From stars to flags: how we rate a model

Catherine Lloyd Auckland Bioengineering Institute





Stars



A star system denotes the curation status of a model



Not curated



Maths consistent with the published paper



No typos, units consistent, model complete, not over-constrained, and reproduces the published results;



Model satisfies physical constraints – eg conservation of mass, momentum, charge, etc.



But... we have problems





1 star and 2 stars are mutually exclusive



 3 star curation status can only be awarded by a "domain expert", who is not the model author themselves – which limits the number of 3 star models



Replacing stars with flags



Minimum Information Requested In the Annotation of Models

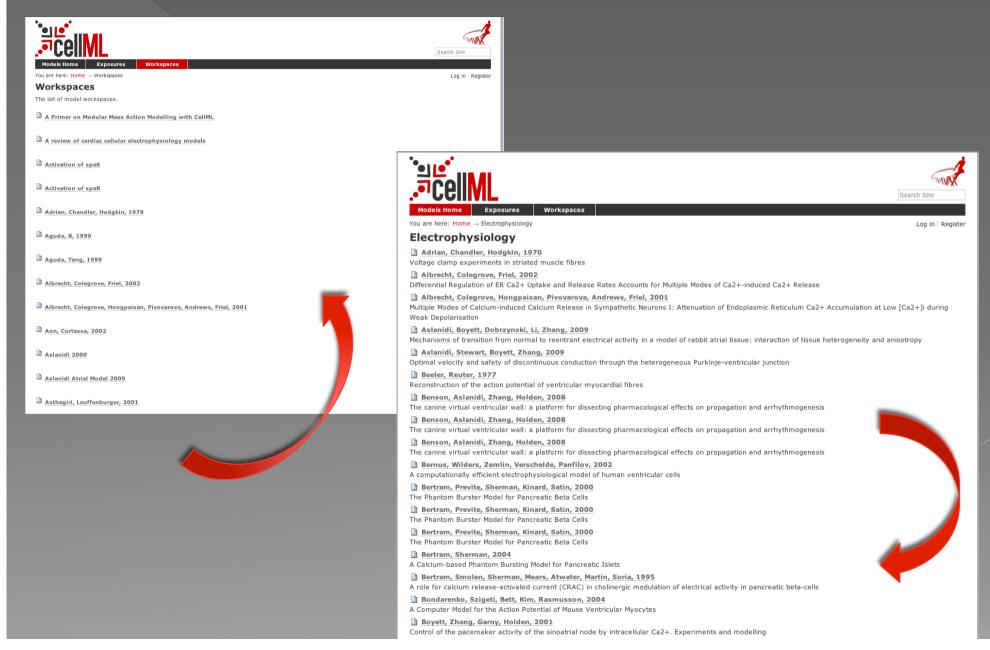


MIRIAM based flags

Application of the MIRIAM Standard as the CellML model curation flags							
	Tyson 1991	Goldbeter 1991	Noble 1998	Sneyd 2004	Cooling 2007	Ostby 2008	Lovell 2004
REFERENCE CORRESPONDENCE							
Is the model encoded in a public, standardized, machine-readable format?	yes	yes	yes	yes	yes	yes	yes
Does the model comply with the standard in which it is encoded?	yes	yes	yes	no	yes	no	yes
Does the model clearly relate to a single reference description. If the model is composed from different parts is there a description of the derived/combined model?	yes	yes	yes	yes	yes	yes	yes
Does the encoded model structure reflect the processes listed in the reference description?	yes	yes	yes	yes	yes	yes	yes
Is the model instantiated in a simulation: if it is are all quantitative attributes defined, including initial conditions?	yes	yes	yes	no	yes	yes	no
When instantiated, does the model reproduce all results given in the reference description within an epsilon (algorithms, round-up errors)?	yes	yes	yes	no	yes	yes	no
ATTRIBUTION ANNOTATION							
Is the model named?	yes	yes	yes	yes	yes	yes	yes
Is there a citation of the reference description associated with the model (complete							
citation, unique identifier, unambigous URL)? Note that this citation should allow identification the authors of the model.	yes	yes	yes	yes	yes	yes	yes
Are the name and contact of the model creators joined?	yes	yes	yes	yes	yes	yes	yes
Are the date and time of creation and last modification specified?	yes	yes	yes	yes	yes	yes	yes
Is the model linked to a precise statement about the terms of distribution?	yes	yes	yes	yes	yes	yes	yes
EXTERNAL RESOURCE ANNOTATION							
Are all the model components unambiguously identified (annotated with external resources		n/a	n/a	n/a	n/a	n/a	n/a
Is this annotation refined/is there a qualifier?	n/a	n/a	n/a	n/a	n/a	n/a	n/a
Does the annotation use MIRIAM URIs?	n/a	n/a	n/a	n/a	n/a	n/a	n/a
- Is the identifier analysed within the framework of the data type?	n/a	n/a	n/a	n/a	n/a	n/a	n/a
		_			_		
CURRENT CURATION STAR RATING	2	2	2	0	3	2	0



Curation flag display





Curation flag display

ves

yes

yes

ves

ves

ves



Contributions of H

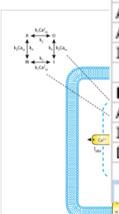
Model Status

This model is known to run in Ope helieve the units are correct in this

Model Structure

dependent channel-mediated curr which a number of point mutation focus for Safety Pharmacology reg changes in the formulations for tw human ventricular action notentia overall goal is to understand the g

Virag, Andras Varro, and Wayne R



REFERENCE CORRESPONDENCE

Is the model encoded in a public, standardized, machine-readable format? ves ventricular action Does the model comply with the standard in which it is encoded? ves

Does the model clearly relate to a single reference description. If the model is composed are balanced and consistent while from different parts is there a description of the derived/combined model?

Does the encoded model structure reflect the processes listed in the reference description? ABSTRACT: Action potential repola Is the model instantiated in a simulation: if it is are all quantitative attributes defined, work has shown that one of the K- including initial conditions?

which a number of point mutation

K+ channel isoform to inhibition b

When instantiated, does the model reproduce all results given in the reference description HERG-mediated K+ currents in re within an epsilon (algorithms, round-up errors)?

excised human ventricular tissue: ATTRIBUTION ANNOTATION

important first step in defining the IS the model named? ves

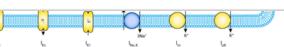
Is there a citation of the reference description associated with the model (complete Contributions of HERG K+ current citation, unique identifier, unambigous URL)? Note that this citation should allow PDF versions of the article are avail identification the authors of the model.

> Are the name and contact of the model creators joined? ves Are the date and time of creation and last modification specified? ves

Is the model linked to a precise statement about the terms of distribution?

EXTERNAL RESOURCE ANNOTATION

Are all the model components unambiguously identified (annotated with external resources n/a Is this annotation refined/is there a qualifier? n/a Does the annotation use MIRIAM URIs? n/a - Is the identifier analysed within the framework of the data type? n/a





Model exchange

Common set of curation flags (MIRIAM)

+

Model annotation

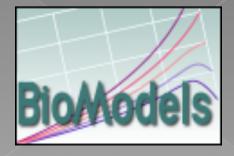
+

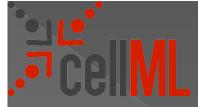
Language converters











Summary

- Current star system can be confusing and misinterpreted
- MIRIAM standard can form the basis of a new set of curation "flags"
- Together with model annotation and improved language converters this should allow model exchange between databases
- Combining limited curation resources