

Reproducible Models and Replicable Implementations

Current Trends in Computational Neuroscience

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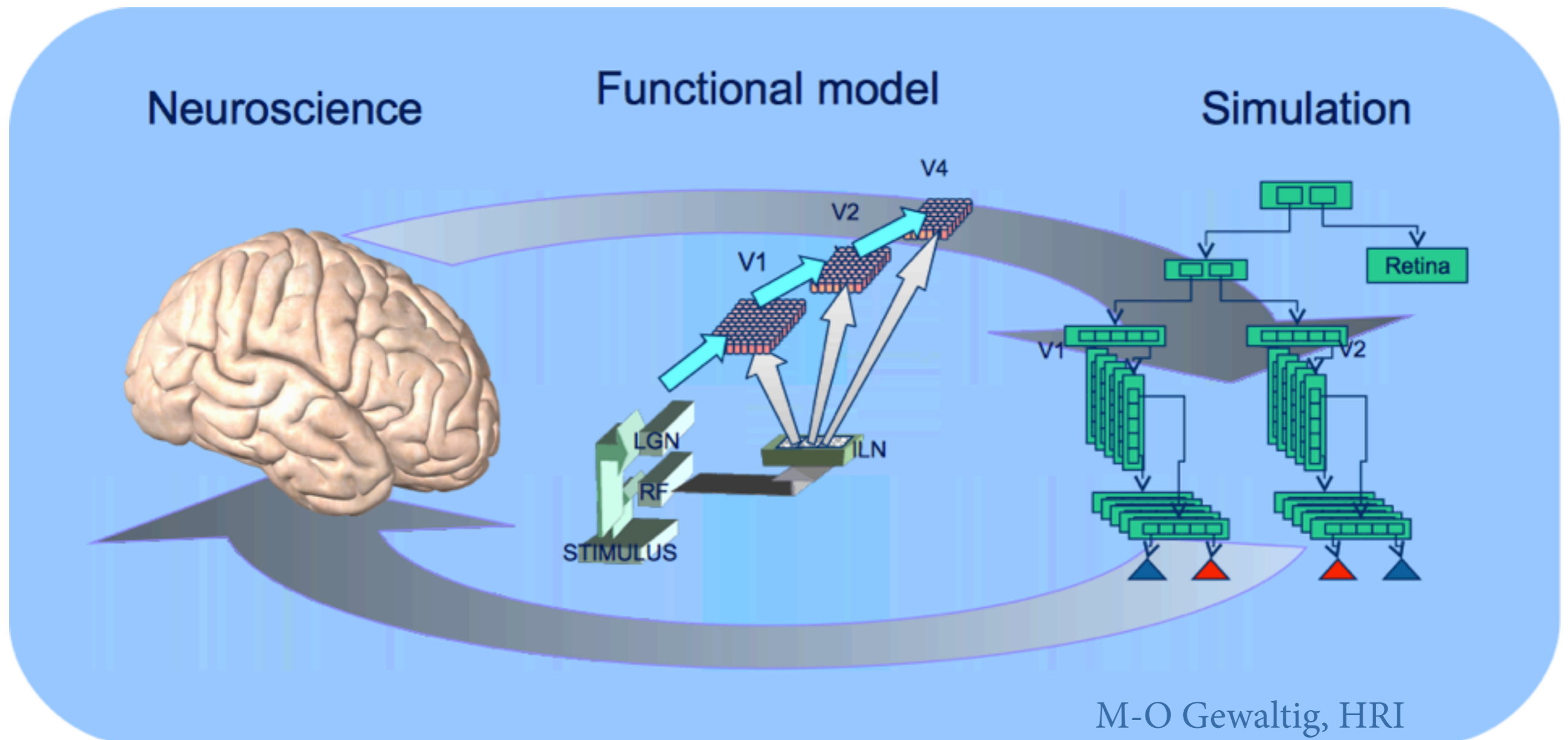
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Computational Neuroscience

What is Computational Neuroscience?

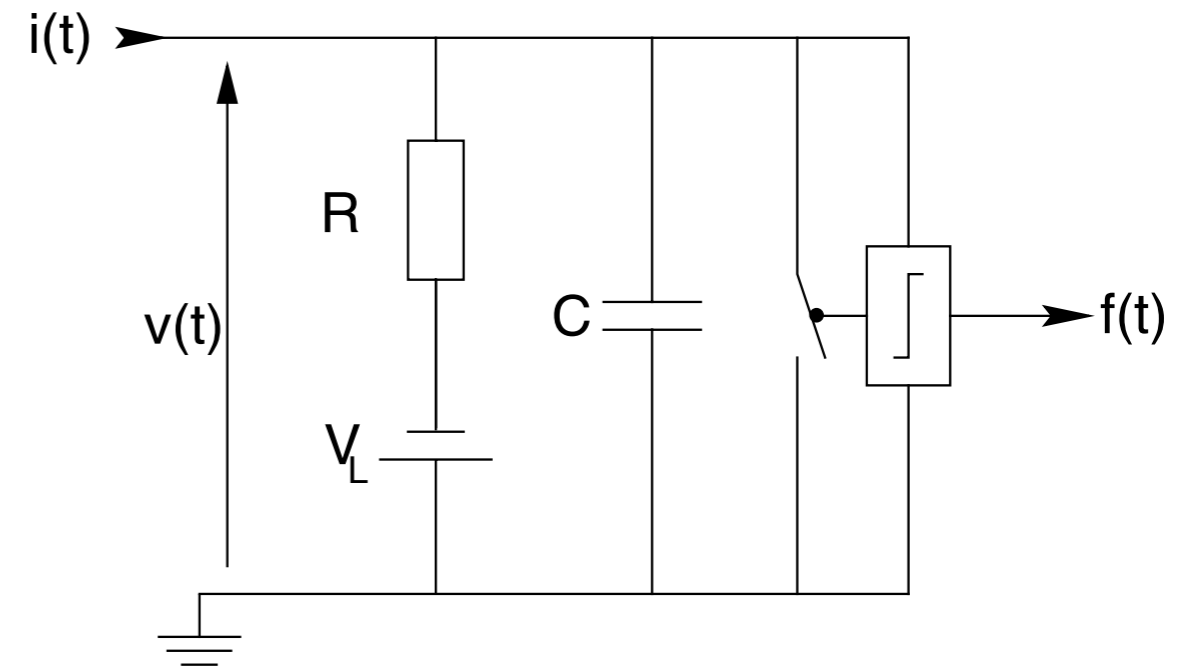
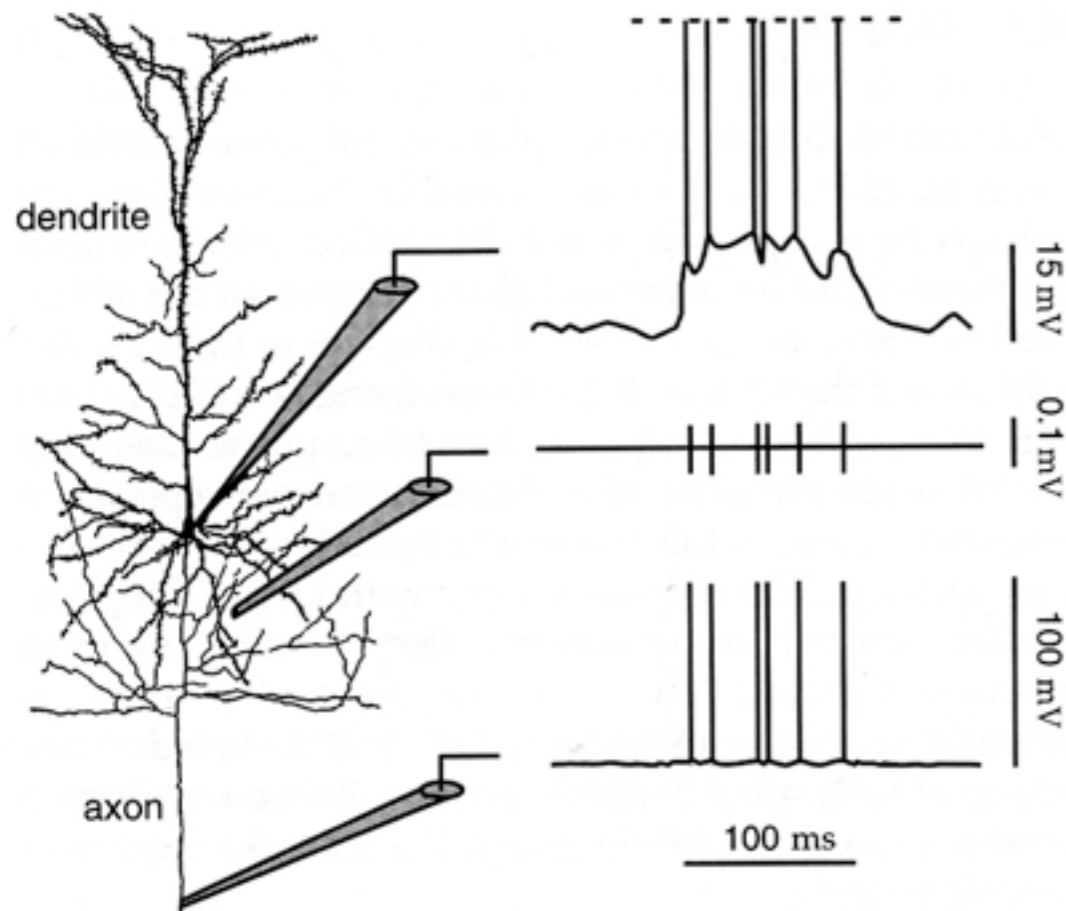


The goal of neural modeling is to relate, in nervous systems, function to structure on the basis of operation.

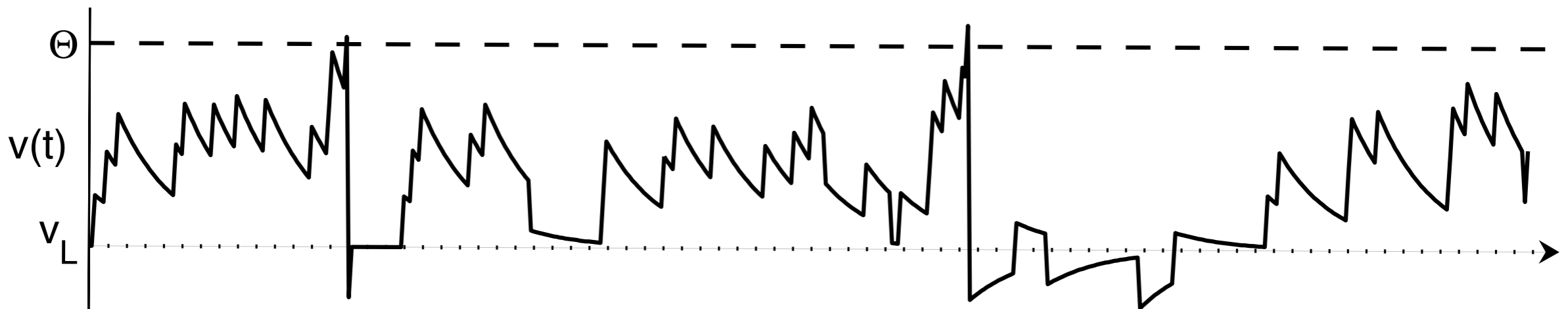
MacGregor & Lewis, 1977

Keeping it simple: point neurons

Complexity of the neuron ... abstracted as simple RC-circuit

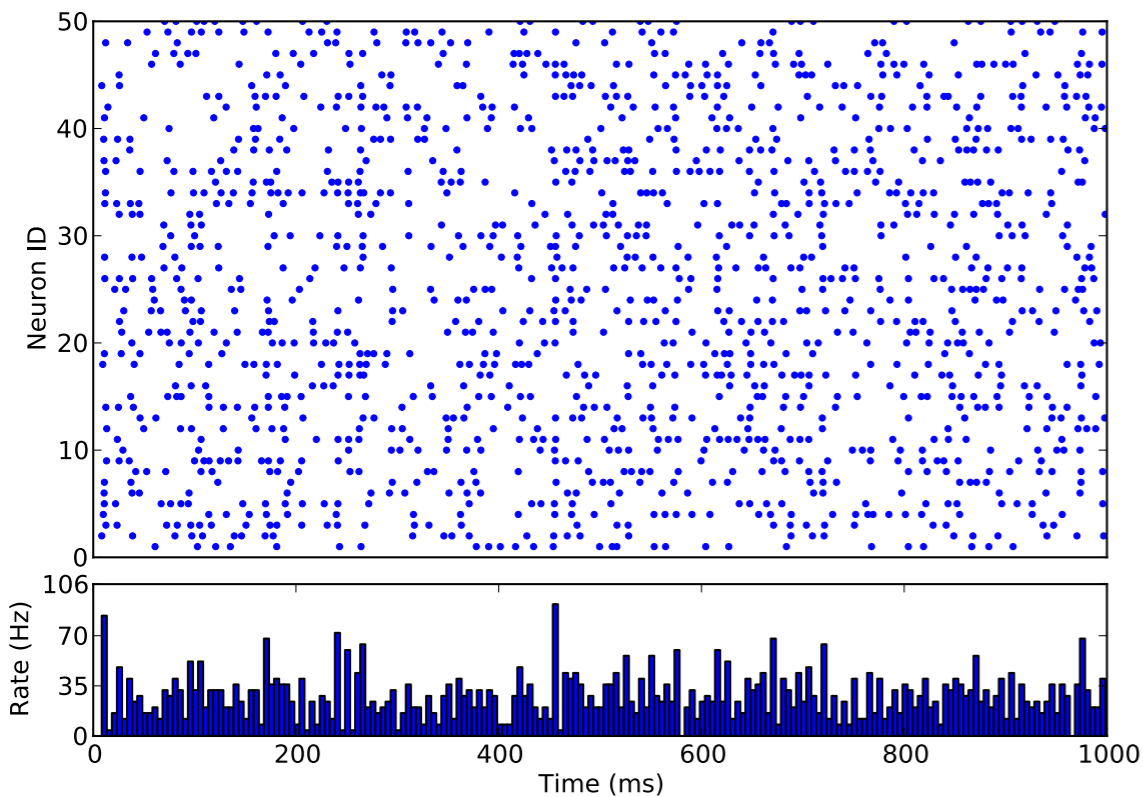
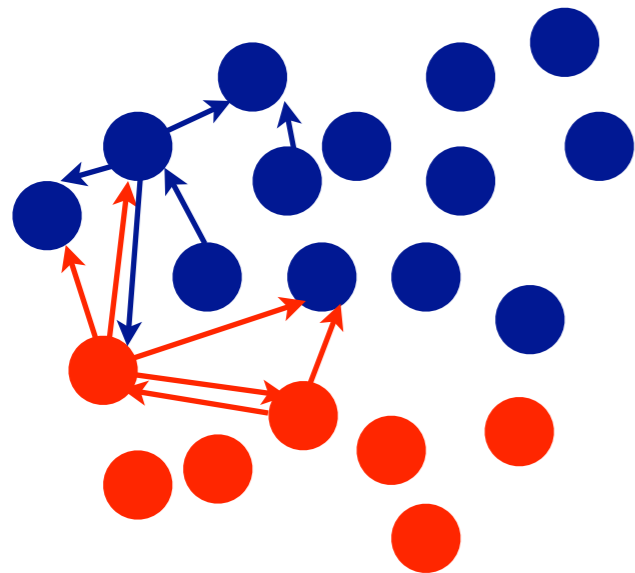


$$C\dot{V} = -\frac{(V - V_L)}{R} + i(t)$$

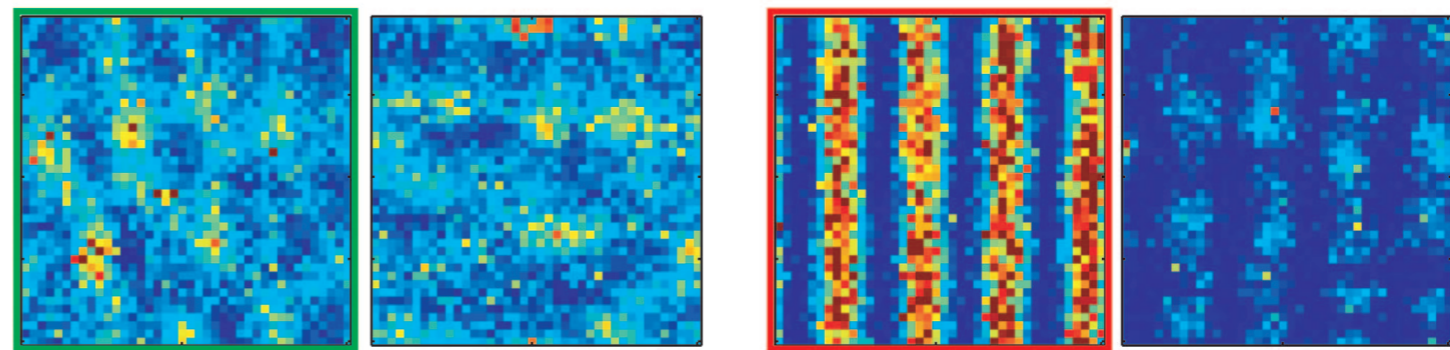
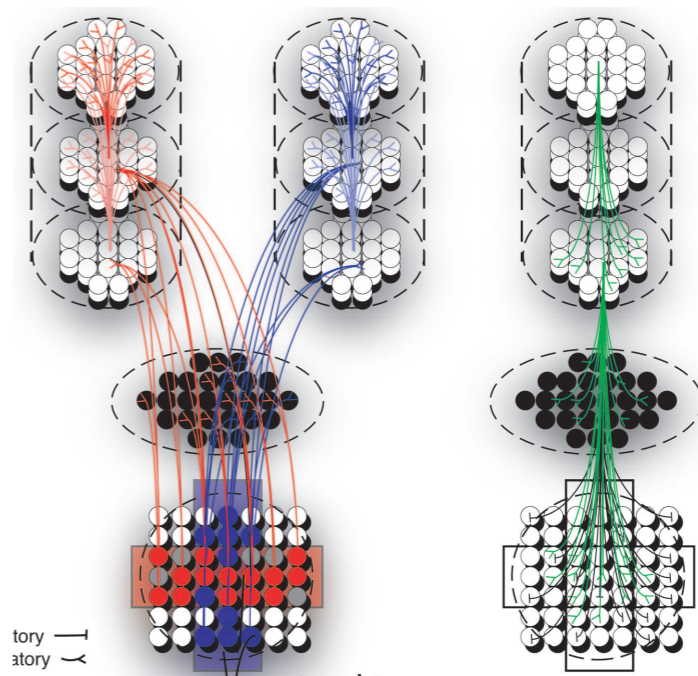


Then explore network dynamics!

Random network



Structured network



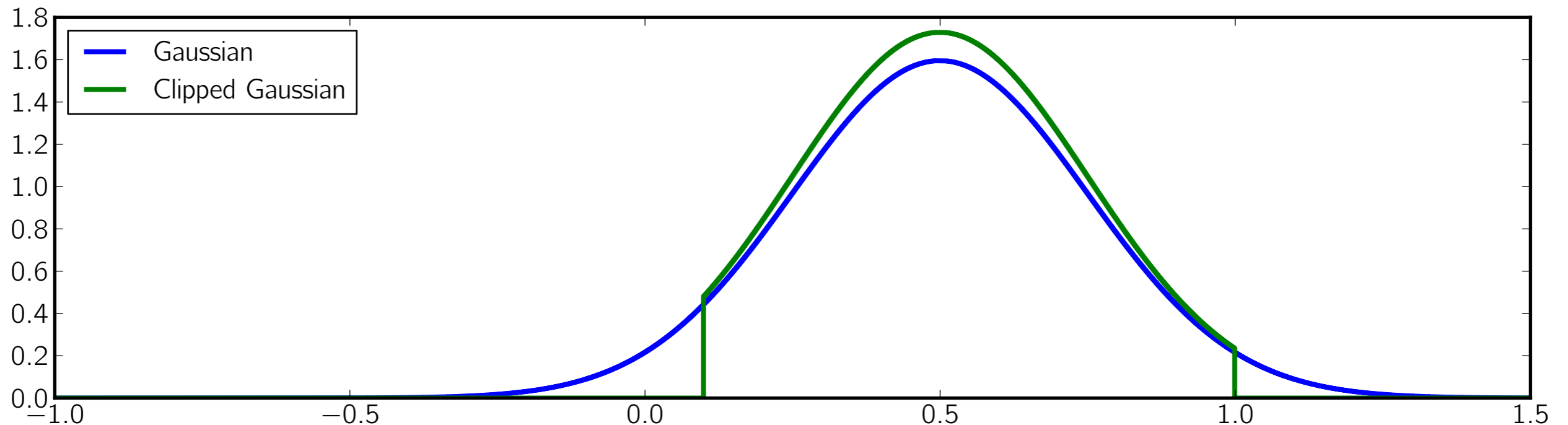
Hill & Tononi (2005)

Simulations are exciting — but reliable?

- Computational neuroscience
 - no conservation laws
 - no clear-cut separation of scales
 - no general agreement on which aspects of network activity are essential (spike rate vs spike time debate)
 - highly abstract models difficult to compare to experimental data quantitatively
- ➔ Highly dependent on *reliable* simulations
- ➔ Let's look at a real-life case

Case 2: The clipped Gaussian

- Well-known paper on plasticity
- Parameters chosen from Gaussian distribution, according to paper
- Results could not be reproduced independently
- Analysis of original C-code provided by authors:
 - ➔ Parameters were chosen from *clipped* Gaussian



The (sad) state of the art

- Few published results can be reproduced independently
- Authors often struggle to replicate their own results
- Systematic comparison and evaluation of models are rare
- Authors rarely discuss why and how models ended up as they are
- Models are seldom re-used

Reproduction *vs* Replication

Chris Drummond

Replicability is not Reproducibility: Nor is it Good Science

ICML Montreal 2009



Replication: necessary, difficult, & insufficient

- Replication
 - Re-create identical results
 - Essentially book-keeping
 - Requires tools & discipline
 - No new insights: tests implementation, not ideas
- *Internal replication*

Joe recreates results on original machine
- *External replication*

Jane recreates Joe's results on her machine using Joe's code
- *Cross replication*

Alice recreates Joe's results using a different simulator, based on a simulator-independent model description

How much detail does replication require?

- Very simple point-neuron model

$$\dot{V} = -\frac{V}{\tau} + \frac{I_E}{C}$$

- Exact updating rule ($a = I_E\tau/C$)

$$V_{k+1} = V_k e^{-h/\tau} + a(1 - e^{-h/\tau})$$

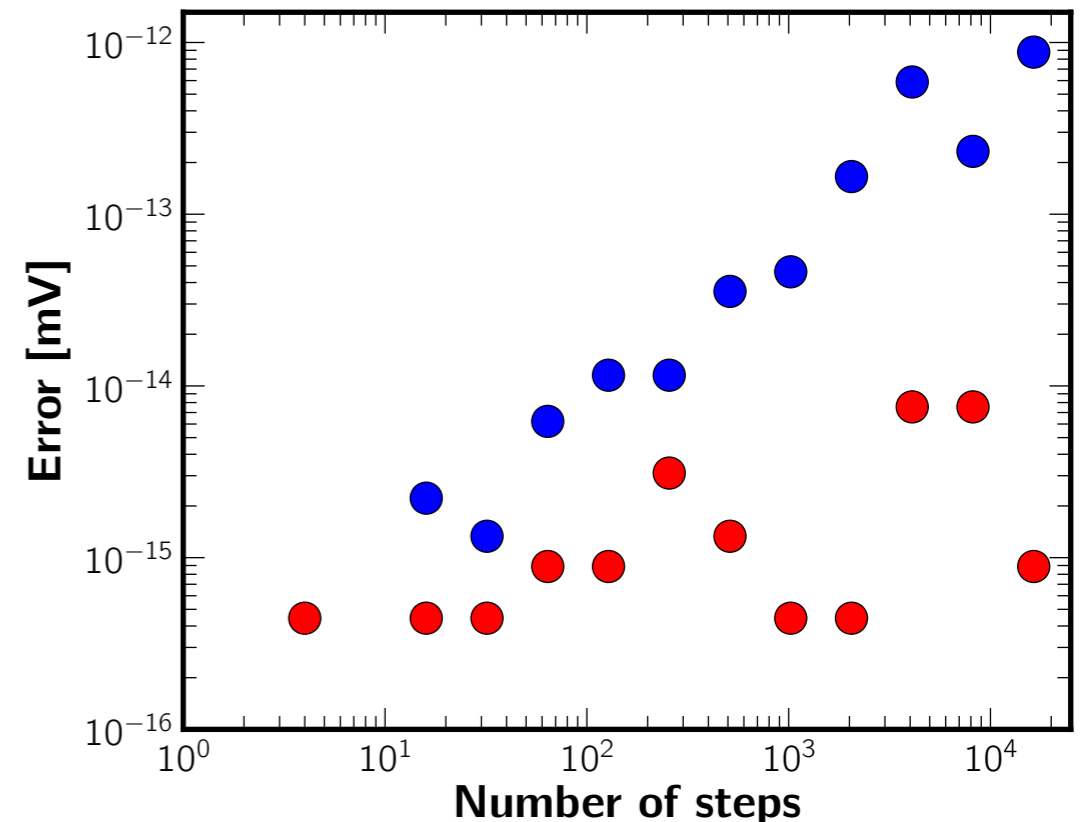
- Two different implementations

$$V[k+1] = V[k] * \exp(-h/\tau) - a * \expm1(-h/\tau)$$

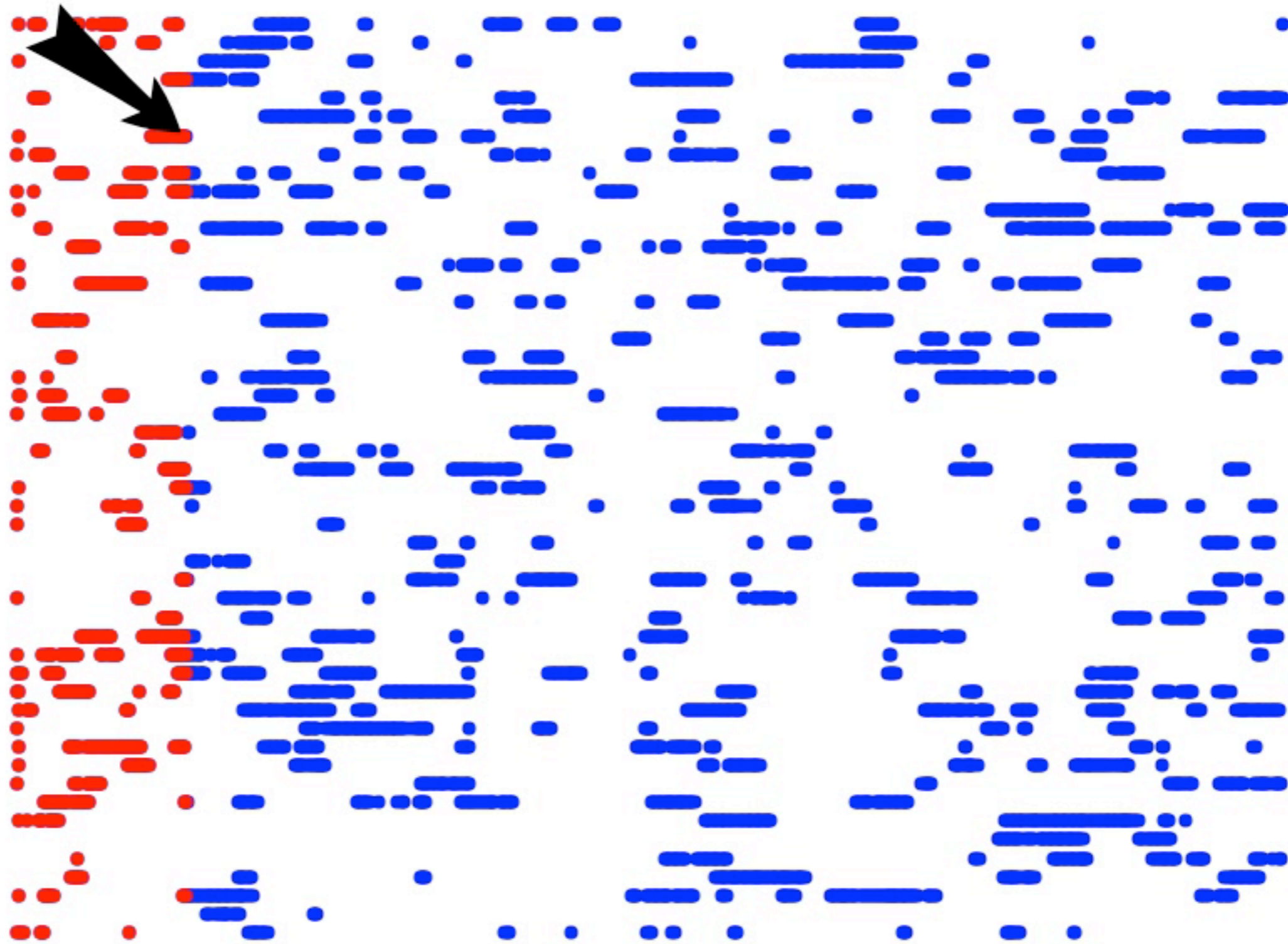
$$V[k+1] = V[k] - (a - V[k]) * \expm1(-h/\tau)$$

- Different numerical properties

➔ Replication requires that we specify implementation!



But do such tiny differences matter?

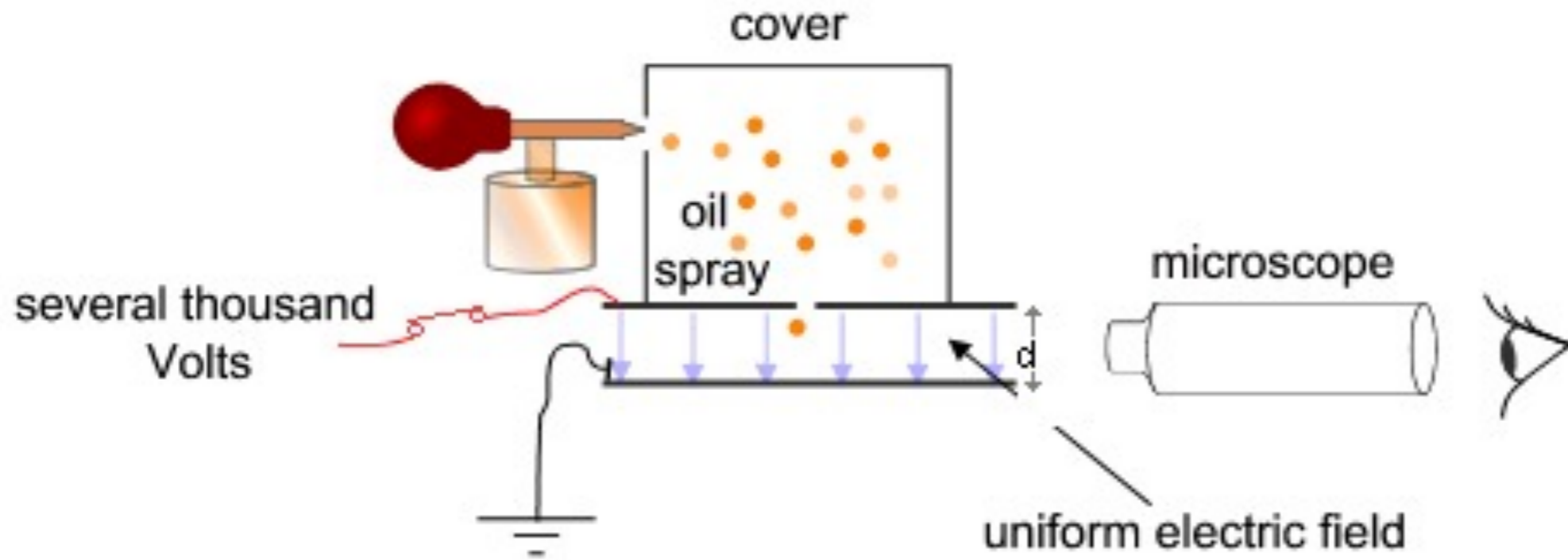


Reproduction

- Independent
- Test hypotheses and models
- Validates concepts
- Requires reflection

From oil drops to first-passage times

Millikan's oil drop experiment



Wikipedia

Erwin Schrödinger

Zur Theorie der Fall- und Steigversuche an Teilchen mit Brownscher Bewegung

Physikalische Zeitschrift **16**:289 (1915)

Reproduction needs replication

- Reproducible models describe scientific ideas
 - Independent reproduction will generally fail to replicate original results precisely
 - Requires learned judgement of discrepancies
 - Requires means to understand failure
- ➔ Replicable implementation

Improving Scientific Practice in Computational Neuroscience

Good model description practice

- Systematic approach to model description in papers
- Standardize tables/checklists
- Standards for graphic representation of models
- Nordlie, Gewaltig, & Plesser
PLoS Comp Biol 5:e1000456 (2009)

A Model Summary	
Populations	Three: excitatory, inhibitory, external input
Topology	—
Connectivity	Random convergent connections
Neuron model	Leaky integrate-and-fire, fixed voltage threshold, fixed absolute refractory time (voltage clamp)
Channel models	—
Synapse model	α -shaped current inputs
Plasticity	—
Input	Independent fixed-rate Poisson spike trains to all neurons (during initial stimulation period)
Measurements	Spike activity

B Populations		
Name	Elements	Size
E	laf neuron	$N_E = 4N_I$
I	laf neuron	N_I
E _{ext}	Poisson generator	$C_E(N_E + N_I)$

C Connectivity			
Name	Source	Target	Pattern
EE	E	E	Random convergent $C_E \rightarrow 1$, weight J , delay D
IE	E	I	Random convergent $C_E \rightarrow 1$, weight J , delay D
EI	I	E	Random convergent $C_I \rightarrow 1$, weight $-gJ$, delay D
II	I	I	Random convergent $C_I \rightarrow 1$, weight $-gJ$, delay D
Ext	E _{ext}	E \cup I	Non-overlapping $C_E \rightarrow 1$, weight J , delay D

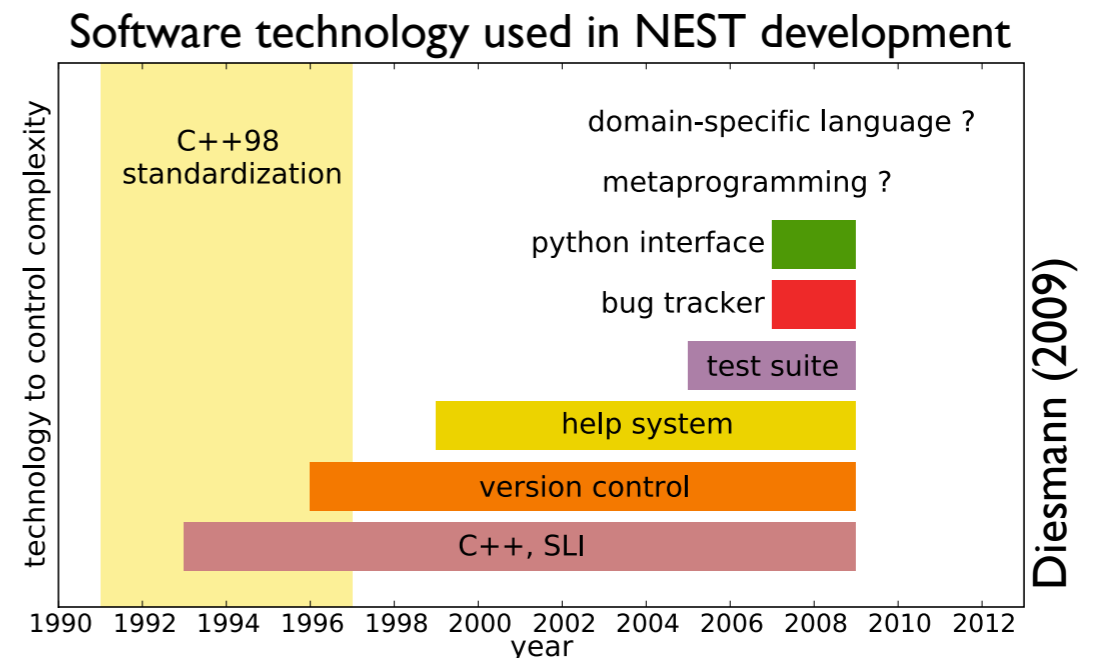
D Neuron and Synapse Model	
Name	laf neuron
Type	Leaky integrate-and-fire, α -current input
Subthreshold dynamics	$\tau \dot{V}(t) = -V(t) + RI(t) \quad \text{if } t > t^* + \tau_{rp}$ $V(t) = V_r \quad \text{else}$ $I(t) = \frac{1}{R} \sum_{\tilde{t}} w \alpha(t - (\tilde{t} + \Delta)) \Theta(t - (\tilde{t} + \Delta))$
Spiking	If $V(t-) < \theta \wedge V(t+) \geq \theta$ <ol style="list-style-type: none"> 1. set $t^* = t$ 2. emit spike with time-stamp t^*

E Input	
Type	Description
Poisson generators	Fixed rate ν_{ext} , C_E generators per neuron, each generator projects to one neuron; active only during initial stimulation period

F Measurements	
Spike activity as raster plots for subset of excitatory neurons	

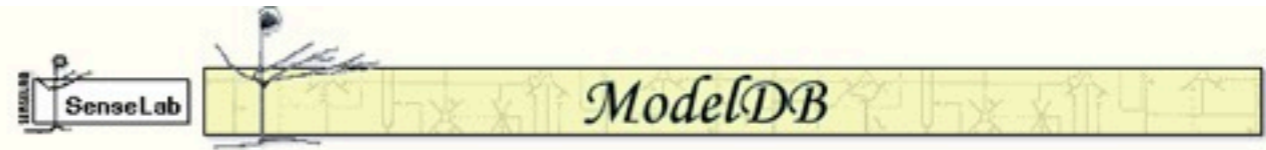
Professional, shared software

- Widely used packages replace homemade *ad hoc* code
- Currently: Neuron, NEST, Genesis, Moose, Brian, PCSim
- “Social” developments
 - Simulator review (Brette *et al*, 2007)
 - Teaching software at summer schools
 - Large-scale *scientific* projects (eg EU FACETS)
 - Neuroinformatics journals
 - Raising awareness among reviewers and editors
- “Technical” developments
 - Version control
 - Test suites



ModelDB: Sharing models

- Curated database of computational neuroscience models
- Only published models
- Open to any software
- Nearly 700 models
- <http://senselab.med.yale.edu/modeldb/>



Sparsely connected networks of spiking neurons (Brunel 2000)

Accession: 42020

The dynamics of networks of sparsely connected excitatory and inhibitory integrate-and-fire neurons are studied analytically (and with simulations). The analysis reveals a rich repertoire of states, including synchronous states in which neurons fire regularly; asynchronous states with stationary global activity and very irregular individual cell activity; and states in which the global activity oscillates but individual cells fire irregularly, typically at rates lower than the global oscillation frequency. See paper for more and details.

Reference: Brunel N (2000) Dynamics of sparsely connected networks of excitatory and inhibitory spiking neurons. *J Comput Neurosci* 8:183-208 [PubMed]

Citations [Citation Browser](#)

Model Information (Click on a link to find other models with that property)

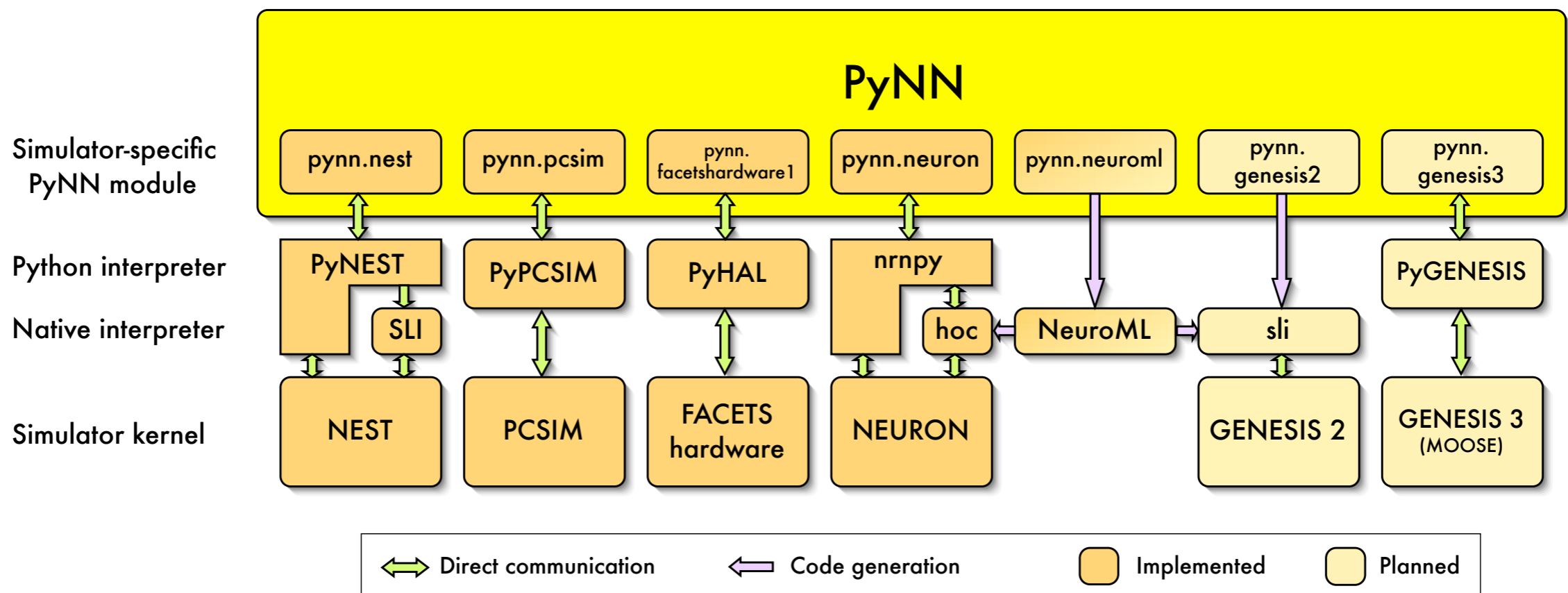
Model Type: [Connectionist Network](#);
Brain Region(s)/Organism:
Cell Type(s):
Channel(s):
Gap Junctions:
Receptor(s):
Gene(s):
Transmitter(s):
Simulation Environment: [NEST \(formerly BLISS/SYNOD\)](#);
Model Concept(s): [Activity Patterns](#); [Oscillations](#); [Spatio-temporal Activity Patterns](#); [Simplified Models](#);
Implementer(s): [Gewaltig, Marc-Oliver](#);

Model files [Download zip file](#) [Help downloading and running models](#)

<ul style="list-style-type: none">└─ \└─ brunel└─ readme.txt└─ brunel.sli	<p>Readme.txt for an implementation of the model associated with the paper:</p> <p>Brunel N (2000) Dynamics of sparsely connected networks of excitatory and inhibitory spiking neurons. <i>J Comput Neurosci</i> 8:183-208</p> <p>The brunel.sli file was supplied by Marc-Oliver Gewaltig and runs under NEST. Please contact Marc-Oliver Gewaltig marc-oliver.gewaltig@honda-ri.de for more information.</p>
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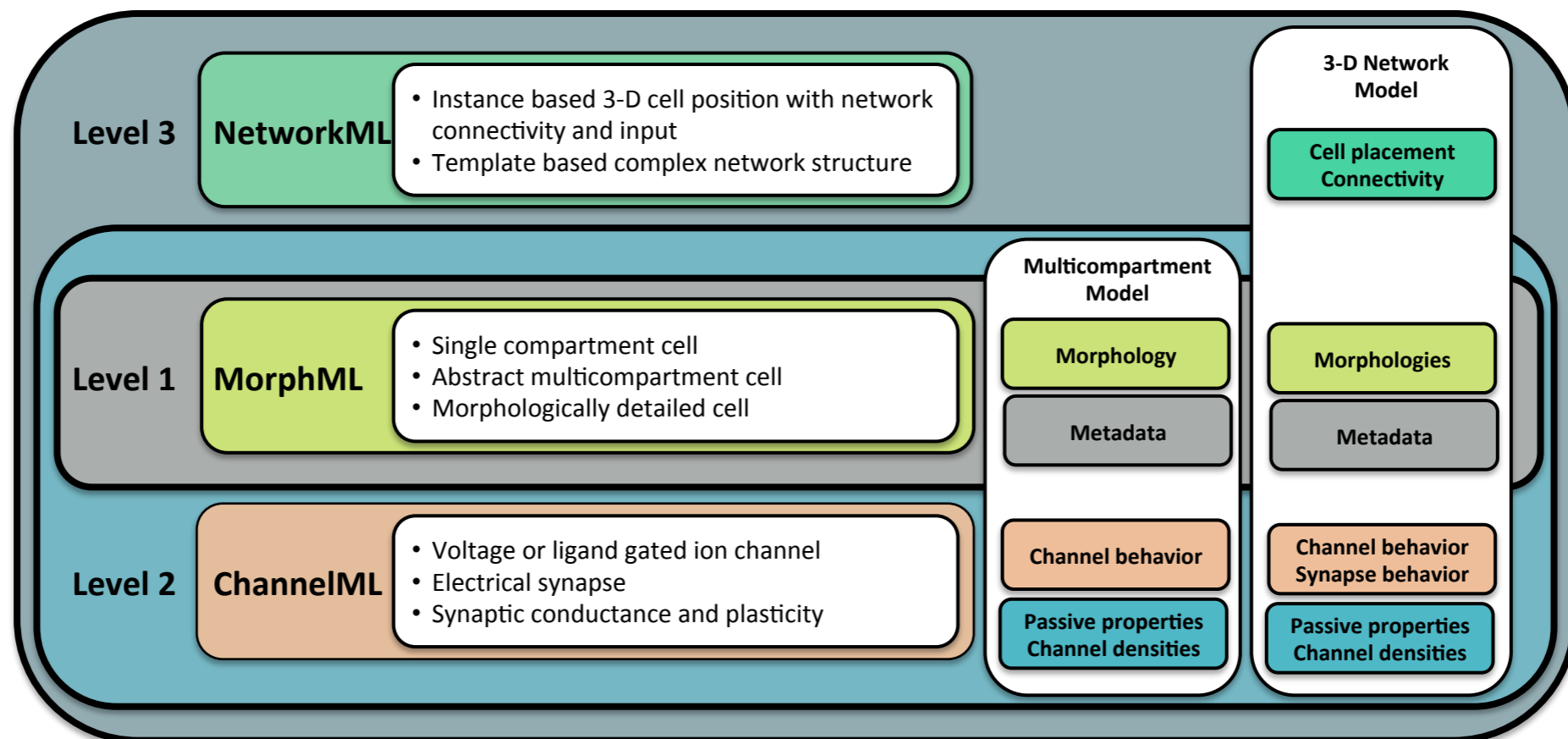
PyNN: A Meta-Simulator

- Python-based wrapper for many simulators
- Write model and simulation code once, run on all
- Facilitates model sharing and cross-validation
- Developed by Andrew Davison for FACETS project
- <http://neuralensemble.org/PyNN>



NeuroML: A model specification language

- XML-based language for model specification
- Multiple layers: channels, neuron morphologies, networks
- Code-generation for several simulators, including PyNN
- Facilitates model sharing and re-use
- <http://www.neuroml.org>



Provenance tracking: Sumatra

- Python package to enable systematic capture of the environment of numerical simulations/analyses
- Tracks simulation code, dependencies, platform information, results
- Developed by Andrew Davison as part of FACETS project
- <http://neuralensemble.org/sumatra>

Delete include data <input type="checkbox"/>	Label	Reason	Outcome	Duration	Processes	Simulator		Script			Date	Time	Tags
						Name	Version	Repository	Main file	Version			
<input type="checkbox"/>	20100709-154255		'Eureka! Nobel prize here we come.'	0.59 s		Python	2.5.2	/Users/andrew/tmp/SumatraTest	main.py	396c2020ca50	09/07/2010	15:42:55	
<input type="checkbox"/>	20100709-154309			0.59 s		Python	2.5.2	/Users/andrew/tmp/SumatraTest	main.py	396c2020ca50	09/07/2010	15:43:09	
<input type="checkbox"/>	haggling	'determine whether the gourd is worth 3 or 4 shekels'	'apparently, it is worth NaN shekels.'	0.59 s		Python	2.5.2	/Users/andrew/tmp/SumatraTest	main.py	396c2020ca50	09/07/2010	15:43:20	foobar

NineML: A model description standard

- Aims for community standard for declarative model descriptions
- Inspired by SBML and CellML
- Focus on networks of point neurons
- Under development by INCF Multi-scale Modeling Task Force

Perspectives

- Community increasingly aware of need for reproducibility and replicability
 - Large-scale projects have led to development of valuable tools
 - Summer schools educate PhD-students and post-docs in use of established modeling tools
 - Neuroinformatics journals allow publication of domain-specific solutions
 - International Neuroinformatics Coordinating Facility (INCF) stimulates debate and development
 - NEST Initiative is devoted to furthering reliable simulations
- ➔ We have a long way to go, but we are (finally) moving!

Collaborators



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Arizona State University



Andrew Davison

Unité de neuroscience, information et
complexité
CNRS
Gif-sur-Yvette



Eilen Nordlie



Marc-Oliver
Gewaltig

Honda Research Institute Europe
Offenbach

Research Council of Norway
(eVita program)

