BioModels Database

Model Curation

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EBI is an Outstation of the European Molecular Biology Laboratory.





Branches of BioModels Database

- Curated
 - checked for MIRIAM compliance
 - curation figure added
 - model elements manually annotated
 - BioModels ID (BIOMDxxxx)
- Noncurated branch
 - only slightly altered by the curators
 - only publication and creation details added automatically and by curators



MIRIAM compliance for BioModels DB

Le Novère N. et al. Nature Biotechnology (2005), 23: 1509-1515

model must :

- be correctly encoded in a standard format (valid SBML)
- contain link to a single reference description (peer reviewed for BioModelsDB)
- contain creator's contact details
- reflect the structure of the processes and formulas described in the reference publication
- be able to reproduce the results given in the reference publication



Non Curated Branch

- valid SBML
- not MIRIAM compliant
 - not kinetic models (eg. FBA, stoichiometric maps)
 - can not reproduce results
- MIRIAM compliant
 - models contain kinetics that we do not curate up to now (eg. boolean models) or parts are not encoded in SBML (eg. spatial information)
 - significant tailback due to insufficient time and workforce → will be moved into curated branch as soon as possible



Curation Guidelines

- read publication
- go through SBML model and compare all elements
 - where possible create reactions out of differential equations
 - add names to unnamed reactions, rules and events
 - add compartments and move species and parameters accordingly
- change names and IDs to correspond to article
- try to reproduce one or two key results of the reference publication and create curation result (figure or table)
- add notes
- move model to the curated branch for annotation and publication

if any problems occure contact submitter first and, if necessary, authors all deviations from the publication have to be documented in the notes whenever the model's SBML file is changed significantly, the curation results have to be redone













Curation

Problems:

- typos and errors in publications both in model or results
- original models used in publication not available
- misleading diagrams or simplified kinetics in model/publication
- automatically generated SBML files can be hard to interpret

only few models stay in the curation pipeline:

- no publication info or never published
- large deviations from the model structure of the publication



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BioModels DB Curation

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