Advanced Research on Biologically Inspired Cognitive Architectures, 2017, pages 117-125

## The BioDynaMo project: Experience report

Bauer R., Breitwieser L., Di Meglio A., Johard L., Kaiser M., Manca M., Mazzara M., Rademakers F., Talanov M., Tchitchigin A. *Kazan Federal University, 420008, Kremlevskaya 18, Kazan, Russia* 

## Abstract

© 2017, IGI Global. All rights reserved. Computer simulations have become a very powerful tool for scientific research. Given the vast complexity that comes with many open scientific questions, a purely analytical or experimental approach is often not viable. For example, biological systems comprise an extremely complex organization and heterogeneous interactions across different spatial and temporal scales. In order to facilitate research on such problems, the BioDynaMo project aims at a general platform for computer simulations for biological research. Since scientific investigations require extensive computer resources, this platform should be executable on hybrid cloud computing systems, allowing for the efficient use of state-of-the-art computing technology.

http://dx.doi.org/10.4018/978-1-5225-1947-8.ch006

## References

- [1] Kane, Hohman, Cerami, McCormick, Kuhlmman, & Byrd. (2006). Agile methods in biomedical software development: a multi-site experience report. BMC Bioinformatics, 7(1).
- [2] Kelly, D. (2015). Scientific Software Development Viewed as Knowledge Acquisition: Towards Understanding the Development of Risk-Averse Scientific Software. Proceedings of the ACM 2011 conference on Computer supported cooperative work. ACM.
- [3] Howison, J., & Herbsleb, J. D. (2015). Scientific software production: incentives and collaboration. Proceedings of the ACM 2011 conference on Computer supported cooperative work. doi:10.1016/j.jss.2015.07.027
- [4] Whitmore, A., Choi, N., & Arzrumtsyan, A. (2015). Open source software: The role of marketing in the diffusion of innovation. Information Technology and Control, 38(2).
- [5] Howison, J., Deelman, E., McLennan, M. J., da Silva, R. F., & Herbsleb, J. D. (2015). Understanding the scientific software ecosystem and its impact: Current and future measures. Research Evaluation.
- [6] Zubler, F., & Douglas, R. (2009). A framework for modeling the growth and development of neurons and networks. Frontiers in Computational Neuroscience, 3. doi:10.3389/neuro.10.025.2009 PMID:19949465
- [7] Sutter, H. (2005). The free lunch is over: A fundamental turn toward concurrency in software. Dr. Dobbs Journal, 30(3), 202-210.
- [8] Khazeev, M., Rivera, V., & Mazzara, M. (2016). Usability of AutoProof: a case study of static debugging. The 5th international Conference in Software Engineering for Defense Applications.
- [9] Meyer, B. (2009). Touch of Class: Learning to Program Well with Objects and Contracts. Springer Publishing Company, Incorporated. doi:10.1007/978-3-540-92145-5
- [10] Cousot, P., & Cousot, R. (1977). Abstract Interpretation: A Unified Lattice Model for Static Analysis of Programs by Construction or Approximation of Fixpoints. Proceedings of the 4th ACM SIGACT-SIGPLAN Symposium on Principles of Programming Languages. ACM. doi:10.1145/512950.512973
- [11] Clarke, J., Grumberg, O., & Peled, D. A. (1999). Model Checking. MIT Press.
- [12] Loveland, D. W. (1978). Automated Theorem Proving: A Logical Basis (Fundamental Studies in Computer Science). Elsevier North-Holland.

- [13] Bauer, R., Zubler, F., Pfister, S., Hauri, A., Pfeiffer, M., Muir, D. R., & Douglas, R. J. (2014). Developmental selfconstruction and self-configuration of functional neocortical neuronal networks. PLoS Computational Biology, 10(12), e1003994. doi:10.1371/journal.pcbi.1003994 PMID:25474693
- [14] Freund. (2014). Numerical simulation of flowing blood cells. Annual Review of Fluid Mechanics.
- [15] Izhikevich & Edelman. (2008). Large-scale model of mammalian thalamocortical systems. Proceedings of the national academy of sciences.
- [16] Eklund, Nichols, & Knutsson. (2016). Cluster failure: Why fmri inferences for spatial extent have inflated falsepositive rates. Proceedings of the National Academy of Sciences.