

POSTER PRESENTATION

Open Access

From laptops to supercomputers: a single highly scalable code base for spiking neuronal network simulations

Susanne Kunkel^{1,2,3*}, Maximilian Schmidt², Jochen M Eppler², Hans E Plessner⁴, Jun Igarashi⁵, Gen Masumoto⁶, Tomoki Fukai⁵, Shin Ishii⁷, Abigail Morrison^{1,2,3,8}, Markus Diesmann^{1,2,3,9}, Moritz Helias^{2,5}

From Twenty Second Annual Computational Neuroscience Meeting: CNS*2013
Paris, France. 13-18 July 2013

Over the last couple of years, supercomputers such as the Blue Gene/Q system JUQUEEN in Jülich and the K computer in Kobe have become available for neuroscience research. These massively parallel systems open the field for a new class of scientific questions as they provide the resources to represent and simulate brain-scale networks, but they also confront the developers of simulation software with a new class of problems. Initial tests with our neuronal network simulator NEST [1] on JUGENE (the predecessor of JUQUEEN) revealed that in order to exploit the memory capacities of such machines, we needed to improve the parallelization of the fundamental data structures. To address this, we developed an analytical framework [2], which serves as a guideline for a systematic and iterative restructuring of the simulation kernel. In December 2012, the 3rd generation technology was released with NEST 2.2, which enables simulations of 10^8 neurons and 10,000 synapses per neuron on the K computer [3].

Even though the redesign of the fundamental data structures of NEST is driven by the demand for simulations of interacting brain areas, we do not aim at solutions tailored to a specific brain-scale model or computing architecture. Our goal is to maintain a single highly scalable code base that meets the requirements of such simulations whilst still performing well on modestly dimensioned lab clusters and even laptops.

Here, we introduce the 4th generation simulation kernel and describe the development workflow that yielded

the following three major improvements: the self-collapsing connection infrastructure, which takes up significantly less memory in the case of few local targets, the compacted node infrastructure, which causes only negligible constant serial memory overhead, and the reduced memory usage of synapse objects, which does not affect the precision of synaptic state variables. The improved code does not compromise on the general usability of NEST and will be merged into the common code base to be released with NEST 2.4. We show that with the 4g technology it will be possible to simulate networks of 10^9 neurons and 10,000 synapses per neuron on the K computer.

Acknowledgements

Partly supported by the early access to the K computer at the RIKEN Advanced Institute for Computational Science, by the VSR computation time grant JINB33 on the JUGENE and JUQUEEN supercomputers in Jülich, the Alliance on Systems Biology, Initiative and Networking Fund and Portfolio theme SMHB of the Helmholtz Association, the Jülich-Aachen Research Alliance (JARA), the Next-Generation Supercomputer Project of MEXT, EU Grant 269921 (BrainScaleS), Research Council of Norway Grant 178892/V30 (eNeuro) and access to NOTUR supercomputing facilities. All network simulations carried out with NEST (<http://www.nest-initiative.org>).

Author details

¹Simulation Laboratory Neuroscience - Bernstein Facility Simulation and Database Technology, Institute for Advanced Simulation, Jülich Aachen Research Alliance, Jülich Research Centre, Germany. ²Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6), Jülich Research Centre and JARA, Germany. ³Bernstein Center Freiburg, Albert-Ludwig University of Freiburg, Germany. ⁴Department of Mathematical Sciences and Technology, Norwegian University of Life Sciences, Aas, Norway. ⁵Laboratory for Neural Circuit Theory, RIKEN Brain Science Institute, Wako, Japan. ⁶High-Performance Computing Team, RIKEN Computational Science Research Program, Kobe, Japan. ⁷Integrated Systems Biology Laboratory, Department of Systems Science, Graduate School of Informatics, Kyoto University, Japan. ⁸Institute of Cognitive Neuroscience, Faculty of Psychology, Ruhr-University Bochum, Germany. ⁹Medical Faculty, RWTH University, Aachen, Germany.

* Correspondence: kunkel@fz-juelich.de

¹Simulation Laboratory Neuroscience - Bernstein Facility Simulation and Database Technology, Institute for Advanced Simulation, Jülich Aachen Research Alliance, Jülich Research Centre, Germany

Full list of author information is available at the end of the article

Published: 8 July 2013

References

1. Gewaltig MO, Diesmann M: **NEST (NEural Simulation Tool)**. *Scholarpedia* 2007, 2(4):1430.
2. Kunkel S, Potjans TC, Eppler JE, Plesser HE, Morrison A, Diesmann M: **Meeting the memory challenges of brain-scale simulation**. *Front Neuroinform* 2012, 5:35.
3. Helias M, Kunkel S, Masumoto G, Igarashi J, Eppler JE, Ishii S, Fukai T, Morrison A, Diesmann M: **Supercomputers ready for use as discovery machines for neuroscience**. *Front Neuroinform* 2012, 6:26.

doi:10.1186/1471-2202-14-S1-P163

Cite this article as: Kunkel et al.: From laptops to supercomputers: a single highly scalable code base for spiking neuronal network simulations. *BMC Neuroscience* 2013 14(Suppl 1):P163.

**Submit your next manuscript to BioMed Central
and take full advantage of:**

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

Submit your manuscript at
www.biomedcentral.com/submit

