Workspaces, exposures, and multiscale modelling

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Te Whare Wānanga o Tāmaki Makaurau

PMR2

- Workspace data agnostic mercurial repository
- Changeset a representation of a single revision of the content of a workspace
- Exposure a permanent link to a specific changeset with data rendered for the web
- Exposure plug-ins an extensible framework for rendering workspace content for web presentation
- Plone CMS workflow manager; user access controls; web presentation; etc.

An example from computational physiology

- Build up a multiscale model of the renal nephron
 - ion transporters, cellular models, segmental models, whole nephron...
- Share the various models with collaborators
- Publish the model along the way

 Disclaimer: not all the following features are implemented/integrated in either language specifications and/or supporting software tools – and such features may change considerably before they are supported.

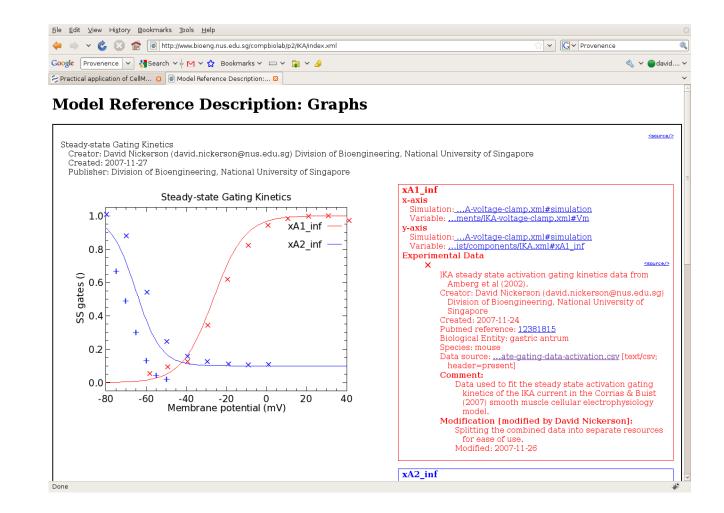
Membrane transporters



(Workspace)

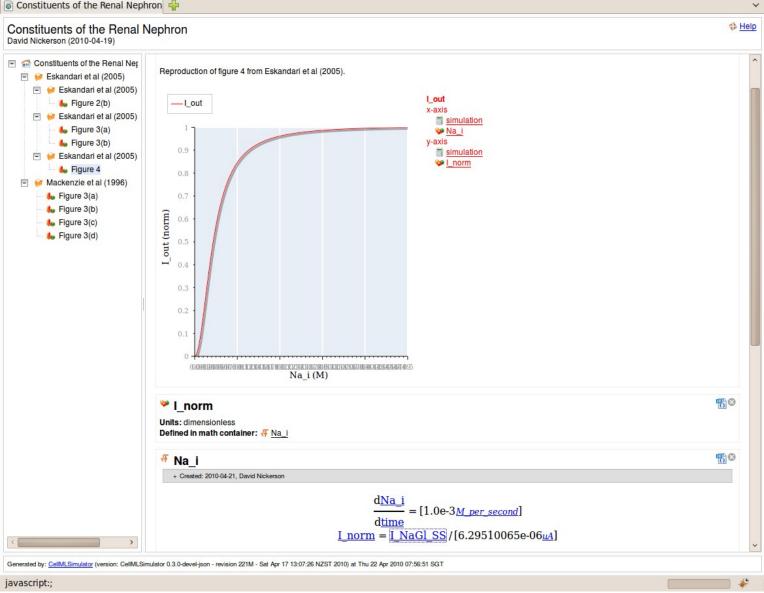


Membrane transporters

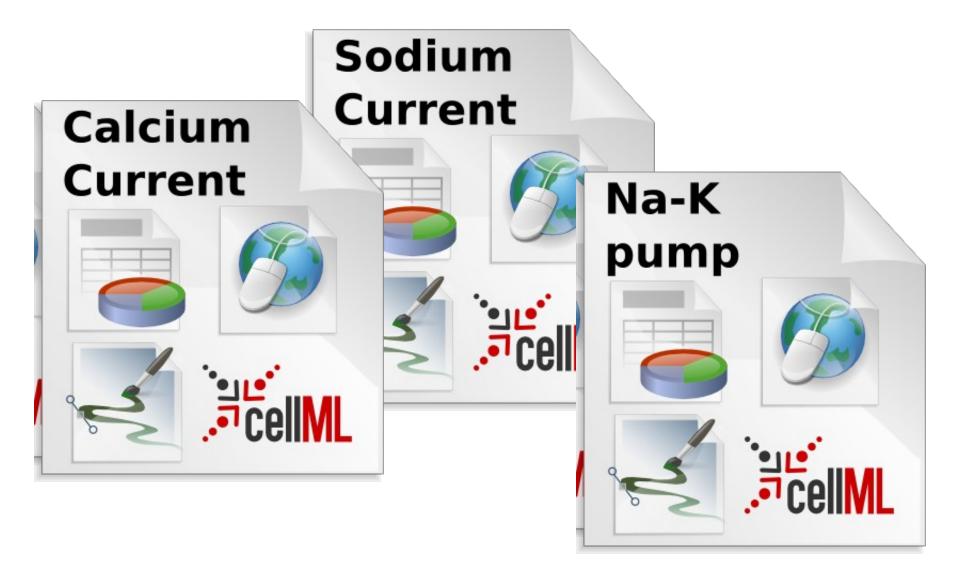


File Edit View History Bookmarks Tools Help

💿 Constituents of the Renal Nephron 🚽



Membrane transporters

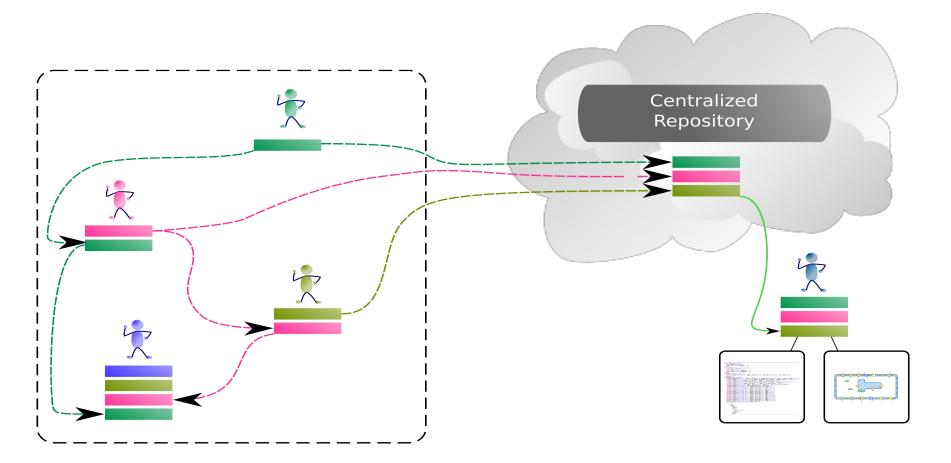


Assemble a cell model **Epithelial Cell** Sodium Current Calcium Na-K Current pump .**•** Ce

Embedded workspaces

- Intended to manage the separation of modules which are integrated to create a model
- Facilitate the sharing and reuse of model components independently from the source model
- Enables the development of the modules to proceed independently, thus the version of the workspaces embedded is also tracked
- Allows authors to make use of relative URIs when linking data resources providing a file system agnostic method to describe complex module relationships in a portable manner

Collaborative model development



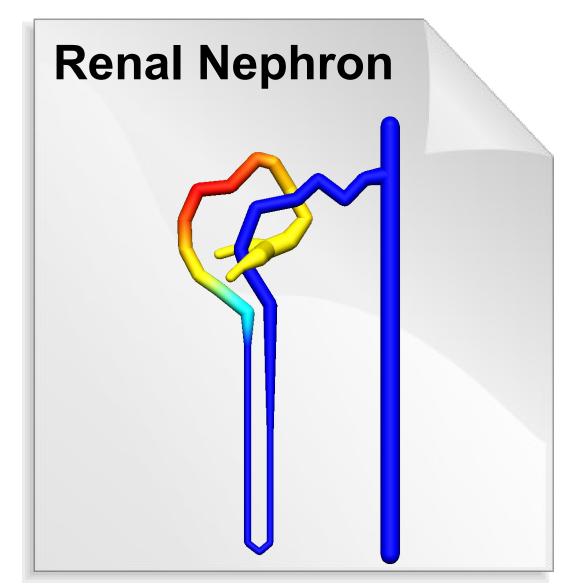
Versioning embedded workspaces

- Workspaces can be embedded at a specific revision or set to track the most recent revision of the source workspace
- Changes made to the source workspace will not affect the embedding workspace until the author explicitly chooses to update the embedded workspace
- Provides the author with the opportunity to review the changesets and make an informed decision regarding alterations to embedded revisions

Data agnostic workspaces

- Generic mercurial repositories
- Can contain any format data (currently relatively unrestricted)
 - CellML, SBML, FieldML, SED-ML, PDF, .doc, ...
- No restriction to models only
 - experimental data, simulation results, generated images, ...

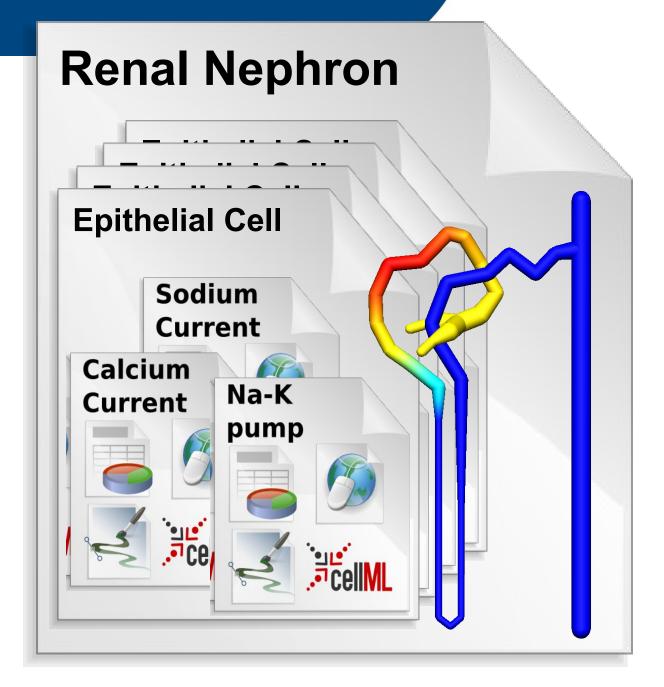
Multiscale models



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Nephron Interface

+ The Renal Nephron Close all 🖃 🗖 Nephron protein Reset view Input mode: Picking Ø **Proximal Tubule** - Anatomy Efferent Arterio Afferent Arterio First of the transporting tubule Glomerulus segments. 🔄 🗔 Proximal Tubule Proximal Co Proximal Str From the Bowman's Capsule, the filtered fluid + The Loop of Henle enters the first of the reabsorptive epithelial 📰 Macula Densa tubule segments - the proximal tubule. The 📰 Distal Convolut proximal tubule consists of a convoluted portion and a straight portion. This segment has a high Connecting Tub transport activity and is responsible for the bulk Collecting Duct of the salt and water reabsorption. Furthermore, 🔄 🛄 Physiology the majority of the key organic molecules Efferent Arterio (glucose and amino acids), as well as other Afferent Arterio important ions (K, Ca, HCO3), are actively Glomerulus reabsorbed in this segment. 🗄 🛅 Proximal Tubul + The second sec Related CellML models: 📰 Macula Densa Weinstein et al (2007) 🖶 🗂 Distal Convolut Thomas and Dagher (1994) 🗄 🛅 Connecting Tub 🗄 💼 Collecting Duct - C Modelling Studies 0 **Proximal Convoluted** - SGLT2 Inhibition 📰 Na-glucose 🕻 Tubule Control Inhibited Convoluted portion of the Proximal Tubule. As its name suggests, the proximal convoluted tubule undergoes a convoluted trajectory through the cortical region of the kidney, primarily the cortical labyrinth. > 0 Manhran April 2010 David Nickerson Done 1



OpenCMISS

- Connecting variables in CellML models to field components in a finite element model
 - Prototype for linking CellML and FieldML models?
- Allows information to flow in both directions
 - Field values can be controlled by the CellML model and CellML model variables can be controlled by field components
- Will have the ability to make use of many different CellML models which can be simulated independently
- Each CellML model may be replicated many millions of times for large scale problems
 - Distributed computing, GPUs, FPGAs, ...

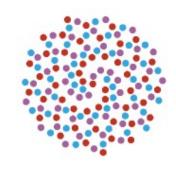
Acknowledgements



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MAURICE WILKINS CENTRE