







### SED-ML support in JWS Online

Using SED-ML and COMBINE archives to reproduce Simulation Results

#### MARTIN PETERS, JACKY SNOEP, DAGMAR WALTEMATH

Department of Systems Biology & Bioinformatics, University of Rostock Department of Biochemistry, Stellenbosch University, South Africa







Reproducing Simulations

About JWS Online

What's new in JWS Online?

Summary – Achievements and Limitations



# Reproducing Figures Running a simulation by reading a paper

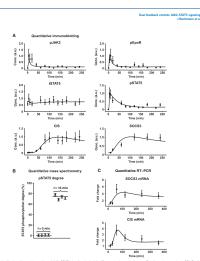


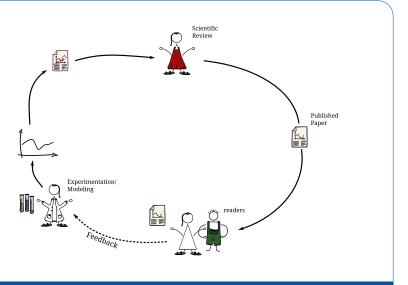
Figure 3. Model calculation with experimental date of AMCS STATS spalling obtained by different experimental behaviours. For all experiments, primary CFUE colds were stated and definital with STATE of AMCS STATS spalling obtained by eight spalling and spalling and



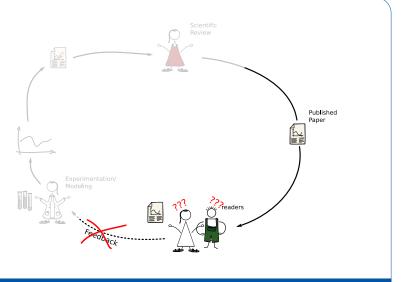
## Demo shows experiment database in JWS Online in action. Try it out here:

https://jjj.bio.vu.nl/models/experiments/ bachmann2011/simulate

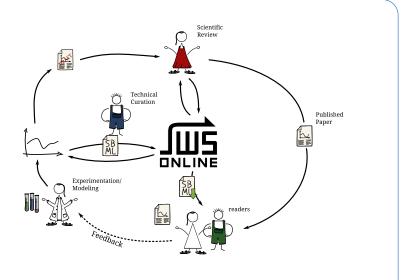






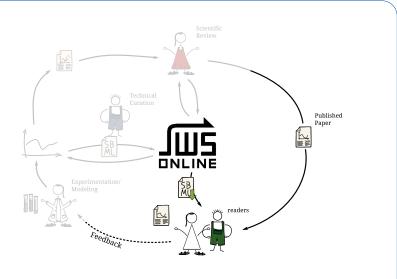






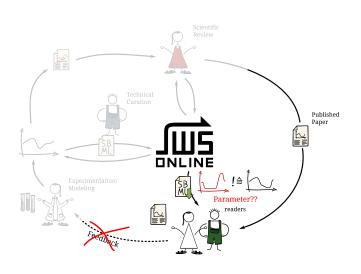


# About JWS Online a database for kinetic models

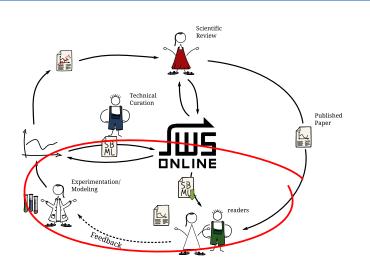


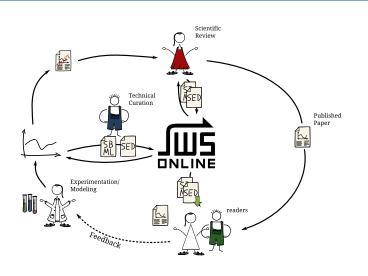


## About JWS Online a database for kinetic models



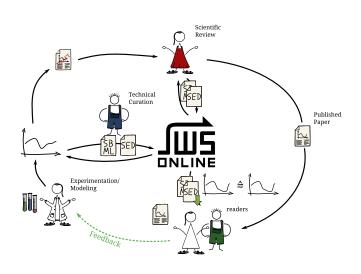




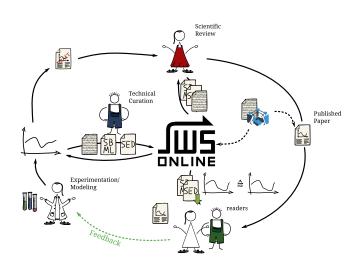


- Simulation Experiment Description Markup Language
   http://identifiers.org/combine.specifications/sed-ml.level-1.version-2
- Enabling reproducible in silico experiments since 2007
- Sophisticated multi-model/multi-experiment setups
- Support for experimental data and post-processing

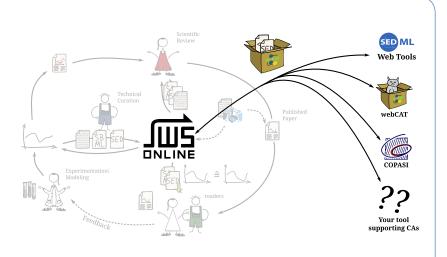












- Container format for bundling belonging to modeling or simulation experiment
- Enriched with meta information according to the OMEX standard
- Enable to bundle all necessary resources for reproduction
- http://co.mbine.org/standards/omex





- Dockerized JWS Online
   https://jws-docs.readthedocs.io/10\_docker.html
- Introduced a basic set of APIs into JWS Online https://jws-docs.readthedocs.io/8\_rest.html
- Widely extended the handling of Manuscript meta information
- Implemented SED-ML database into JWS Online https://jjj.bio.vu.nl/models/experiments/
- Enabled import and export of COMBINE Archives
- Introduced Reproduction of Simulation Experiments by a single click

https://jjj.bio.vu.nl/models/experiments/
bachmann2011/simulate



- Only Time-Course Simulations
- No Repeated-Tasks
- Excel Spreadsheets instead of NuML
- Only 2D Plots









#### That's it!



SEMS Task Force



JWS Team

Fabienne Lambusch Martin Scharm Dagmar Waltemath Mariam Nassar

Tom Gebhardt Vasundra Toure Ron Henkel Olaf Wolkenhauer Dawie van Niekerk Johann Eicher Daniel Palm Jacky Snoep











### That's it!

2016-09-21

### Want to talk? Drop me a message!



martin@freakybytes.net











#### References

- Köhn, Dagmar, and Nicolas Le Novere. 2008. SED-ML—an XML Format for the Implementation of the MIASE Guidelines. In Computational Methods in Systems Biology, 176–190. Springer.
- Olivier, Brett G, and Jacky L Snoep. 2004. Web-Based Kinetic Modelling Using JWS Online. Bioinformatics 20 (13): 2143–2144.
- Waltemath, Dagmar, Richard Adams, Frank T Bergmann, Michael Hucka, Fedor Kolpakov, Andrew K Miller, Ion I Moraru, et al. 2011. Reproducible Computational Biology Experiments with SED-ML-the Simulation Experiment Description Markup Language. BMC Systems Biology 5 (1): 1.
- Wolstencroft, K, Owen, S, Krebs, O, Nguyen, Q, Stanford, NJ, Golebiewski, M, Weidemann, A, Bittkowski, M, An, L, Shockley, D, Snoep, JL, Mueller, W, Goble, C (2015) SEEK: a systems biology data and model management platform. BMC Systems Biology, Issue 9:33, pages 33, 2015. DOI:10.1186/s12918-015-0174-y
- Bergmann, Frank T., Richard Adams, Stuart Moodie, Jonathan Cooper, Mihai Glont, Martin Golebiewski, Michael Hucka, et al. One File to Share Them All: Using the COMBINE Archive and the OMEX Format to Share All Information about a Modeling Project. arXiv:1407.4992 [Cs, Q-Bio], July 18, 2014. http://arxiv.org/abs/1407.4992.