right. Below the logo, navigation tabs for 'Models Home', 'Exposures', and 'Workspaces' are present. The main content area shows the model title 'Reconstruction of the action potential of ventricular myocardial fibres' and its status. A detailed description of the model's origin and purpose is provided. To the right, a 'Model Curation' section shows star ratings for various attributes. Below that, a 'Source' section lists the workspace and changelog. A 'Download Methods' section offers options to download the archive or solve using OpenCell. A 'Views available' section lists metadata, mathematics, procedural code, and OpenCell sessions. A 'Navigation' section provides a link to the model's reconstruction. At the bottom, a diagram of a ventricular myocardial fiber is shown, illustrating the ionic currents involved in the action potential.

cellML

Models Home Exposures Workspaces

You are here: Home → Exposures → Beeler, Reuter, 1977 → Reconstruction of the action potential of ventricular myocardial fibres

Log in | Register

Reconstruction of the action potential of ventricular myocardial fibres

Model Status

This model has been curated by Penny Noble using Flavio Fenton's Java code as a reference (See <http://thevirtualheart.org/> for Java applet rendering of model - Java code is available from Dr Fenton.) An artificial stimulus component has been added to allow it to reproduce the action potential simulation shown in Figure 4 of the publication. The model is known to run and integrate in the PCEnv and COR CellML environments. A PCEnv session file is also associated with this model.

ValidateCellML detects unit inconsistency within this model.

Model Structure

In contrast to the earlier Purkinje fibre ionic current models of D. Noble (1962) and R.E. McAllister, D. Noble and R.W. Tsien (1975), the G.W. Beeler and H. Reuter 1977 model was developed to describe the mammalian ventricular action potential. Not all the ionic currents of the Purkinje fibre model are present in ventricular tissue; therefore, this model is simpler than the MNT model. The total ionic flux is divided into only four discrete, individual ionic currents (see the figure below). The main additional feature of the Beeler-Reuter ionic current model is a representation of the intracellular calcium ion concentration.

The complete original paper reference is cited below:

Reconstruction of the action potential of ventricular myocardial fibres, Beeler, G.W. and Reuter, H. 1977 *Journal of Physiology* , 268, 177-210. PubMed ID: 874889

The diagram illustrates a rectangular ventricular myocardial fiber. It shows four ionic currents flowing across the membrane: I_{Na} (sodium current) entering the cell, I_L (l-type calcium current) entering the cell, I_{Ca} (calcium current) entering the cell, and I_{K1} (potassium current) leaving the cell. The fiber is represented by a blue rectangular shape with a grid pattern.

Model Curation

Curation Status: ★★☆☆
OpenCell: ★★☆☆
JSim: ★★☆☆
COR: ★★☆☆

Source

Derived from workspace [Beeler, Reuter, 1977](#) at changeset [d4ac7e982034](#).

Download Methods:

[Complete Archive as .tgz](#)
[This File](#)
[Solve using OpenCell](#)

Views available

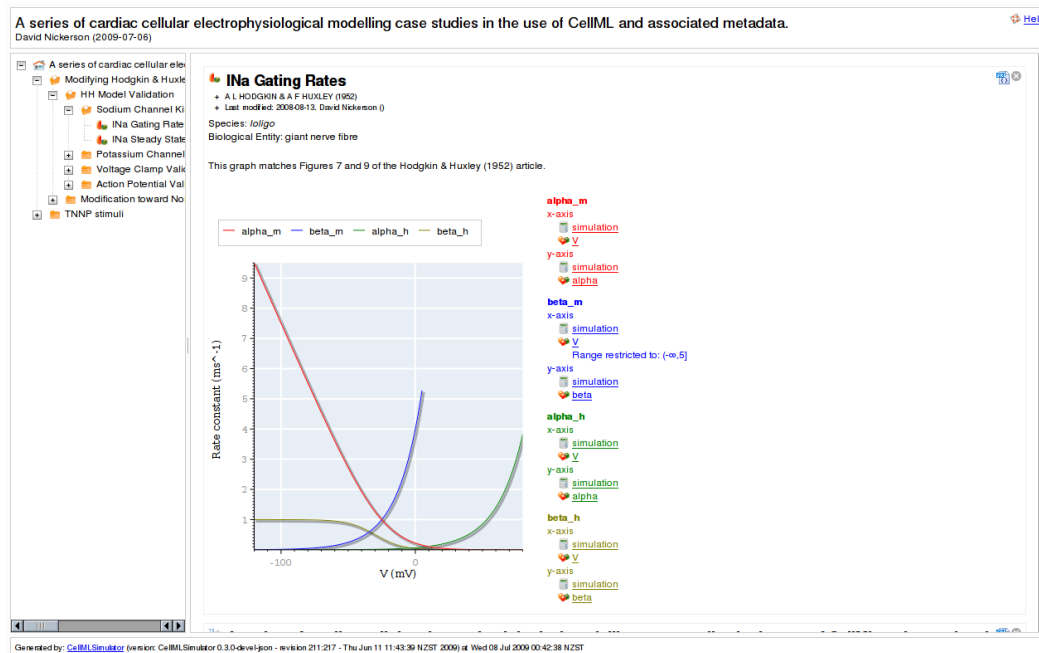
Model Metadata
Mathematics
Procedural C Code
OpenCell Session

Navigation

[Reconstruction of the action potential of ventricular myocardial fibres](#)

Improving presentation of models

- Through the exposure plugin system, it is possible to implement and integrate an interactive model viewer, such as CellMLSimulator.



Searching in PMR2

- Default search engine in Plone limits to text, meaning the RDF metadata within the file is ignored.
- So we will introduce RDF store, and support SPARQL for querying.
- Also provide pre-made search templates so users would not need to construct those SPARQL statements.
- This however depends on a more standardized metadata specification.

Web service access

- JSON/XMLRPC access to repository to gather information about workspaces/exposures.
- Thus other programs and/or websites can access the data stored in PMR2 through this machine friendly interface.
- Query any saved simulation results.

Fin

- Thank you for your attention.