

# Publications

## Research papers

<p>Meyer, K., Ostrenko, O., Bourantas, G., Morales-Navarrete, H., Porat-Shliom, N., Segovia-Miranda, F., Nonaka, H., Ghaemi, A., Verbavatz, J.M., Bruschi, L. and Sbalzarini, I., Kalaidzidis, Y., Weigert, R., Zerial M.</p> <p><a href="#">A Predictive 3D Multi-Scale Model of Biliary Fluid Dynamics in the Liver Lobule.</a> <i>Cell Systems</i> 4 (3), 277–290, 2017.</p>
<p>K.B. Hoffmann, A. Voss-Böhme, J.C. Rink, L. Bruschi.</p> <p><a href="#">A dynamically diluted alignment model reveals the impact of cell turnover on the plasticity of tissue polarity patterns.</a> <i>Journal of the Royal Society Interface</i> 14 (135), 20170466, 2017.</p>
<p>O. Ostrenko, P. Incardona, R. Ramaswamy, L. Bruschi, I.F. Sbalzarini.</p> <p><a href="#">pSSAlib: The partial-propensity stochastic chemical network simulator.</a> <i>PLoS Computational Biology</i> 13(12), e1005865, 2017.</p>
<p>O. Parvu and D. Gilbert.</p> <p><a href="#">A novel method to verify multilevel computational models of biological systems using multiscale spatio-temporal meta model checking.</a> <i>PLoS ONE</i>, 11(5): e0154847, 2016.</p>
<p>M. Herberg, T. Zerjatke, W. de Back, I. Glauche and I. Roeder.</p> <p><a href="#">Image-based quantification and mathematical modeling of spatial heterogeneity in ESC colonies.</a> <i>Cytometry: Part A</i>, 2015.</p>
<p>O. Parvu and D. Gilbert.</p> <p><a href="#">Automatic validation of computational models using pseudo-3D spatio-temporal model checking.</a> <i>BMC Systems Biology</i>, 8:124, 2014.</p>
<p>A. Köhn-Luque, W. de Back, Y. Yamaguchi, K. Yoshimura, M. A. Herrero and T. Miura.</p> <p><a href="#">Dynamics of VEGF matrix-retention in vascular network patterning.</a> <i>Physical Biology</i>, 10:066007, 2013.</p>
<p>W. de Back, R. Zimm, L. Bruschi</p> <p><a href="#">Transdifferentiation of pancreatic cells by loss of contact-mediated signaling.</a> <i>BMC Systems Biology</i>, 7:77, 2013.</p>
<p>W. de Back, J. X. Zhou, L. Bruschi</p> <p><a href="#">On the role of lateral stabilization during early patterning in the pancreas.</a> <i>Journal of the Royal Society Interface</i> 10(79):20120766, 2012.</p>
<p>A. Köhn-Luque, W. de Back, J. Starruß, A. Mattiotti, A. Deutsch, J. M. Pérez-Pomares, M. A. Herrero</p> <p><a href="#">Early embryonic vascular patterning by matrix-mediated paracrine signalling.</a> <i>PLoS ONE</i> 6(9):e24175, 2011.</p>
<p>J. Starruß, T. Bley, L. Søgaard-Andersen, A. Deutsch</p> <p><a href="#">A new mechanism for collective migration in <i>Myxococcus xanthus</i>.</a> <i>Journal of Statistical Physics</i>, 128, 269-286, 2007.</p>

## Also cited in

<p>Appleton, E., Madsen, C., Roehner, N. and Densmore, D.,</p> <p><a href="#">Design Automation in Synthetic Biology.</a> <i>Cold Spring Harbor Perspectives in Biology</i>, 2017.</p>
<p>Cytowski, M., Szymańska, Z., Umiński, P., Andrejczuk, G. and Raszkowski, K.</p> <p><a href="#">Implementation of an Agent-Based Parallel Tissue Modelling Framework for the Intel MIC Architecture.</a> <i>Scientific Programming</i>, 2017.</p>
<p>P. Macklin, H. Frieboes, J. Sparks, A. Ghaffarizadeh, S. Friedman, E. Juarez, E. Jonckheere, S. Mumenthaler</p> <p><a href="#">Progress Towards Computational 3-D Multicellular Systems Biology</a> In: <i>Systems Biology of Tumor Microenvironment</i> (edited by: K. Rejniak), 2016.</p>

J. S. Yu, N. Bagheri. <a href="#">Multi-class and multi-scale models of complex biological phenomena</a> <i>Current Opinion in Biotechnology</i> , 39:167-173, 2016.
O. Chara, E. Tanaka, L. Brusch. <a href="#">Mathematical Modeling of Regenerative Processes</a> . In: <i>Current Topics in Developmental Biology: Mechanisms of Regeneration</i> (edited by: B. Galliot) Volume 108, 2014.
S.J. Parker, K. Raedschelders and J. E. Van Eyk. <a href="#">Emerging proteomic technologies for elucidating context-dependent cellular signaling events: A big challenge of tiny proportions</a> . <i>Proteomics</i> , 2014.
L.A. D'Alessandro, S. Hoehme, A. Henney, D. Drasdo and U. Klingmüller. <a href="#">Unraveling liver complexity from molecular to organ level: Challenges and perspectives</a> . <i>Progress in biophysics and molecular biology</i> , 2014.
S. Kang, S. Kahan, J. McDermott, N. Flann and I. Shmulevich. <a href="#">Biocellion: accelerating computer simulation of multicellular biological system models</a> . <i>Bioinformatics</i> 30(2):3101-3108, 2014.

Beyond this selection, 44 sources have so far cited our paper in total (as of Dec. 2017).

## Education

Morpheus was used in the following courses:

<a href="#">OpenMultiMed Training School</a> at the Friedrich Alexander University Erlangen-Nürnberg, Germany February 22, 2018.
<a href="#">Multi-scale Biology Summer School</a> in Nottingham, organized by Prof. Markus Owen from the multi-scale biology network, UK 13 September 13, 2017.
Workshop on multicellular modeling at <a href="#">Institute for Computational Biology</a> , Helmholtz Zentrum München, Germany May 30, 2017.
Graduate school workshop on Systems Biology at <a href="#">HZI in Braunschweig</a> , Germany May 10, 2017.
<a href="#">ESMTB/EMS Summer School on Tissue Mechanics</a> at the Lorentz center in Leiden, the Netherlands July 25-29, 2016.
<a href="#">GSCN</a> workshop on <a href="#">Computational Stem Cell Biology</a> 1-2 December 2014.
Annual <a href="#">DIPP</a> 5-day course on Spatio-temporal Pattern Formation in Cells and Tissues Autumn 2012 till 2017.
<a href="#">ECMI modeling week European Summer School in Industrial Mathematics and Modelling Week</a> (ESSIM2012) August 12-22, 2012.
Described in this paper: F. Rost, A. Quintero, M. Myllykoski, A. Igolkina, A. Rohde O'Sullivan Freltoft, N. Dixit <a href="#">Morphogenesis and Dynamics of Multicellular Systems</a> ECMI Newsletter, 52, October 2012.

## Conferences

Talks and poster presentations at the following conferences and workshops:

W. de Back, Morpheus 2: Modeling and simulation platform for multicellular systems biology. And MorpheusML: declarative markup language for multicellular systems biology, Workshop "Towards a unified framework for benchmarking multi-cellular models and simulation software" (organized D. Drasdo and S. Hoehme et al., Leipzig, March 2016 (invited talks)

W. de Back, [Spatial multiscale computational systems biology](#), Schloss **Dagstuhl** - Leibniz-Zentrum für Informatik, Germany, November 2014 (talk)

L. Brusch, W. de Back, J. Starruß, A. Deutsch, Morpheus: a User-friendly Modeling Environment for Multiscale and Multicellular Systems Biology, Symposium on "*Executable cell biology for tissue engineering and regenerative medicine*", **TERMIS-EU**, Genova, Italy, 2014. ([abstract](#)) (talk)

W. de Back, Integrative modeling in developmental systems biology, Institute for Medical Informatics and Biometry, Medical Faculty „Carl Gustav Carus“, Technische Universität Dresden, 2013. (talk)

W. de Back, A. Deutsch, Morpheus: Modeling and Simulation in Multicellular System Biology, 1st Sino-German Workshop on Simulation, Changsha, China, 2011. (talk)

From:

<https://imc.zih.tu-dresden.de/wiki/morpheus/> - **Morpheus**

Permanent link:

<https://imc.zih.tu-dresden.de/wiki/morpheus/doku.php?id=documentation:publications&rev=1629203754>

Last update: **14:35 17.08.2021**

