

Necessity and feasibility of brain-scale simulations at cellular and synaptic resolution

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www.csn.fz-juelich.de
www.nest-initiative.org

History of brain-scale simulations on K computer

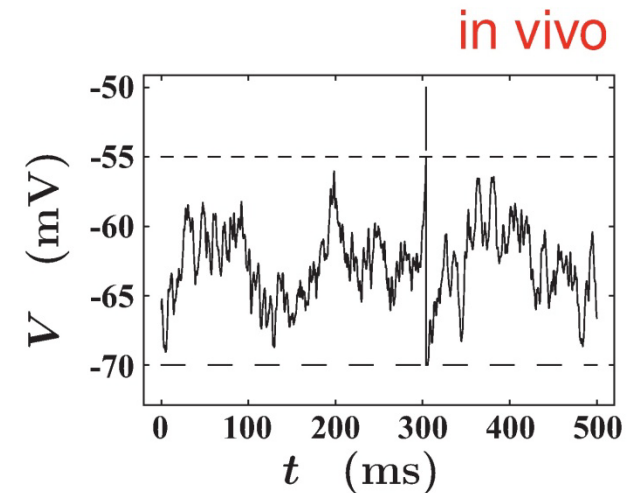
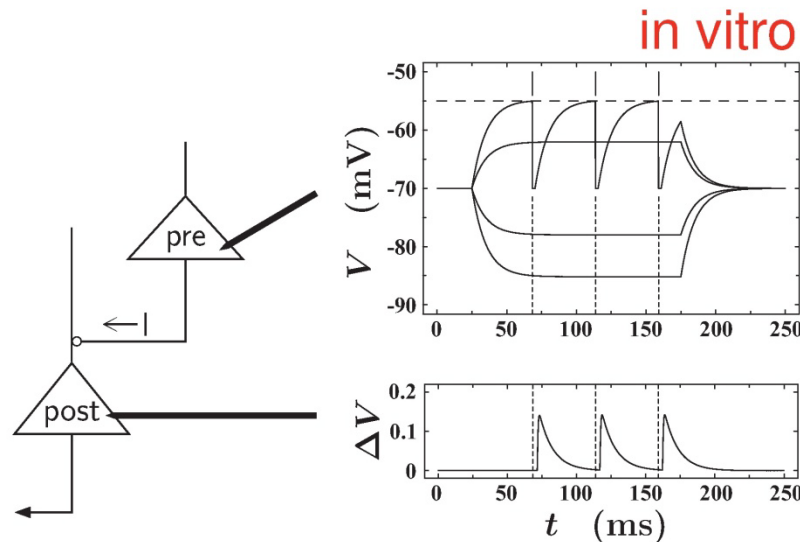
- work reported started in 2006
- Next-Generation Supercomputing project of MEXT
- group at RIKEN Brain Science Institute (BSI) 2006-2011
- from March 2011 Juelich Research Centre

- special thanks to
 - Ryutaro Himeno
 - Mitsuhisa Sato
 - Naoya Maruyama

Diesmann, M (2012) Proceedings of the 4th Biosupercomputing Symposium Tokyo 83–85

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Fundamental interaction between neurons



- current injection into pre-synaptic neuron causes excursions of membrane potential
- supra-threshold value causes spike transmitted to post-synaptic neuron
- post-synaptic neuron responds with small excursion of potential after delay
- inhibitory neurons (20%) cause negative excursion

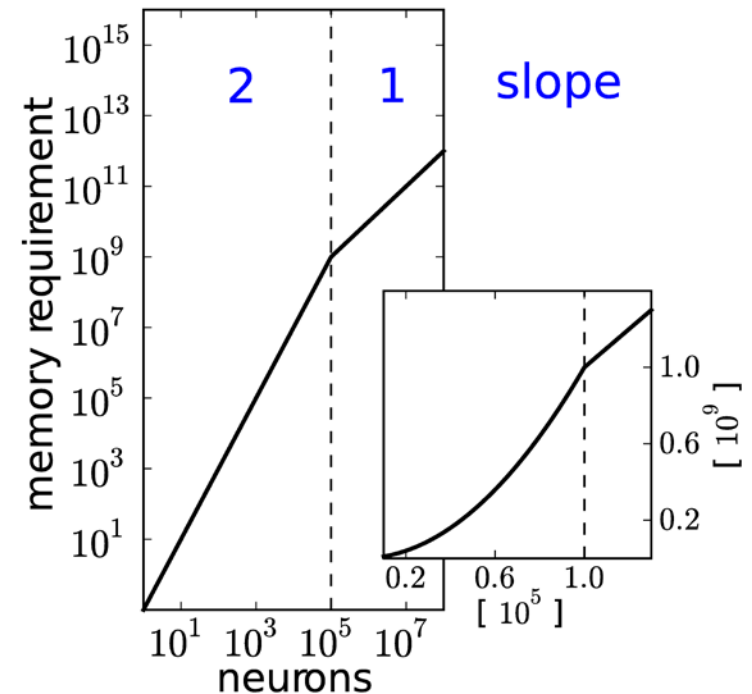
- each neuron receives input from 10,000 other neurons
- causing large fluctuations of membrane potential
- emission rate of 1 to 10 spikes per second

Realistic local cortical networks

- connectivity $c = 0.1$
- synapses per neuron = 10^4
- ⇒ minimal network size = 10^5

- network $N = 10^5$
 - considered **elementary unit**
 - corresponding to 1 mm^3

- total number of synapses = $(cN) \cdot N$ ⇒ possible



⇒ possible

Morrison A, Mehring C, Geisel T, Aertsen A, Diesmann M (2005) Neural Comput 17(8):1776-1801

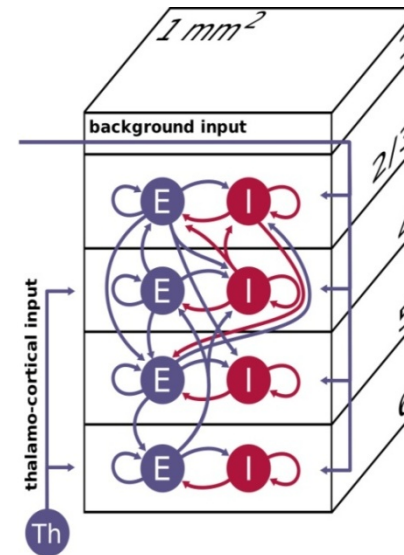
Morrison A, Straube S, Plesser HE, Diesmann M (2007) Neural Comput 19(1):47-79

Local cortical microcircuit

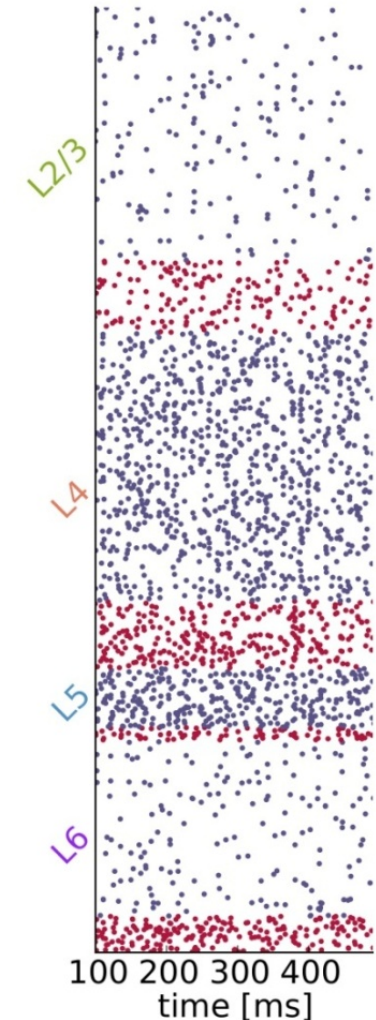
taking into account layer and neuron-type specific connectivity is sufficient to reproduce experimentally observed:

- asynchronous-irregular spiking of neurons
- higher spike rate of inhibitory neurons
- correct distribution of spike rates across layers
- integrates knowledge of more than 50 experimental papers

Potjans TC & Diesmann M (2014) The cell-type specific connectivity of the local cortical network explains prominent features of neuronal activity. *Cerebral Cortex* 24 (3): 785-806



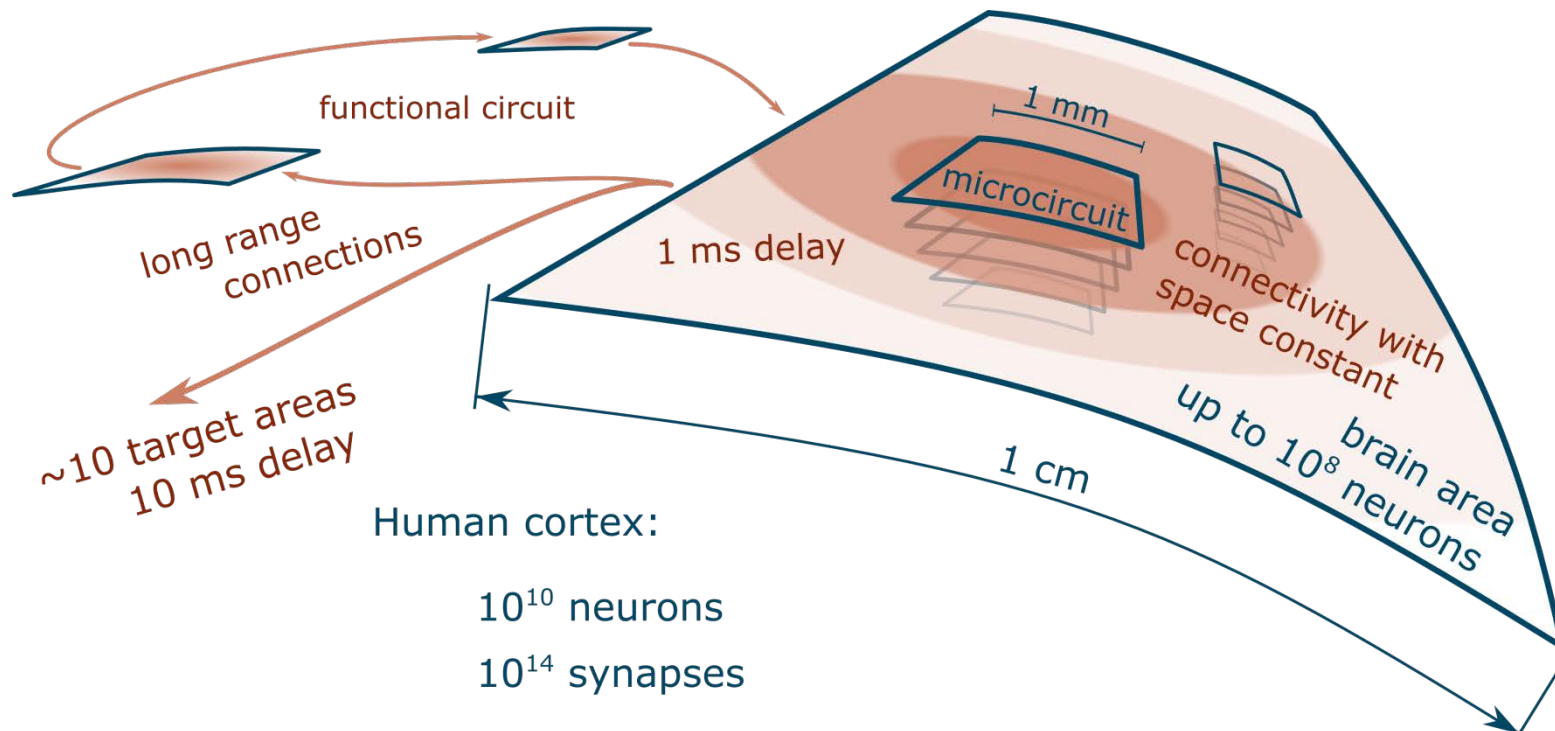
10⁵ neurons
10⁹ synapses



available at: www.opensourcebrain.org

Critique of local network model

a **network of networks** with at least three levels of organization:



- neurons in local microcircuit models are missing 50% of synapses
- e.g., power spectrum shows discrepancies, slow oscillations missing
- solution by taking brain-scale anatomy into account

Meso- and macro-scale measures

brain-scale networks basis for:

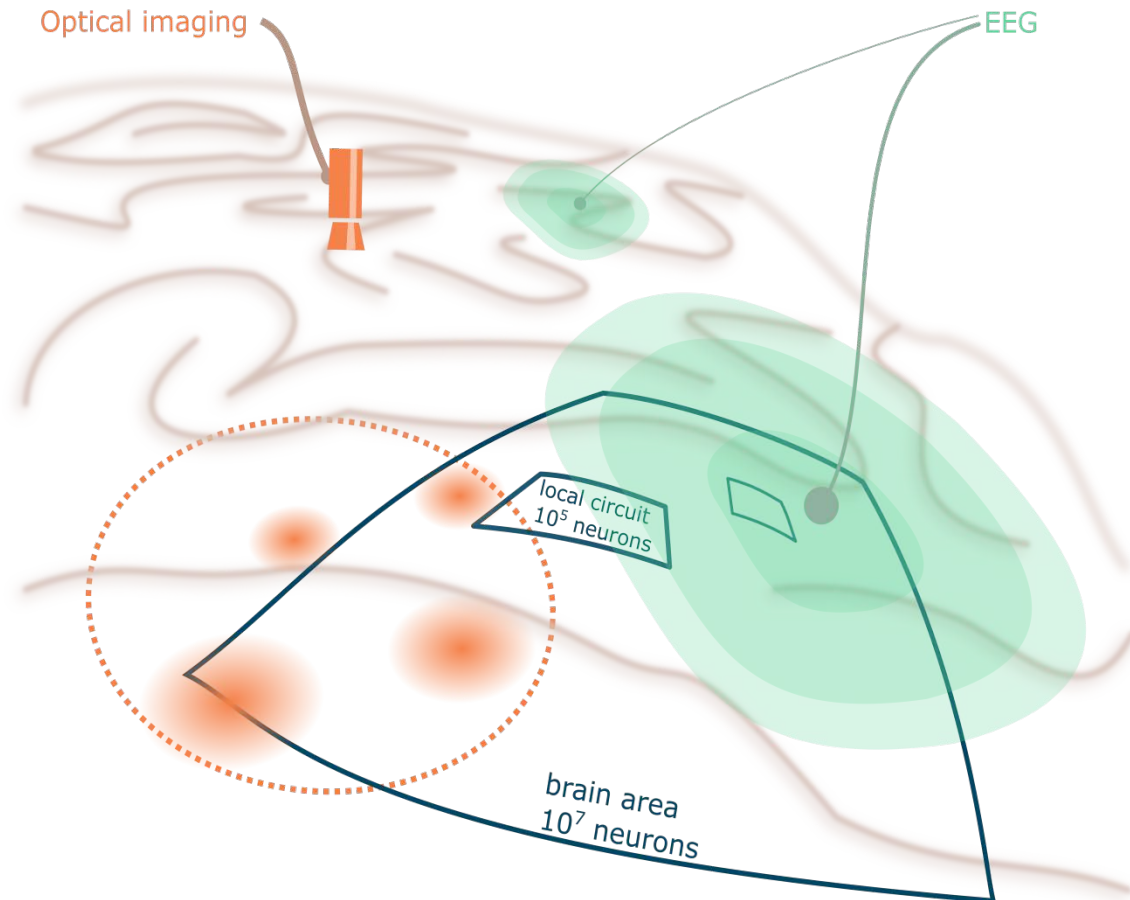
- further measures by forward modeling
- comparison with mean-field models

mesoscopic measures

- local field potential (LFP)
- voltage sensitive dyes (VSD)

and macroscopic measures

- EEG, MEG
- fMRI resting state networks



Feasibility and necessity

- Can we do simulations at the brain scale?
- Do we need to simulate full scale (at cellular resolution)?



Supercomputers ready for use as discovery machines for neuroscience

Moritz Helias^{1,2*}, Susanne Kunkel^{1,3,4}, Gen Masumoto⁵, Jun Igarashi⁶, Jochen Martin Eppler¹, Shin Ishii⁷, Tomoki Fukai⁶, Abigail Morrison^{1,3,4,8} and Markus Diesmann^{1,2,4,9}

¹ Institute of Neuroscience and Medicine (INM-6), Computational and Systems Neuroscience, Jülich Research Centre, Jülich, Germany

² RIKEN Brain Science Institute, Wako, Japan

³ Simulation Laboratory Neuroscience – Bernstein Facility for Simulation and Database Technology, Institute for Advanced Simulation, Jülich Aachen Research Alliance, Jülich Research Centre, Jülich, Germany

⁴ Bernstein Center Freiburg, Albert-Ludwig University of Freiburg, Freiburg, Germany

makes supercomputers accessible for neuroscience

provides the evidence that neuroscience can exploit petascale systems



Spiking network simulation code for petascale computers

Susanne Kunkel^{1,2*}, Maximilian Schmidt³, Jochen M. Eppler³, Hans E. Plesser^{3,4}, Gen Masumoto⁵, Jun Igarashi^{6,7}, Shin Ishii⁸, Tomoki Fukai⁷, Abigail Morrison^{1,3,9}, Markus Diesmann^{3,7,10} and Moritz Helias^{2,3}

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

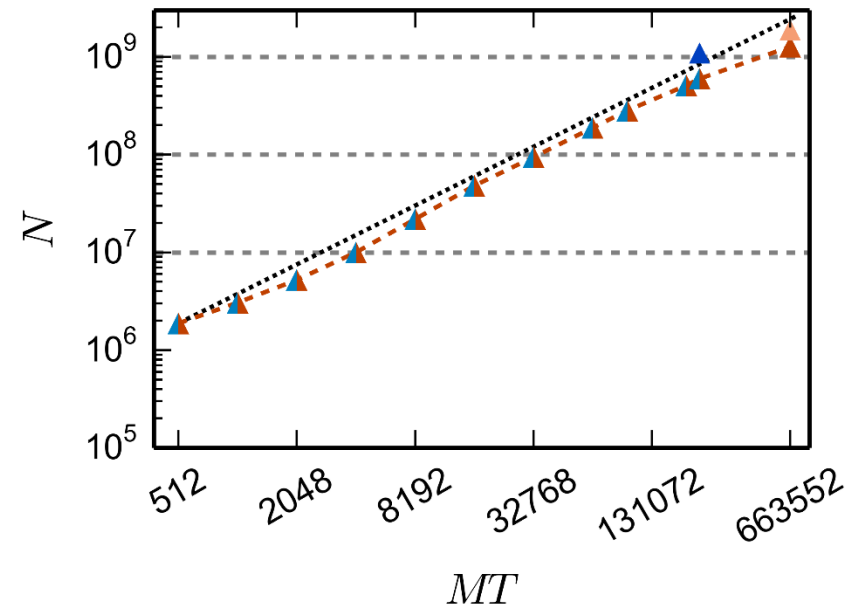
² Programming Environment Research Team, RIKEN Advanced Institute for Computational Science, Kobe, Japan

³ Institute of Neuroscience and Medicine (INM-6), Institute for Advanced Simulation (IAS-6), Jülich Research Centre and JARA, Jülich, Germany

⁴ Department of Mathematical Sciences and Technology, Norwegian University of Life Sciences, Aas, Norway

NEST – Maximum network size

- using 663,552 cores of K
- using 229,376 cores of JUQUEEN
- worst case: random network
- exc-exc STDP



- largest general network simulation performed to date:
 - 1.86x10⁹ neurons, 6000 synapses per neuron
 - 1.08x10⁹ neurons, 6000 synapses per neuron



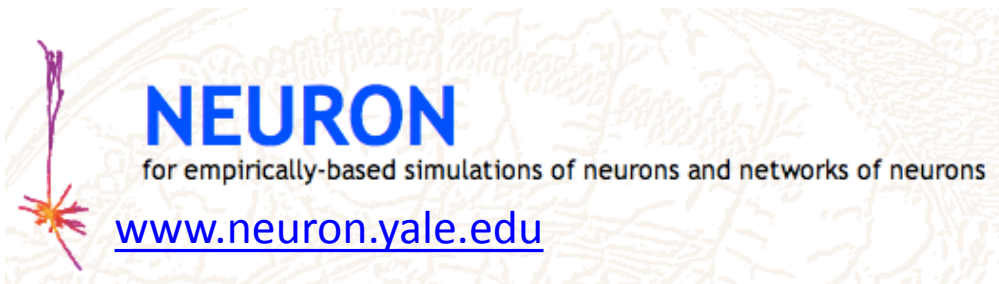
NEST simulation software

European Human Brain Project

- simulation engines in ramp-up phase



www.nest-initiative.org

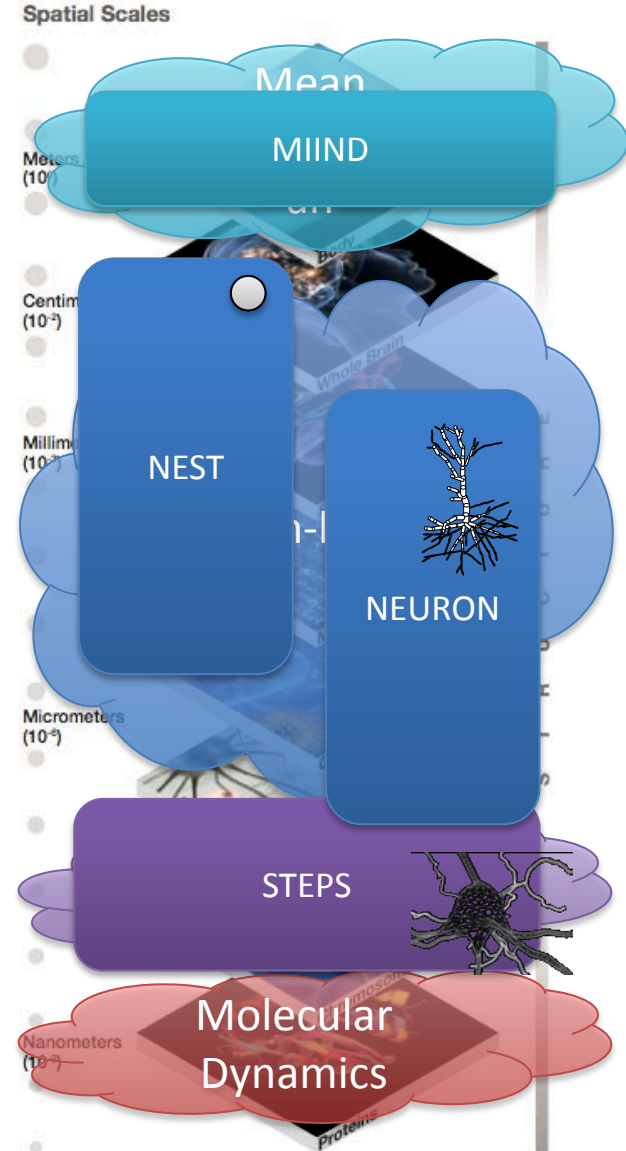


STEPS
Stochastic Engine for Pathway Simulation



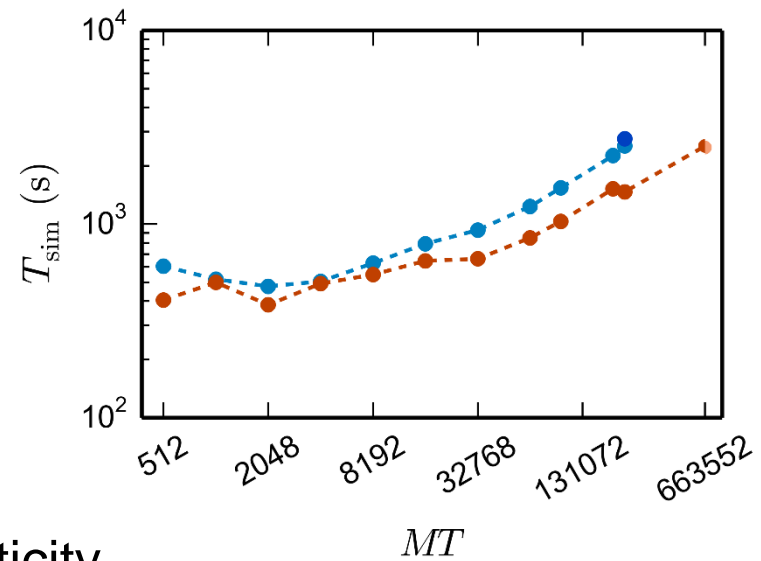
Erik De Schutter (OIST Okinawa)

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NEST – Scaling of run time

- runtime for 1 second biological time:
 - between 6 and 42 min on K computer
 - between 8 and 41 min on JUQUEEN
- wiring: between 3 and 15 min



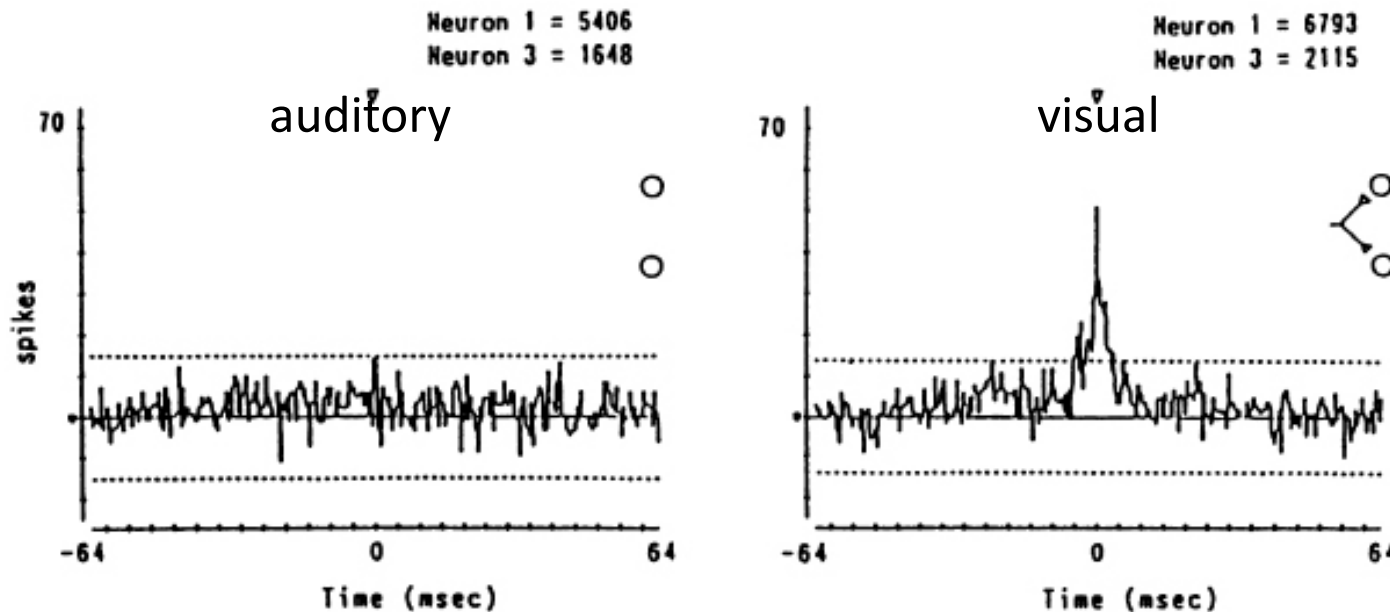
- still not fast enough for studies of plasticity
- need to increase multi-threading on compute nodes

- is self-contained benchmark application for HBP prototype system

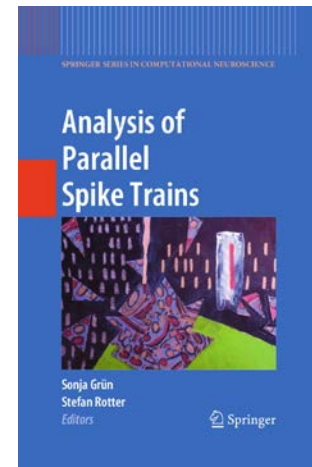
Feasibility and necessity

- Can we do simulations at the brain scale? ✓
- Do we need to simulate full scale (at cellular resolution)?

Functional Correlation




Sakurai, Y. (1999) Neuroscience & Biobehavioral Reviews 23: 785-796



- two neurons in CA1 of a rat performing an auditory or visual discrimination task
 - **cross-correlation function**: probability of neuron 2 emitting spike at delay after 1
 - task related correlation only observed for visual task
- ⇒ interpretation: neurons belong to a cell assembly processing visual information

Networks generally not reducible


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RESEARCH ARTICLE

Scalability of Asynchronous Networks Is Limited by One-to-One Mapping between Effective Connectivity and Correlations

Sacha Jennifer van Albada  Moritz Helias, Markus Diesmann

Published: September 1, 2015 • DOI: 10.1371/journal.pcbi.1004490

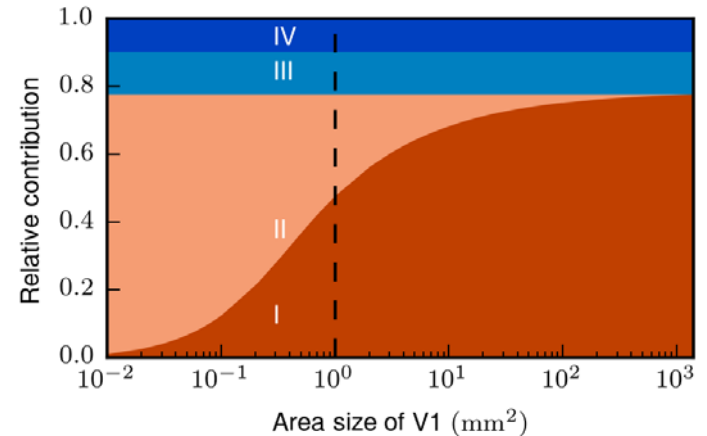
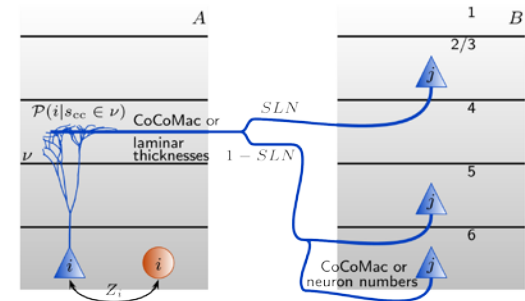
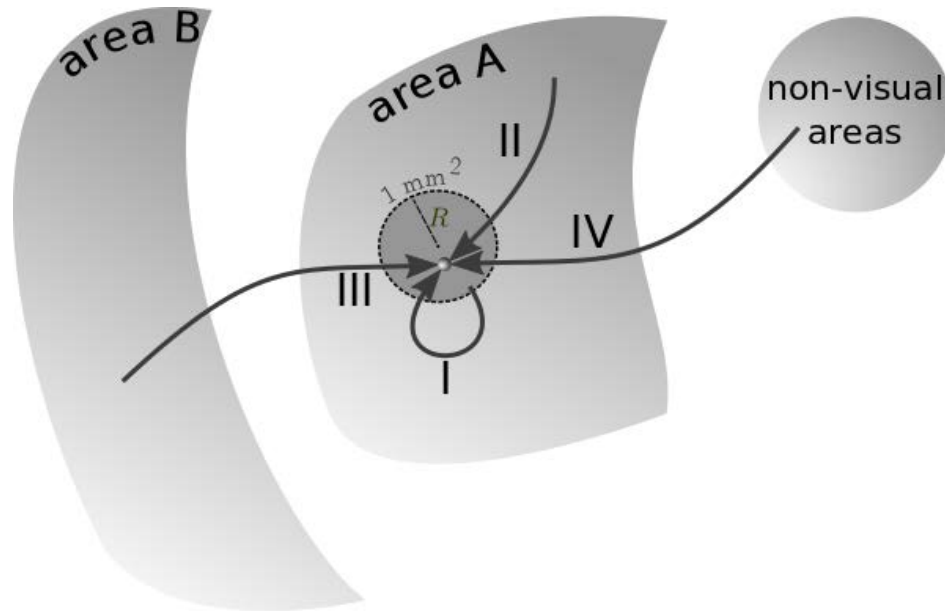
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- downscaling works well for first order statistics like spike rate
- severe constraints already for second order like spike correlation
- spike correlation drives mesoscopic measures like LFP and EEG

Feasibility and necessity

- Can we do simulations at the brain scale? ✓
- Do we need to simulate full scale (at cellular resolution)? ✓

Toward a self-consistent model

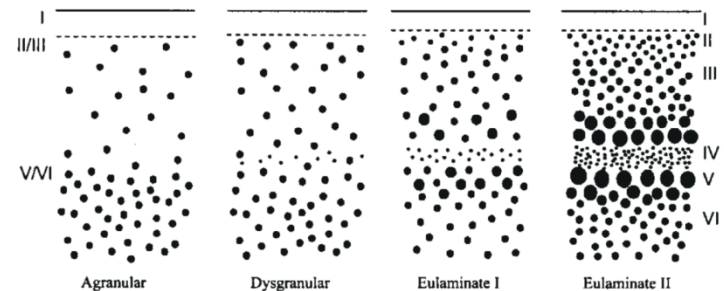
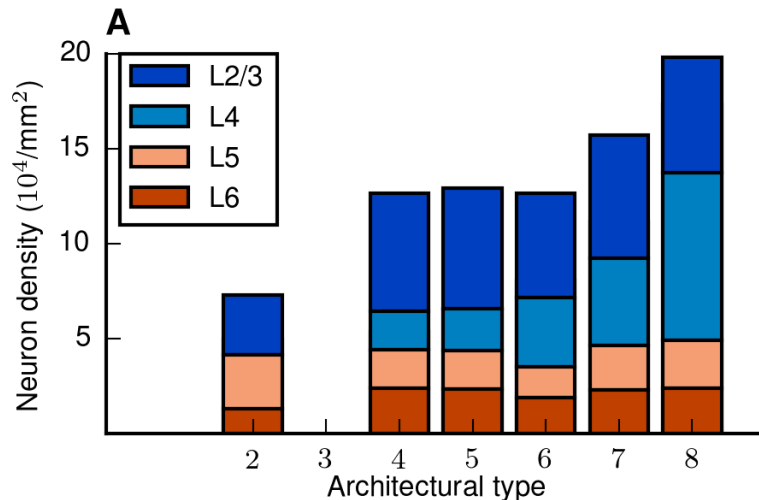


- I. Intra-areal synapses
- II. Intra-areal synapses replaced by random input
- III. Cortico-cortical synapses
- IV. External input represented by random input

- Sacha van Albada
- Maximilian Schmidt
- Rembrandt Bakker

Multi-area model of macaque visual cortex

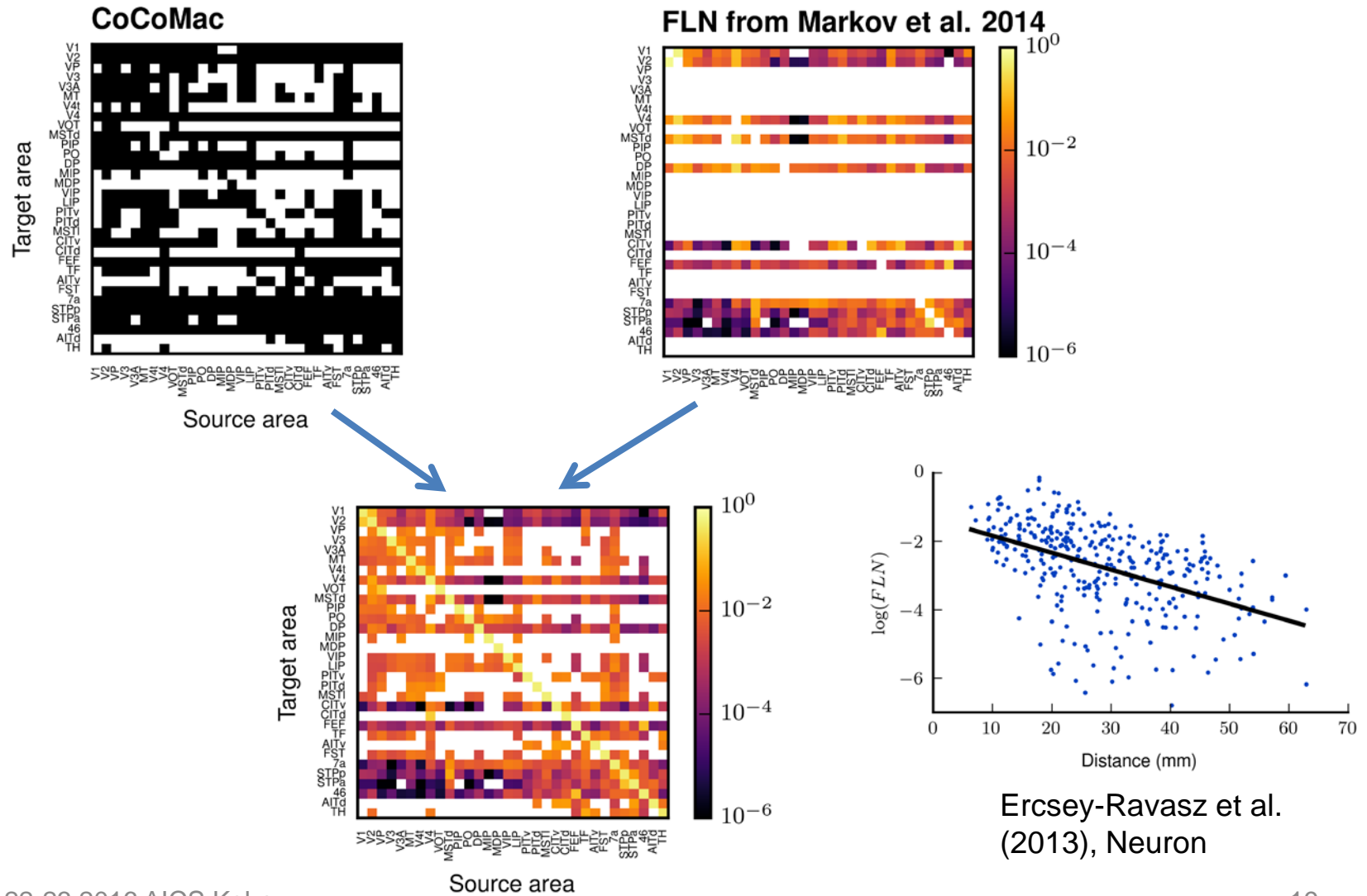
- rich anatomical data sets available (e.g CoCoMac)
- close to human
- 32 areas structured in layers comprising $8 \cdot 10^8$ neurons
- downscaled model with $4.1 \cdot 10^6$ neurons and $3.9 \cdot 10^{10}$ synapses



From Dombrowski et al. (2001), Cereb Cortex

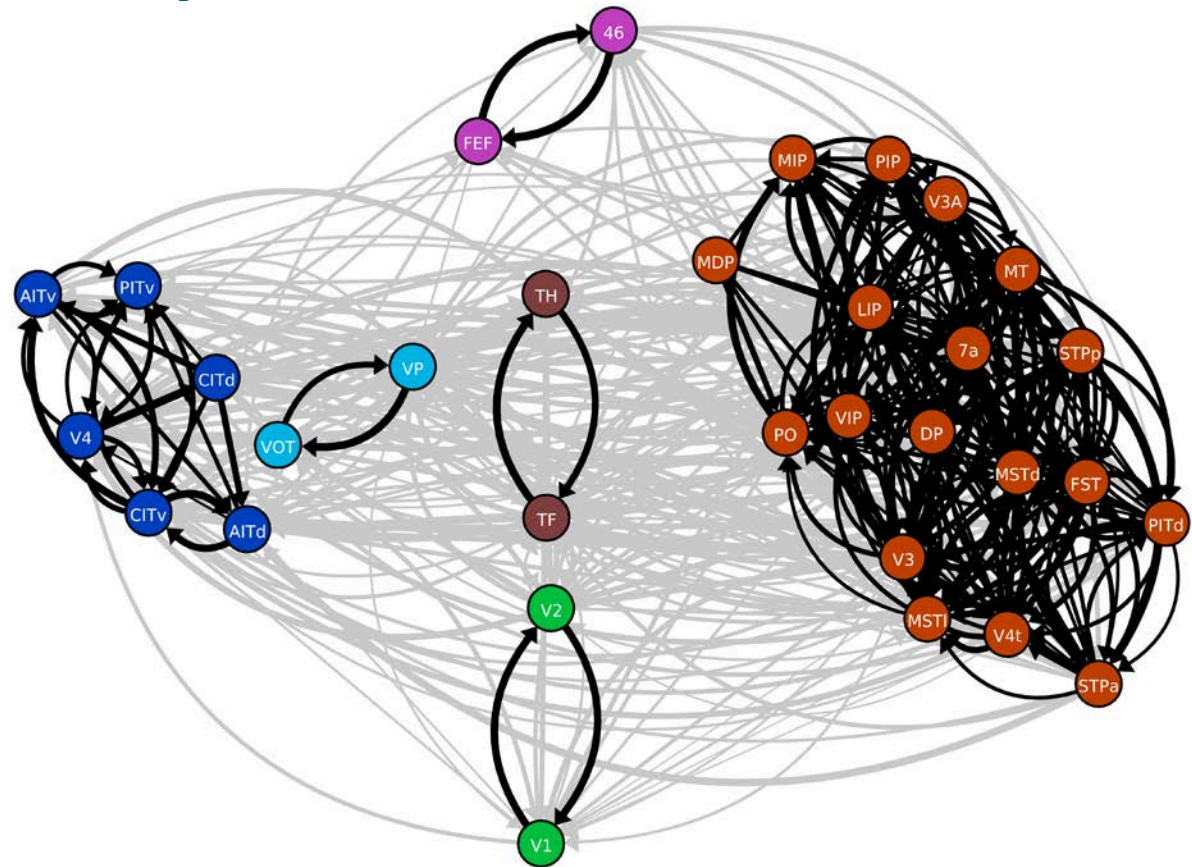
architectural types from Hilgetag et al. (2015)
with data by Helen Barbas

Construction of cortico-cortical connectivity



Ercsey-Ravasz et al. (2013), Neuron

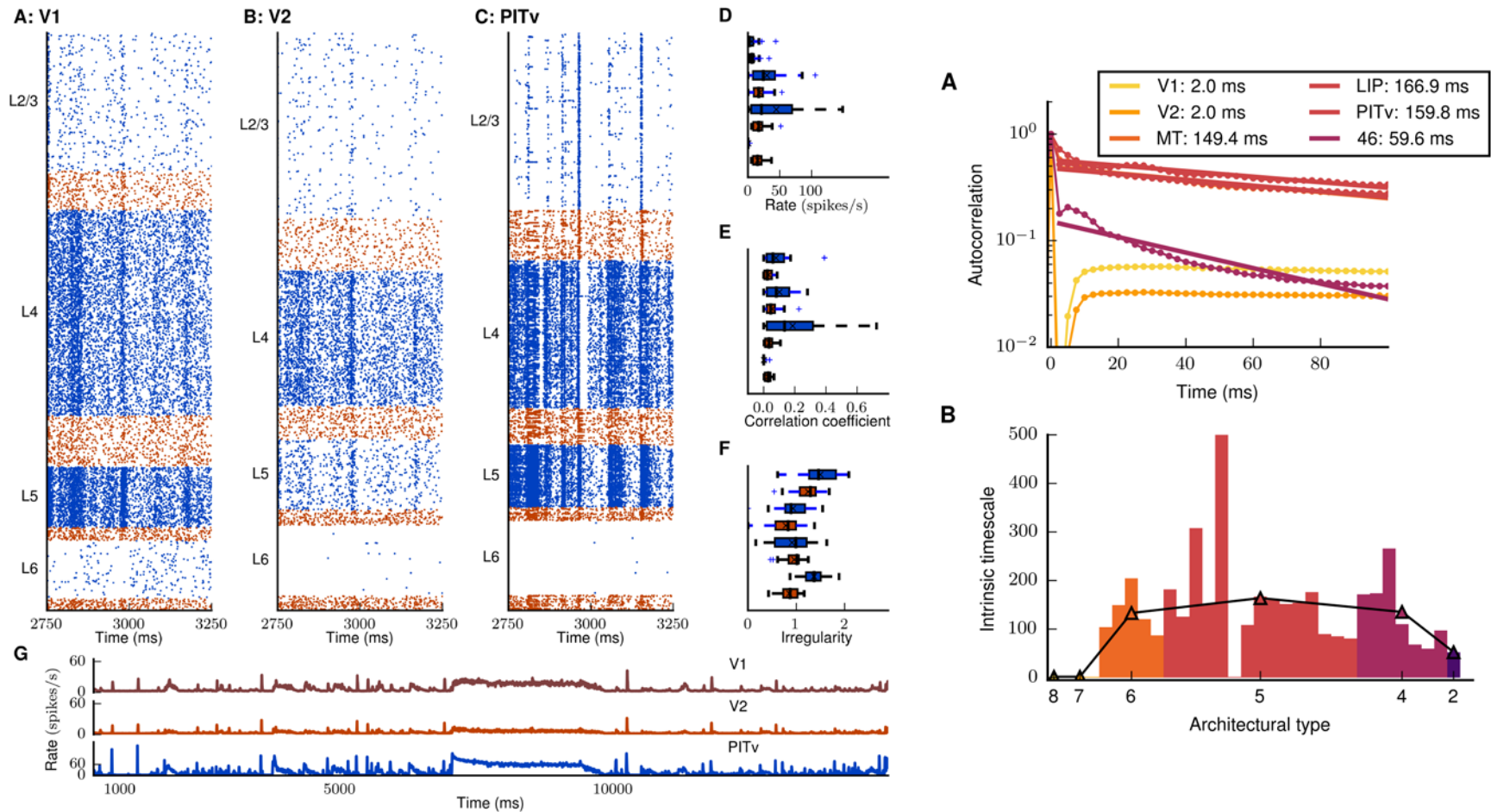
Structural connectivity reveals functionally relevant community structure



clustering by map equation method (Rosvall et al. 2010)

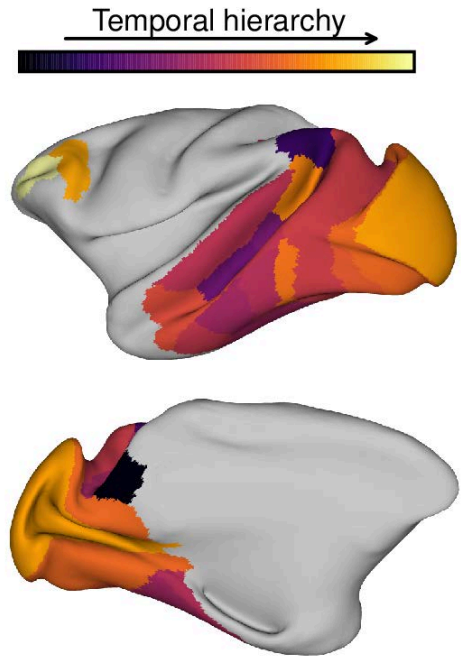
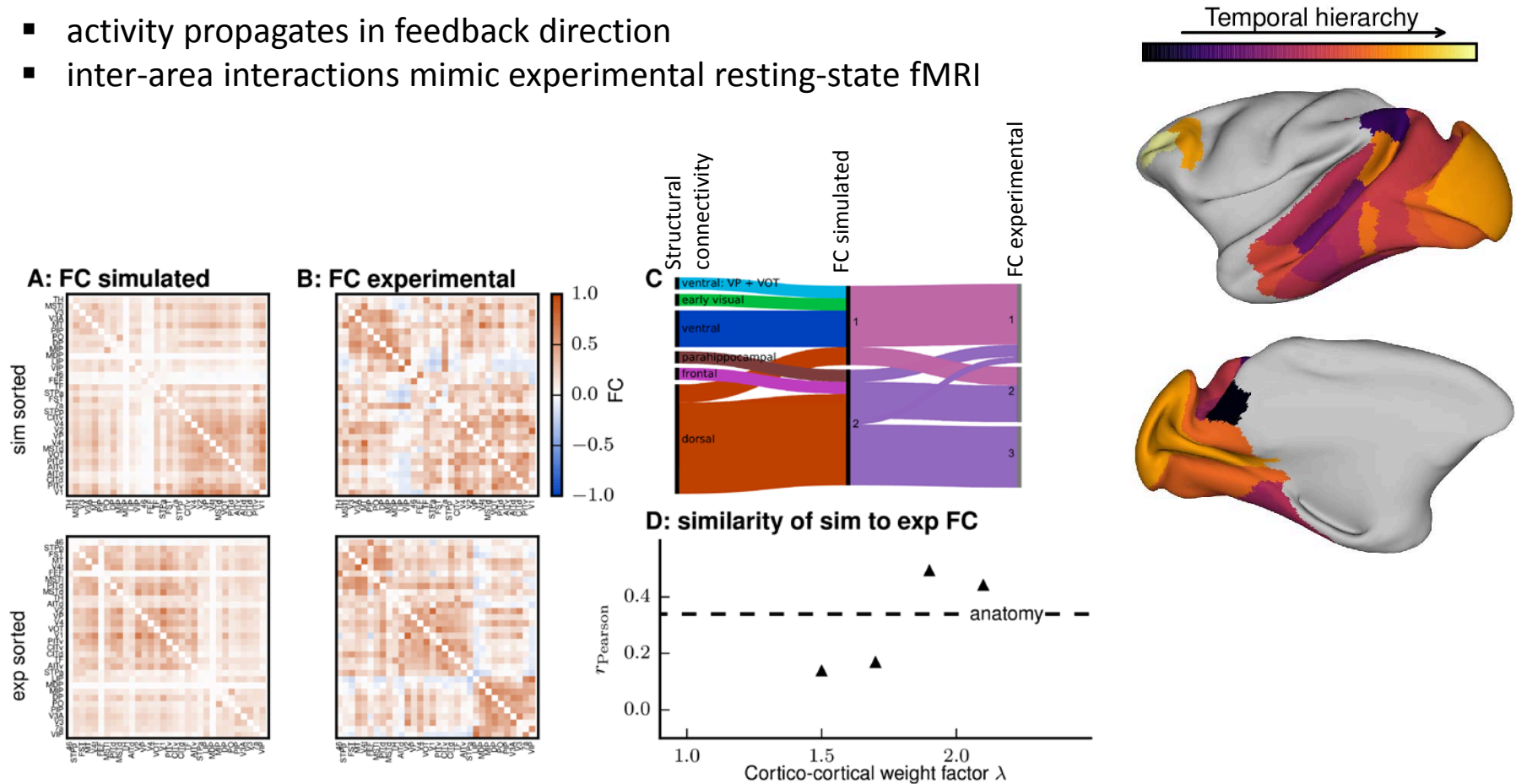
Multi-area model: Dynamical results

- stable resting state with heterogeneous laminar rate patterns and irregular firing
- cortico-cortical interactions trigger increased time scales in higher visual areas



Multi-area model: Dynamical results

- activity propagates in feedback direction
- inter-area interactions mimic experimental resting-state fMRI



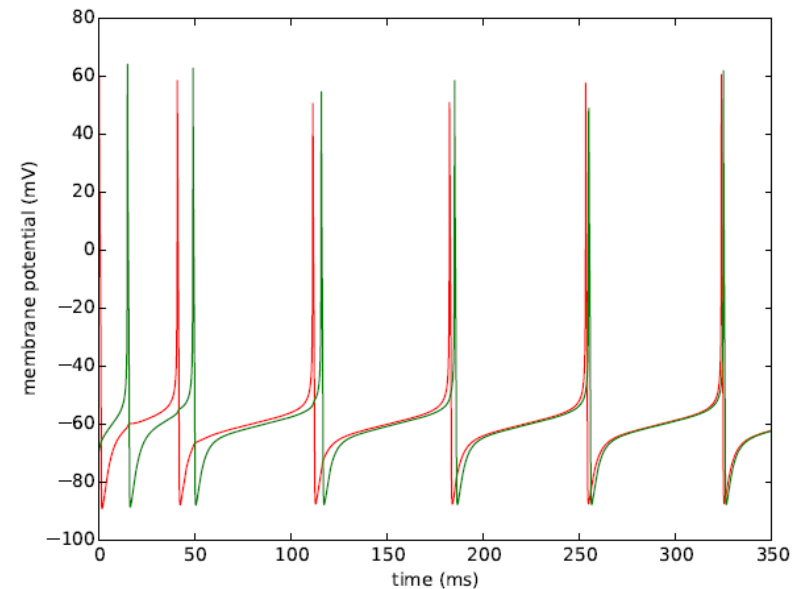
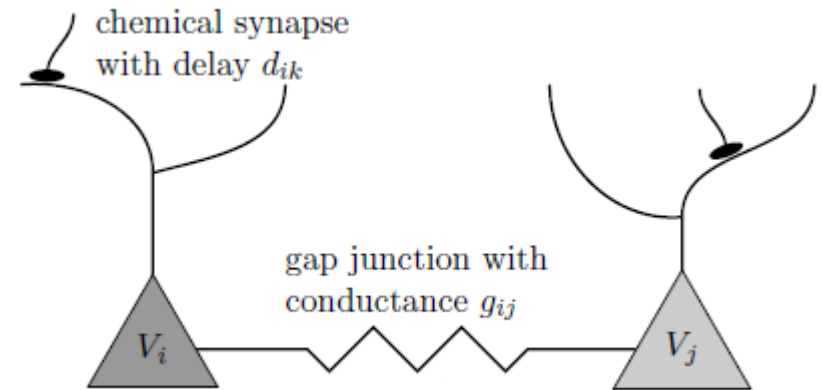
FC sorted according to Louvain clustering (Blondel et al. 2008)

Schmidt M, Bakker R, Shen K, Bezgin G, Hilgetag CC, Diesmann M, van Albada SJ (2016) arXiv:1511.09364

Biophysics of gap junctions

- electrical synapses between neurons
- widespread in nervous system
- crucial for synchronization and generation of rhythmic activity
- mechanism: instantaneous gap current

$$I_{\text{gap}}(t) = g_{ij} (V_i(t) - V_j(t))$$



time evolution of the membrane potentials of two neurons with constant current input and gap-junction coupling

- only very small networks studied so far

Gap junctions in NEST simulator

A unified framework for spiking and gap-junction interactions in distributed neuronal network simulations



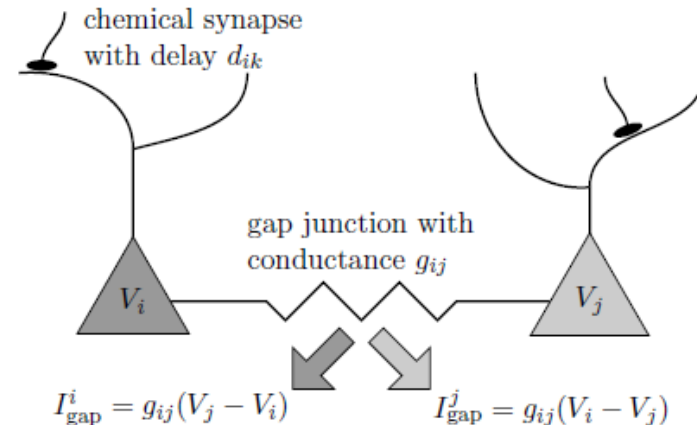
Jan Hahne^{1}, Moritz Helias^{2,3}, Susanne Kunkel^{3,4}, Jun Igarashi^{5,6}, Matthias Bolten¹, Andreas Frommer¹ and Markus Diesmann^{2,7,8}*

¹ Department of Mathematics and Science, Bergische Universität Wuppertal, Wuppertal, Germany, ² Institute of Neuroscience and Medicine (INM-6), Institute for Advanced Simulation (IAS-6), JARA BRAIN Institute I, Jülich Research Centre, Jülich, Germany, ³ Programming Environment Research Team, RIKEN Advanced Institute for Computational Science, Kobe, Japan, ⁴ Simulation Laboratory Neuroscience, Bernstein Facility for Simulation and Database Technology, Institute for Advanced Simulation, Jülich Aachen Research Alliance, Jülich Research Centre, Jülich, Germany, ⁵ Neural Computation Unit, Okinawa Institute of Science and Technology, Okinawa, Japan, ⁶ Laboratory for Neural Circuit Theory, RIKEN Brain Science Institute, Wako, Japan, ⁷ Department of Psychiatry, Psychotherapy and Psychosomatics, Medical Faculty, RWTH Aachen University, Aachen, Germany, ⁸ Department of Physics, Faculty 1, RWTH Aachen University, Aachen, Germany

- technology included in NEST 2.10.0
- collaboration of K computer and JUQUEEN teams
- benchmarking with Itaru Kitayama (AICS) and Brian Wylie (JSC Juelich)

Distributed solver for gap-junction dynamics

- instantaneous interaction couples ODE-systems of single neurons
- iterative approach based on waveform relaxation technique required
- cubic approximation of membrane potentials

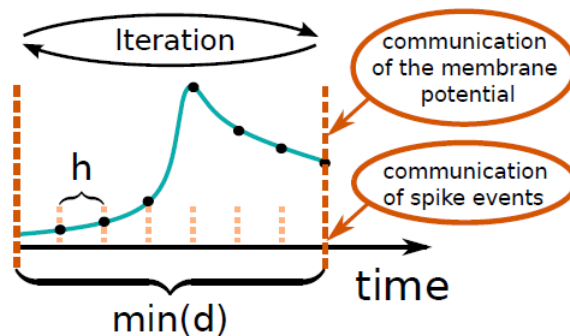


Neuron i (hh_psc_alpha_gap)

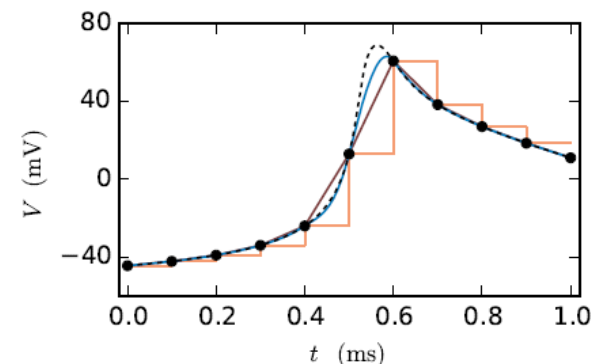
$$\frac{\dot{V}_i}{C_m} = -I_{\text{ionic}}^i(V_i, m_i, h_i, n_i, p_i) + I_{\text{applied}}^i(I_{\text{ex}}^i, I_{\text{in}}^i) + I_{\text{gap}}^i(V_i, V_j)$$

Neuron j (hh_psc_alpha_gap)

$$\frac{\dot{V}_j}{C_m} = -I_{\text{ionic}}^j(V_j, m_j, h_j, n_j, p_j) + I_{\text{applied}}^j(I_{\text{ex}}^j, I_{\text{in}}^j) + I_{\text{gap}}^j(V_j, V_i)$$



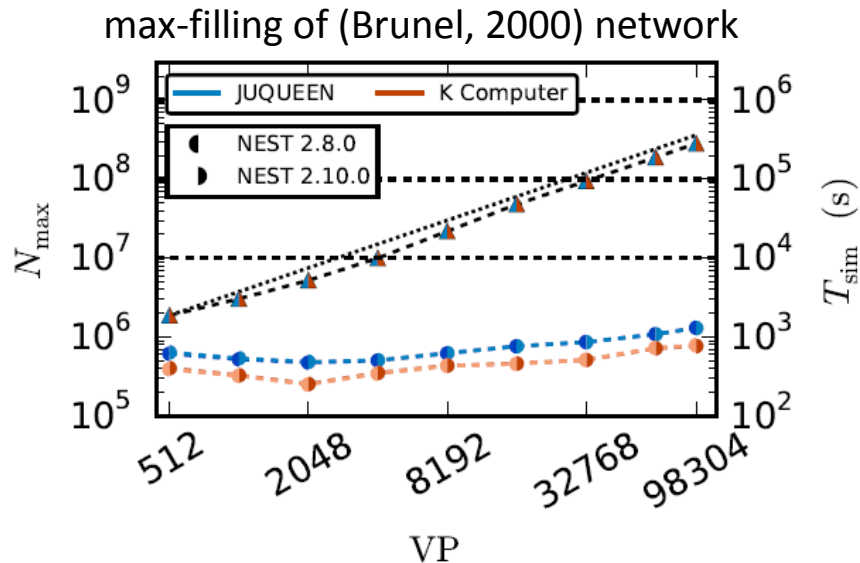
membrane potential evolution during minimal delay of spike interactions, solved repeatedly until stopping criterion is met



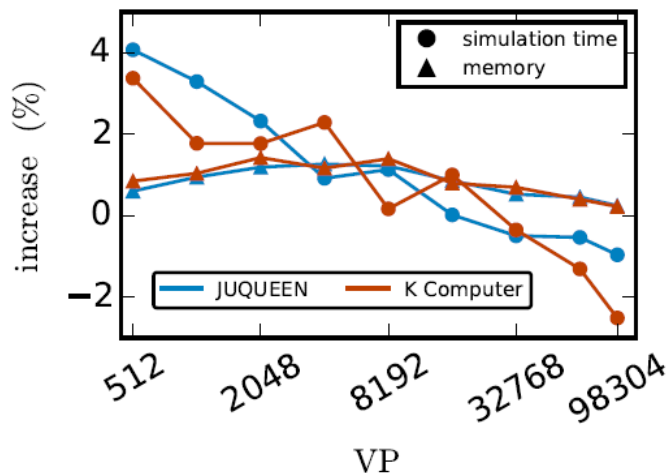
possible approximations of the membrane potential (dashed black curve) representing an action potential (spike), black dots indicate grid points

Performance of gap junction code

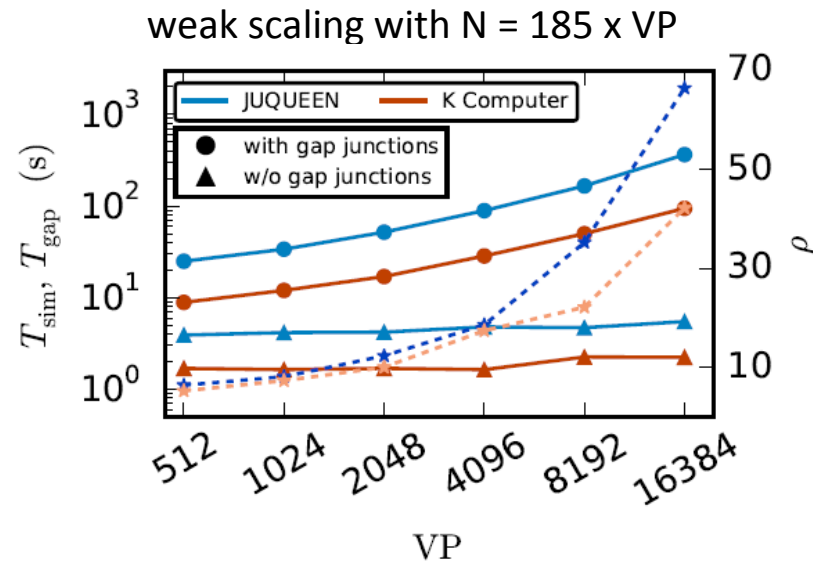
influence on simulations w/o gap junctions



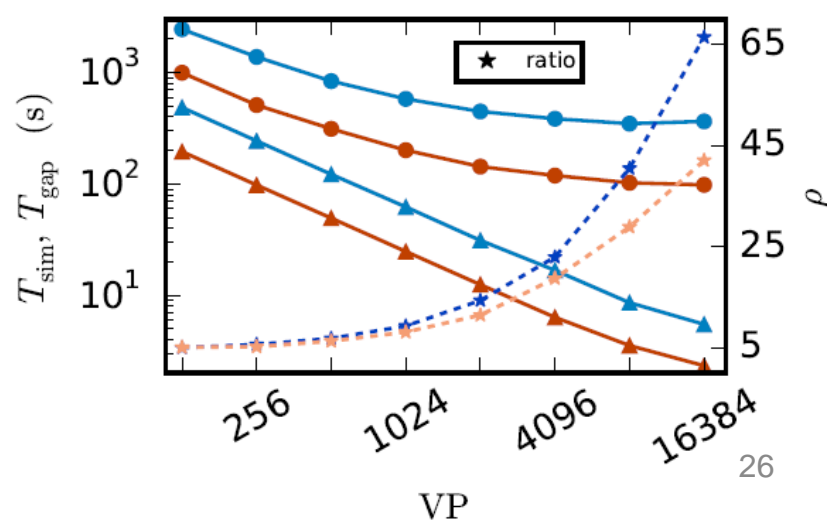
increase of memory/time in percent



costs of gap-junction dynamics



strong scaling with $N = 3,031,040$



Summary

- need for brain-scale models
 - increase self consistency
 - compute meso- and macroscopic measures of activity
- need for full-scale models
 - irreducibility of second order statistics
 - verify mean-field results
- machines ready for use by neuroscience (www.nest-initiative.org)
- K computer and JUQUEEN well suited for brain simulations
- neuroscience results for model of macaque monkey visual cortex
- biological mechanism “gap junction” (electrical synapse) available
 - hard problem due to continuous interaction
 - evaluated in MoU AICS–Jülich

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