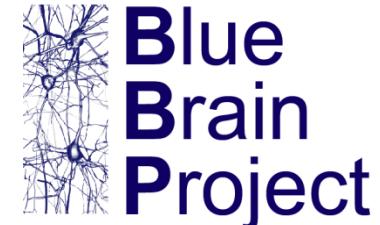


BRAIN MIND INSTITUTE **BMI**



ÉCOLE POLYTECHNIQUE
FÉDÉRALE DE LAUSANNE



Human Brain Project resources for the integrative modeling community



Human Brain Project

Eilif Muller
eilif.mueller@epfl.ch
<http://bluebrain.epfl.ch>

“At any large neuro-science meeting, one is struck by the pace of discovery, with **50,000 or more practitioners heading away from each other in all directions, in a sort of scientific Big Bang**. Although this independence is necessary, it has prevented neuroscience from entering a more mature phase, which would involve developing common standards and collaborative projects.

Neurophysiologists are more likely to use each other’s toothbrushes than each other’s data and software [and models].

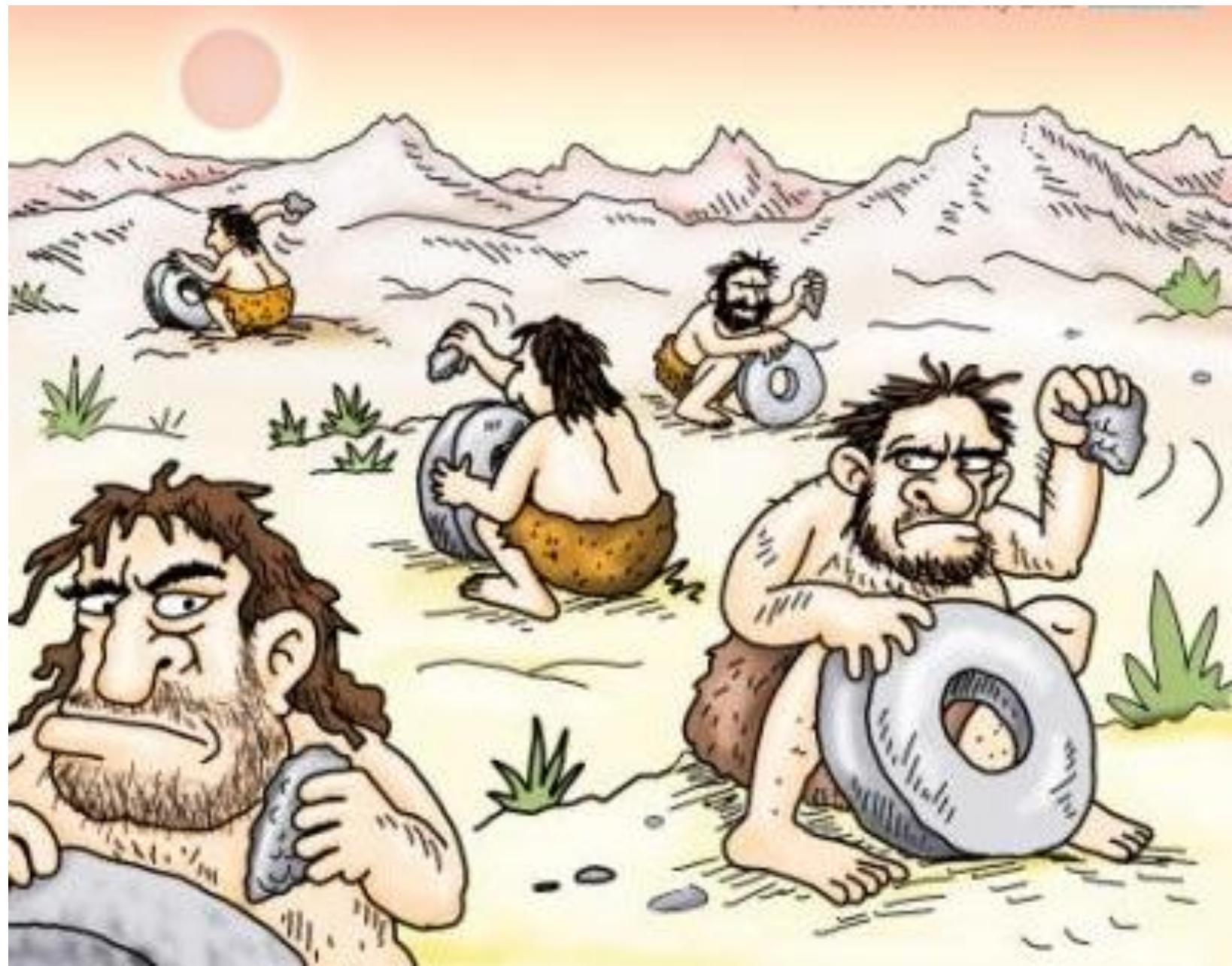
Koch and Reid, 2012

“We find that the major obstacle that hinders our understanding the brain is the fragmentation of brain research and the data it produces.

Our most urgent need is thus a concerted international effort that can integrate this data in a unified picture of the brain as a single multi-level system...”

The HBP-PS Consortium Report 2012:8

<https://goo.gl/xjjoSB>



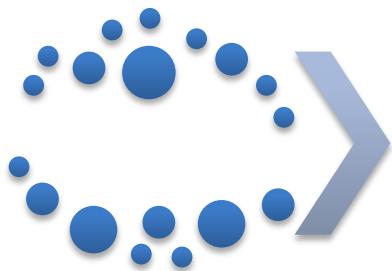
A Whole Community Approach

- Challenge is to bring experimental and theoretical & computational neuroscience closer.
- While the latter seek minimal models, the former want hard earned experimental facts not to be ignored adhoc.
- Adhoc simplification is a matter of taste.

" New databases can be used to study biologically realistic circuit models and test the relevance of the microscopic complexity to computations at the level of local circuits. These studies might demonstrate that the molecular specificity of cell and synaptic types with their complex biophysics plays a major role in neuronal computations"

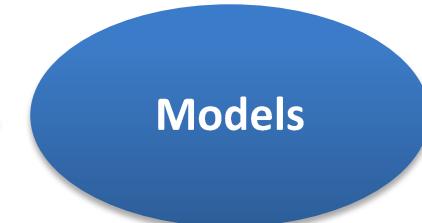
Sompolinsky, 2014

Microcircuit – Somatosensory Cortex

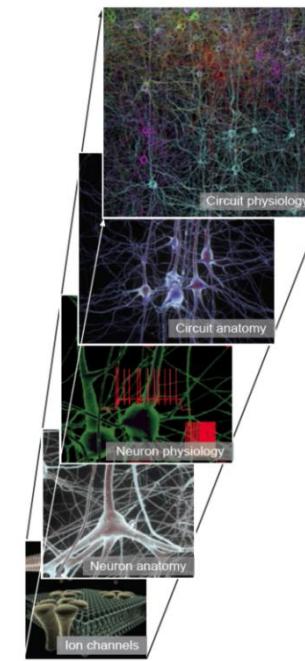


Sparse
Data

Inductive
Principles
Algorithms



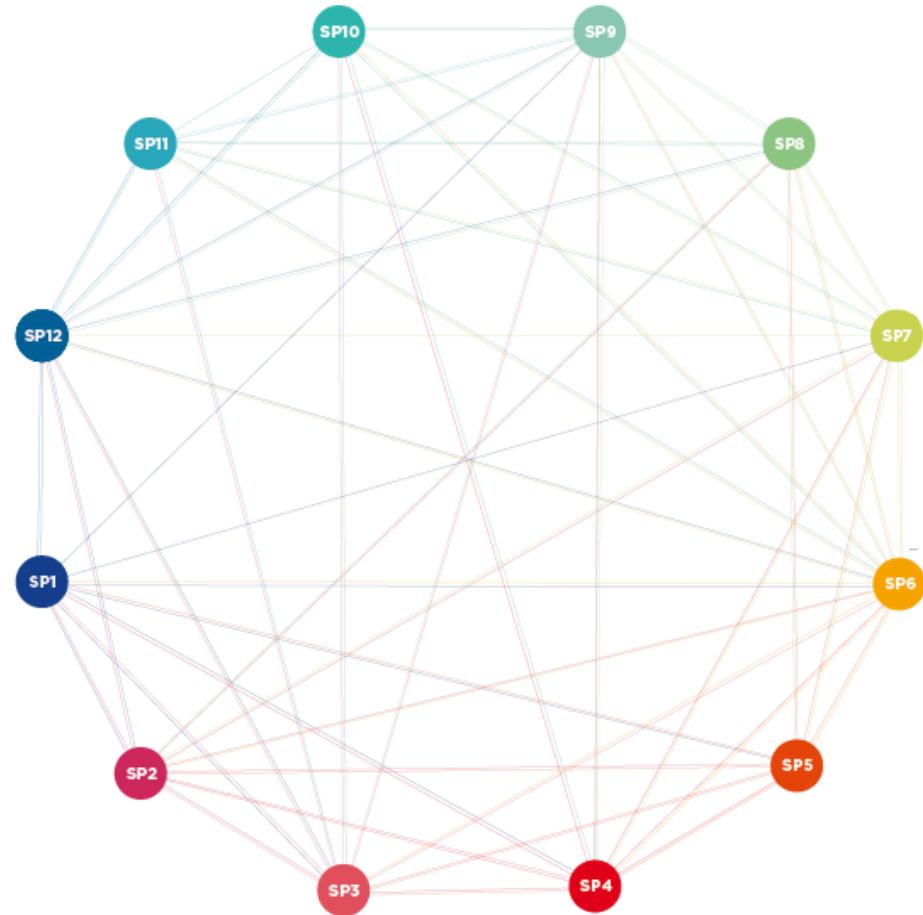
“Massively Collaborative Modeling”



2nd round review: *Markram et. al.*, “Reconstruction and simulation of neocortical microcircuitry”; Collaborative science: 78 authors.

HBP Overview

The HBP is organized in twelve subprojects, spanning the development of six new informatics-based platforms, plus brain organisation, cognitive neuroscience, theory, ethics and society and management



The informatics-based platforms

Neuroinformatics

searchable atlases and analysis of brain data

Brain Simulation

building and simulating multi-level models of brain circuits and functions

Medical Informatics

analyzing clinical data to better understand brain diseases

Neuromorphic Computing

brain-like functions implemented in hardware

Neurorobotics

testing brain models and simulations in virtual environments

High Performance Computing

providing the necessary computing power

Members (WP and Task Leaders)

Data-driven reconstruction of brain models

Henry Markram

Idan Segev
Marc-Oliver Gewaltig
Felix Schürmann



Brain Simulation Platform: integration and operations

Henry Markram

Jeffrey Muller

Brain Simulation Platform: user support and community building

Felix Schürmann

Molecular dynamics simulation

Paolo Carloni

Richard Lavery
Rebecca Wade



Brain simulation engines

Felix Schürmann

Erik De Schutter
Julian Shillcock
Michael Hines
Markus Diesmann
Fabien Delalondre



Brain Simulation Platform: scientific coordination

Felix Schürmann



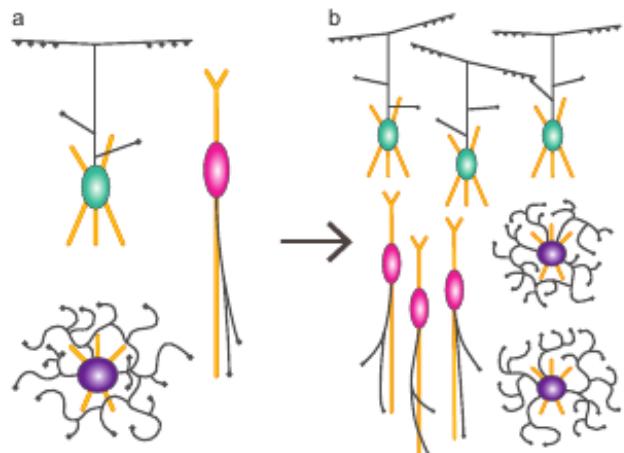
Initial brain models

Jeanette Hellgren Kortaleski

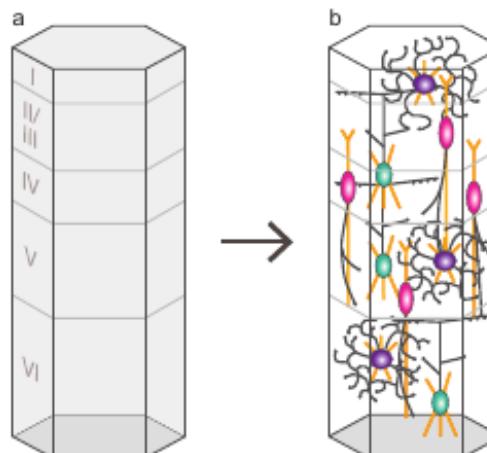
Antoine Triller
Pierre Magistretti
Alex Thomson
Eilif Muller
Egidio D'Angelo
Sten Grillner

Biological Microcircuit Reconstruction Workflow

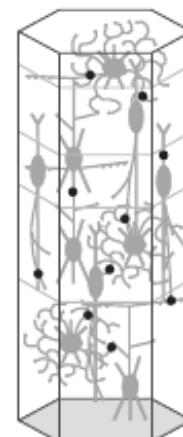
Morphological diversity of neurons:
(a) m-types, (b) cloning



Circuit anatomy: (a) circuit dimensions,
(b) m-type distribution and morphology selection

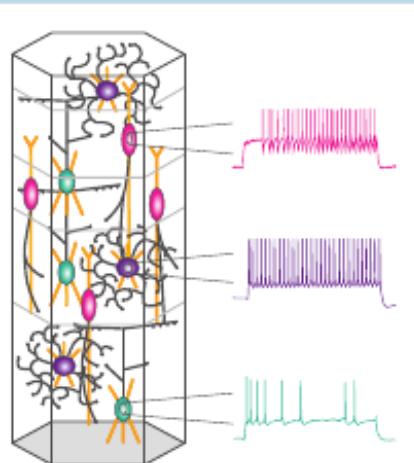


Connectome derivation

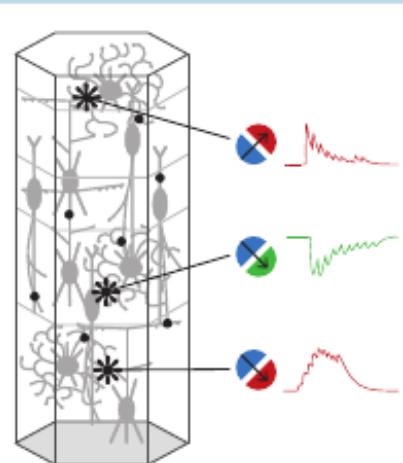


A N A T O M Y

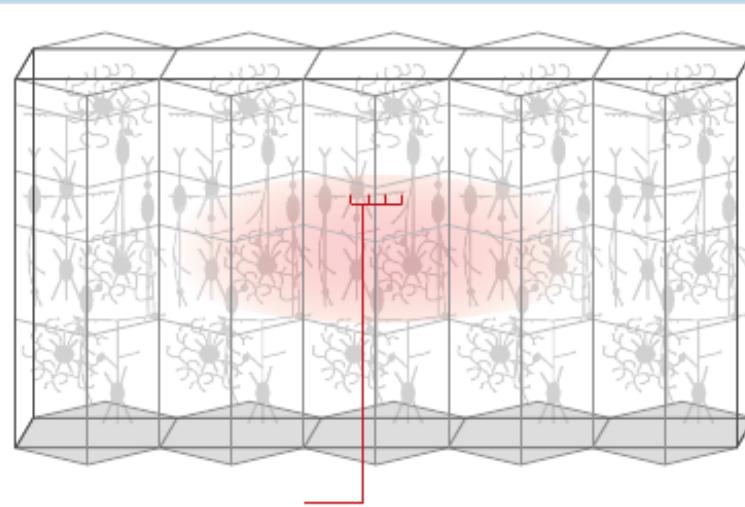
P H Y S I O L O G Y



Electrical diversity of neurons:
e-types



Synaptic diversity of neurons:
s-types



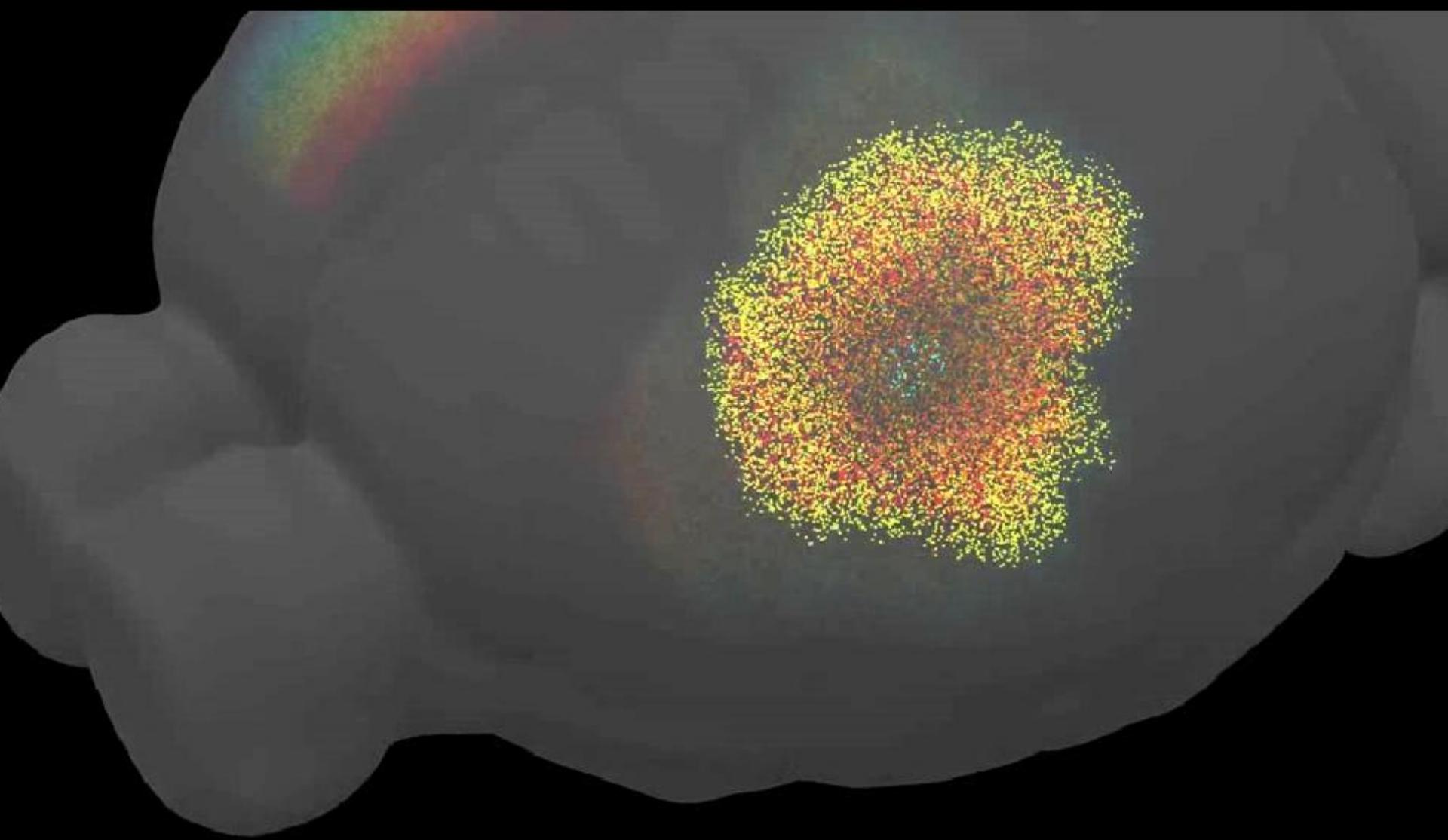
Reconstructing a virtual slice for in-silico experimentation

Generated

Original



Slice comparison with original Nissl stain



Brain Simulation Platform

- The BSP is a collaborative internet-accessible user interface for brain reconstruction and simulation
- High-level Ramp-up Goals:
 - Allow multiple users to collaborate on brain reconstruction and simulation
 - Expose brain building tools and workflows to users
 - Drive user engagement (M18 for HBP members and M30 for community)

WP6.4 Initial Brain Models

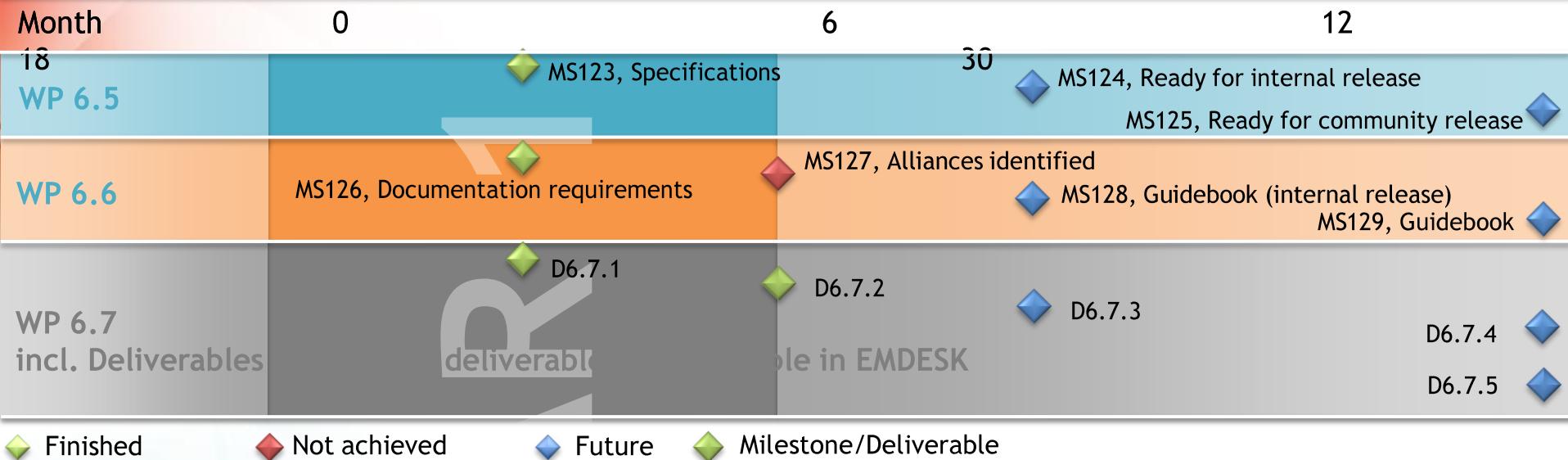
Develop initial models across the scales:

- Macro (whole-brain)
- Meso (brain regions)
- Microcircuits (unitary networks)
- Cells (neurons and glia)
- Sub-cellular (neurons, glia, synapses, vasculature)

to ...

- Exercise the building workflows
- Guide platform development
- Be made available for the community to build-on

Summary to M12



- D6.7.1 - Platform Specification - M6 (June 2014)
- D6.7.2 - Platform Setup Document - M12 (Dec 2014)
- Documentation Requirements
- First developer seminar (Summit 2014)
- To be released to HBP on M18
- To be released to the public in M30

Human Brain Project

Project Number:	28946	Project Title:	Human Brain Project
Document Title:	HBP Brain Simulation Platform Specification		
Document Filename:	HBP_SPh_310516_06_7.1_BSP_Specification_v8.docx		
Deliverable Number:	D6.7.1		
Deliverable Type:	Platform Specification		
Work Package(s):	WP 6.1, WP 6.2, WP 6.3, WP 6.4, WP 6.5, WP 6.6, WP 6.7		
Dissemination Level:	PU		
Planned Delivery Date:	M 6 / 31 March 2014		
Actual Delivery Date:	at 9 /		
Authors:	Henry MARKRAM, EPFL (P1), Jeanette HELLGREN-KOTALESKI, KTH (P33)		
Compiling Editors:	EPFL (P1); Jeffrey MULLER, EPFL MULLER		
Contributors:	EPFL (P1); Sebnis AMEYER, Benoît-Claude COURCOL, Fabien DELALOUCHE, Marc-Olivier GERMALIN, Dan KELLER, Daniel POPPICELLI, Srikar RAJAMONY, Michael REIMAHN, Felix SCHURMANN, Werner VAN GETT, Stefano ZANINETTA		
Reviewers:	JULIUS (P17); Markus DESMAUBL, Abigail MORRISON, Moritz HELIAS, Jochen MARTH EPPLER, Susanne KUNKEL		
Abstract:	The Brain Simulation Platform (BSP) is a collection of ICT performances in the Human Brain Project (HBP), which will be made accessible over the Internet via an HBP Unified Portal (HBP-UP or UP). The HBP's Subproject 6 is developing both the UP and the BSP. This document sets out the specifications for the UP and the BSP, including the models that will run on the BSP. It includes Key Performance Indicators (KPIs) that can be used to track the progress in the UP and BSP development.		

Public Deliverables

https://www.humanbrainproject.eu/en_GB/ec-deliverables

SP6

- Brain Simulation Platform

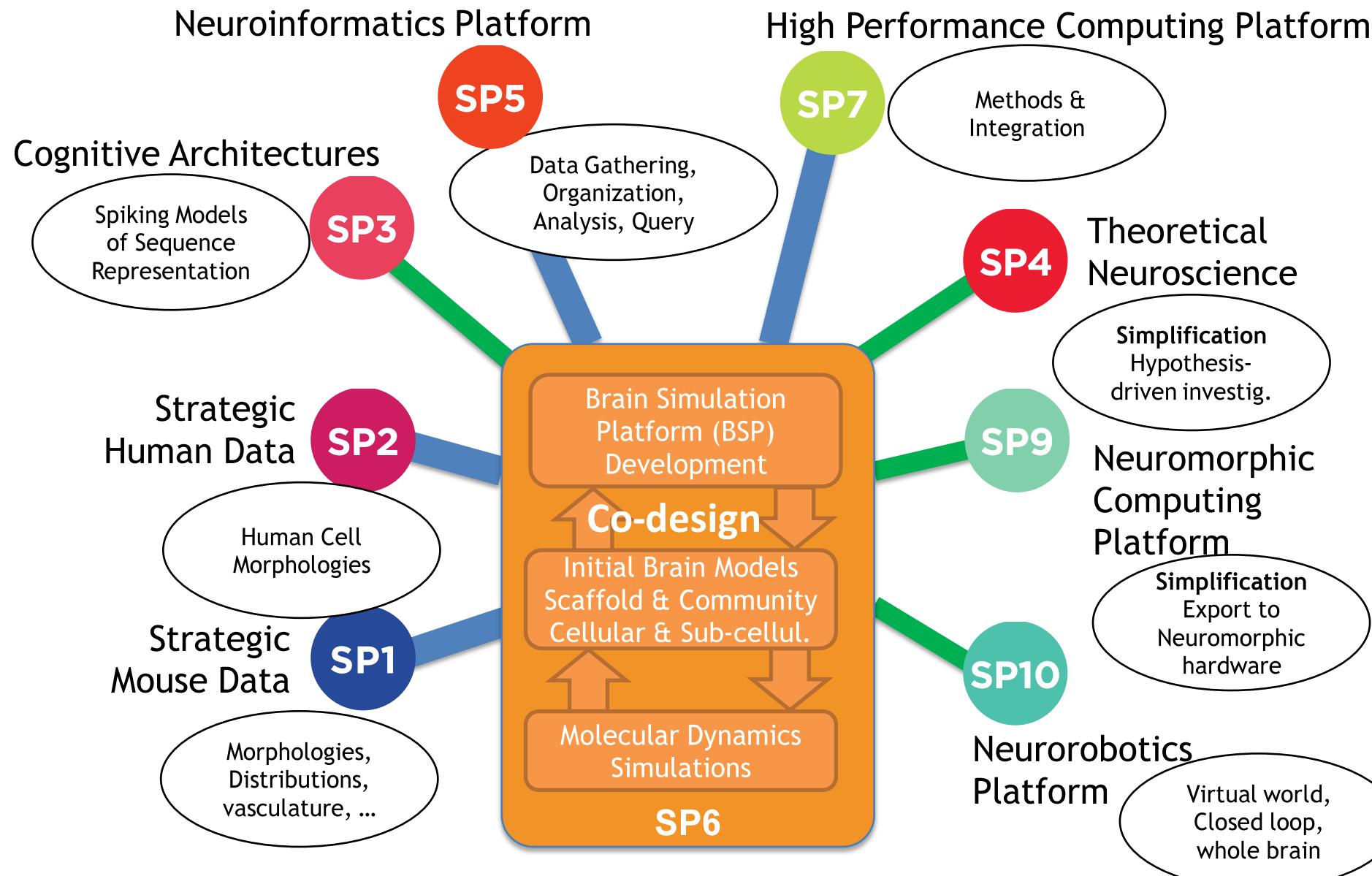
D6.7.1: Brain Simulation Platform v1 - specification document (report).

D6.7.2: Brain Simulation Platform v1 - set-up document (report).

And others from other SPs ...

SP6 - Brain Simulation Platform

Interactions within HBP



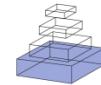
1st HBP release of network simulator

frontiers in
NEUROINFORMATICS

ORIGINAL RESEARCH ARTICLE

published: 10 October 2014

doi: 10.3389/fninf.2014.00078



Spiking network simulation code for petascale computers

Susanne Kunkel^{1,2*}, Maximilian Schmidt³, Jochen M. Eppler³, Hans E. Plessner^{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

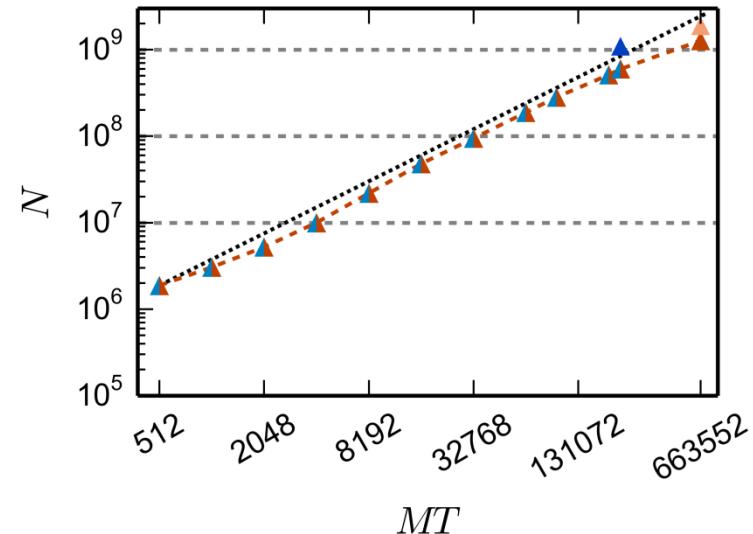
⁴ Department of Mathematical Sciences and Technology, Norwegian University of Life Sciences, Aas, Norway

- scale 100,000 compute nodes
- public open source release Dec 23rd 2014, NEST 2.6.0
- demonstrates that neuroscience can exploit petascale systems

Slide credit: M. Diesmann

HBP Year 1: NEST - Maximum network size

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



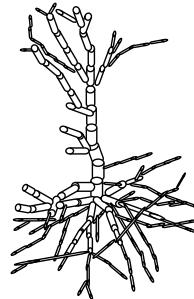
- largest general network simulation performed to date:
- 1.86×10^9 neurons, 6000 synapses per neuron
- 1.08×10^9 neurons, 6000 synapses per neuron

Slide credit: M. Diesmann

HBP Year 1: NEURON Refactoring for HPC



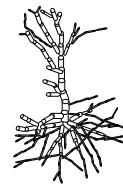
NEURON (300k Lines of Code)



18MB (w/ introspection)



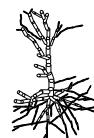
coreNeuron (15k LoC)



MS113
(Month 12)

2.2MB (w/o introspection)

Optimized coreNeuron



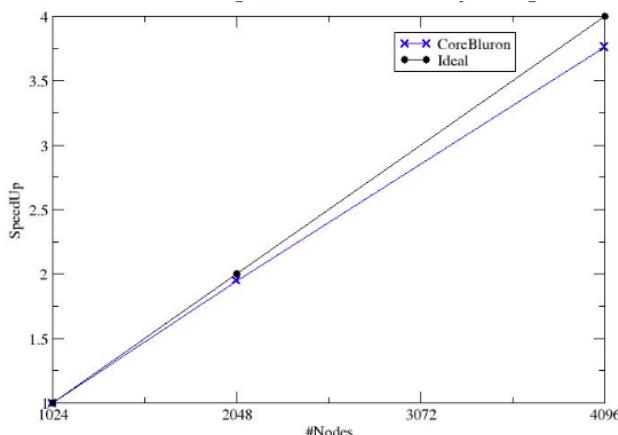
1MB (w/o introspection)

Slide credit: F. Schuermann

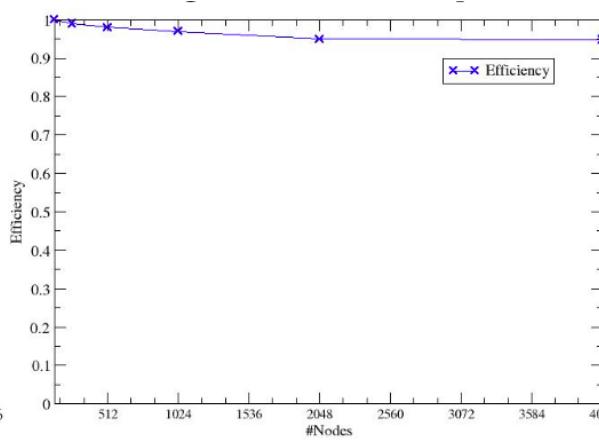
HBP Year 1: NEURON Scaling

- Largest simulated cellular network to date
 - ~25 Million neurons, ~76 Billion synapses
 - HBP Development Computer (4 rack BGQ) @ CSCS
 - Filling ~61TB (of 64TB total) memory; ~370 neurons/core, ~2.5MB/neuron

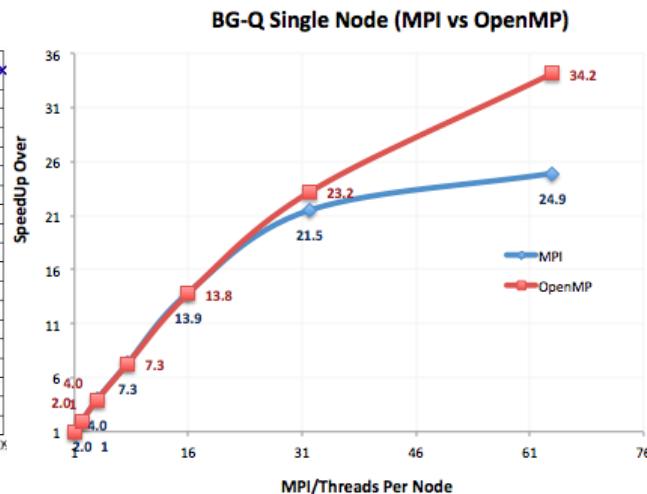
• Scaling



a) Strong Scaling



b) Weak Scaling



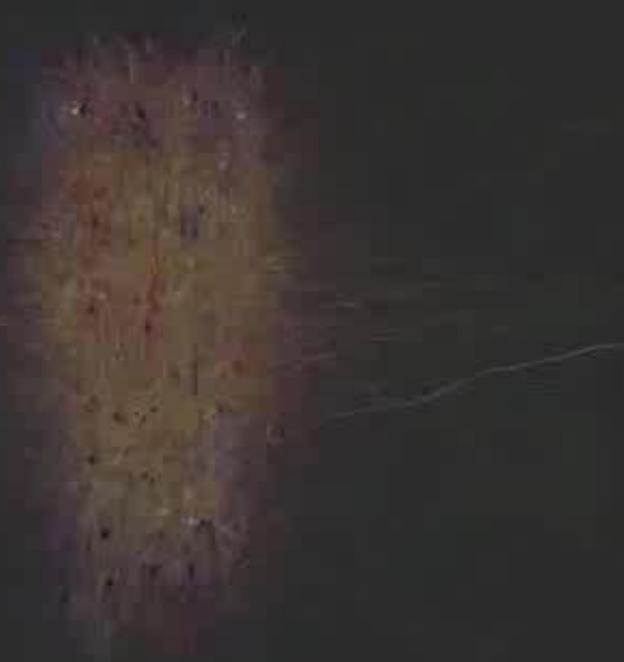
- Scaling to full JUQUEEN (28-racks) achieved
- Planned for Open-source release before FENS 2016

Reconstruction of a neocortical microcircuit



Home Neocortical Microcircuit Literature Consistency Videos Gallery Tools Downloads Copyright

Sign Out



Neural Microcircuit Collaboration Portal

This website provides an online public resource for the Blue Brain Project's first release of a reconstruction of a neocortical microcircuit, rat P14 non-barrel somatosensory cortex. This portal provides the following functionality to support and engage community involvement to size and improve the reconstruction:

- **Neocortical Microcircuit** - an interactive browser of the anatomical and physiological properties of the microcircuit reconstruction. This includes detailed exemplar analyses, simulation video, and model and data downloads.
- **Literature Consistency** - a database of experimental papers which were used either to constrain parameters of the reconstruction directly, or against which the reconstruction was found to be qualitatively or quantitatively consistent. This is intended to be an active list that can be discussed and extended by the community.
- **Videos** - a collection of computer generated animations of actual simulations or *in vivo* experiments.
- **Gallery** - a collection of images telling the story of the reconstruction process.
- **Tools** - tools and documentation for simulating and analyzing the exemplar data provided in this site.
- **Downloads** - downloadable models from the reconstructed neocortical microcircuit.

For the associated scientific papers, please refer to the following links: Blue Brain Project Consortium, Reconstruction of a neocortical microcircuit, 2014; Blue Brain Project Consortium, The Neural Microcircuit Collaboration Portal 2014.

All material is copyright of EPFL/BBP 2005-2014 unless otherwise specified. Further details are available on the [copyright page](#). If you want to use this material, please contact bbp-licensing@bluebrain.ch.

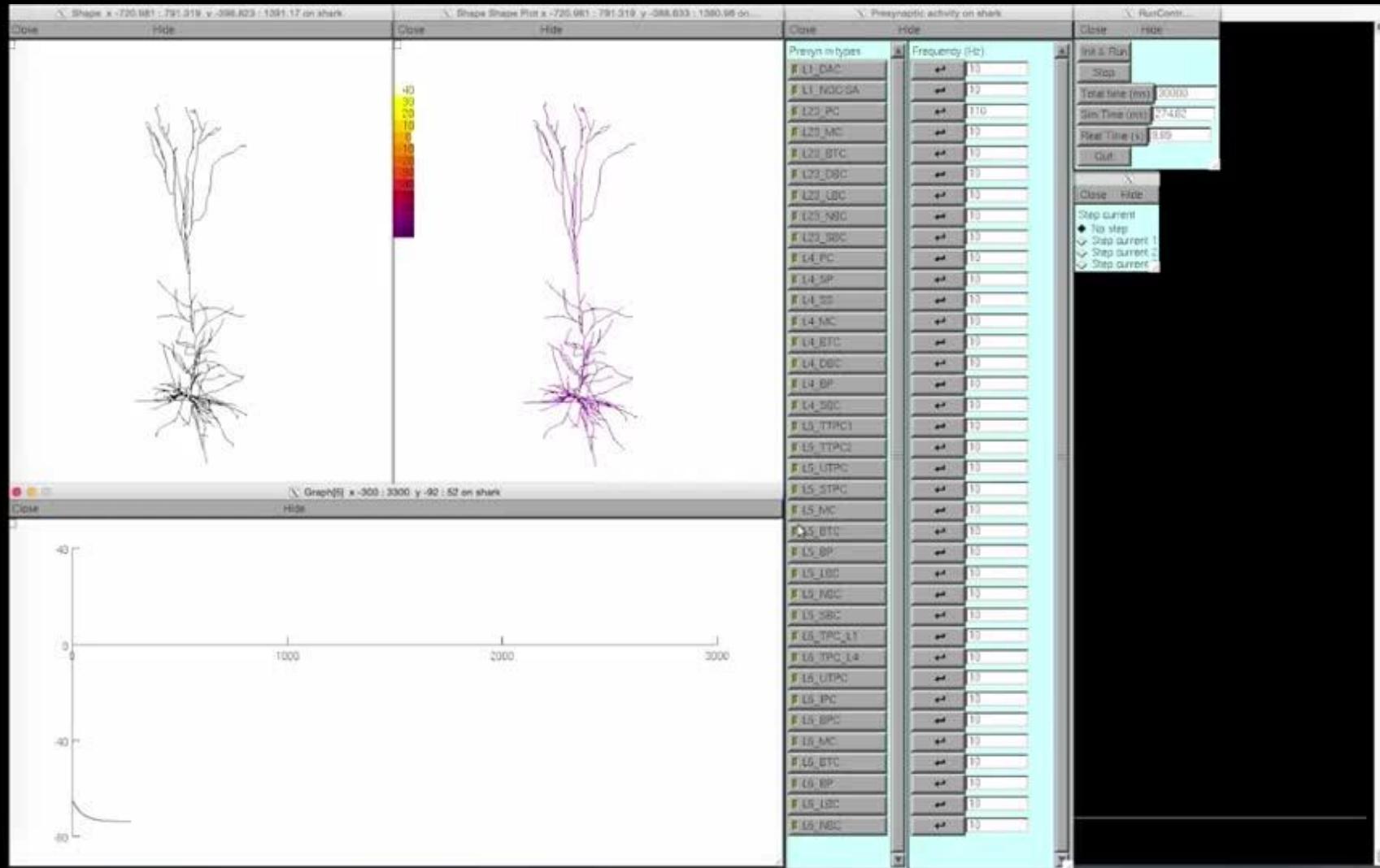
In order to access the content of this site, you will have to log on with your reviewer account.

Latest comments

Public BETA: <https://bbp.epfl.ch/nmc-portal>

Slide credit: J.-D. Courcol, S. Ramaswamy

>1000 Neuron models of 55 excitatory and inhibitory neuron types and incoming synapses released in compact self-contained open-source bundles for NEURON



Working with NeuroML devs to allow export of
models to NeuroML 2.0 , OSB



[NeuroML]

Slide credit: Werner Van Geit

Open Software Repositories

 **The Blue Brain Project**
Lausanne, Switzerland <http://bluebrain.epfl.ch/> bbp-open-source@googl...



<https://github.com/BlueBrain>

 **Human Brain Project**
<https://www.humanbrainproject.eu/>

<https://github.com/HumanBrainProject>

Electrophys Feature Extraction Library

The Electrophys Feature Extract Library (eFEL) allows neuroscientists to automatically extract eFeatures from time series data recorded from neurons (both *in vitro* and *in silico*). Examples are the action potential width and amplitude in voltage traces recorded during whole-cell patch clamp experiments. The user of the library provides a set of traces and selects the eFeatures to be calculated. The library will then extract the requested eFeatures and return the values to the user.

The core of the library is written in C++, and a Python wrapper is included. At the moment we provide a way to automatically compile and install the library as a Python module. Soon instructions will be added on how to link C++ code directly with the eFEL.

The source code of the eFEL is located on github: [BlueBrain/eFEL](#)

Using the eFEL, pyNeuron and the DEAP optimisation library one can very easily set up a genetic algorithm to fit parameters of a neuron model.

http://bluebrain.github.io/eFEL/deap_optimisation.html

Circuit and Simulation I/O

The screenshot shows a web browser window with the URL bluebrain.github.io/Brion-1.5/index.html in the address bar. The search bar contains the text "human brain project github". The page title is "Brion 1.5.0" and the subtitle is "The Blue Brain C++ I/O library". The navigation menu includes "Main Page" (which is selected), "Related Pages", "Classes", and "Files". A sidebar on the left is titled "Brion" and contains links for "Changelog", "Deprecated List", "Classes", and "Files". The main content area is titled "Brion Documentation". It features the Blue Brain Project logo, which consists of a purple neuron-like structure next to the text "Blue Brain Project". Below the logo, there is a welcome message: "Welcome to Brion, a C++ library for read and write access to Blue Brain data structures, including BlueConfig/CircuitConfig, Circuit, CompartmentReport, Mesh, Morphology, Synapse and Target files." There is also a note: "Brion can be retrieved by cloning the [source code](#)".

A Python API is also coming in next months ...

More software developed in part by HBP members



NeuralEnsemble

http://neuraleンsemble.org

<https://github.com/NeuralEnsemble>



PyNN

Python ★ 53 ⚡ 35

A Python package for simulator-independent specification of neuronal network models.

Updated 22 days ago

python-neo

Python ★ 41 ⚡ 55

Neo is a package for representing electrophysiology data in Python, together with support for reading a wide range of neurophysiology file formats

Updated 13 days ago

libNeuroML

Python ★ 21 ⚡ 13

This package provides Python libNeuroML, for working with neuronal models specified in NeuroML

Updated on Jun 26

elephant

Python ★ 11 ⚡ 32

ElePhAnT is the ElectroPhysiology Analysis Toolkit

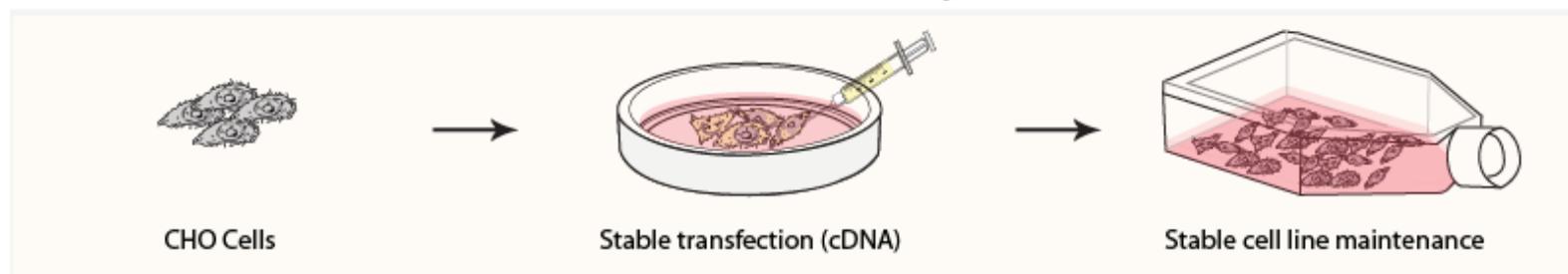
Updated 6 days ago

This is not an exhaustive list ...

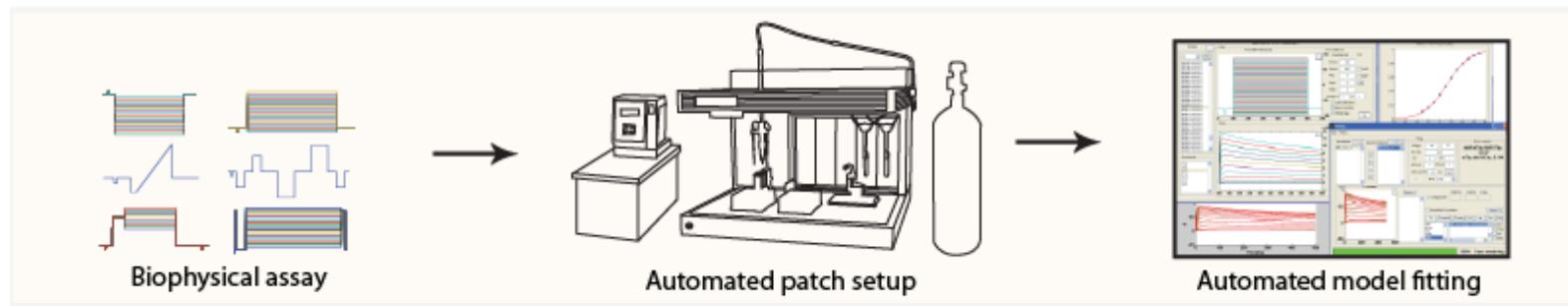
Channelpedia: Ion Channels

Channelome

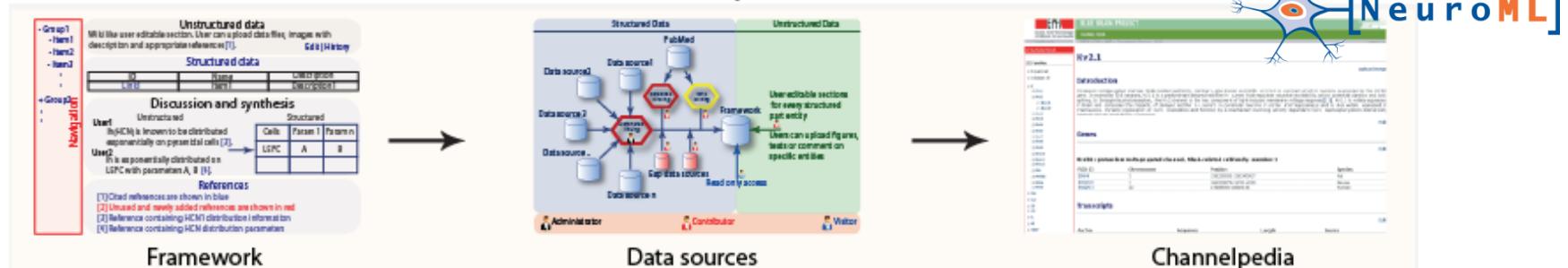
Stable cell line library



Automated biophysics characterization



Channelpedia



Contact: ranjan.rajnish@epfl.ch, emmanuelle.logette@epfl.ch, severine.petitprez@epfl.ch, mirja.herzog@epfl.ch

URL: <http://channelpedia.epfl.ch>

Slide credit: Rajnish Ranjan

HBP Collaboratory

“A ‘dating’ site for [scientific collaboration]”
– Dr. Christine Aicardi

The screenshot shows the HBP Collaboratory platform. At the top, there's a navigation bar with the HBP logo, 'COLLABORATORY', 'HOME', 'COLLABS', and 'HELP'. On the right, there's a user profile for 'JEFFREY CHRISTOPHER MULLER' with options for 'MY COLLABS' and a log-out arrow.

The main area is titled 'Neuroinformatics Platform'. It has tabs for 'Edit' and 'OFF'. Below this is a 'Navigation' sidebar with 'Overview', 'Team', and 'Settings' (which is selected). The main workspace is titled 'Collab settings' and contains fields for 'Title' (Neuroinformatics Platform) and 'Short description (max 140 chars)' (The Neuroinformatics Platform will organize and index a diverse set of neuroscientific data sources connecting researchers with data). There are 'Save' and 'Delete!' buttons at the bottom. To the right is a 'Settings' sidebar with tabs for 'Collaboration' (showing users SH, MT, CZ), 'Activity', 'Provenance', and 'Collaboration' (with 'FEEDBACK' and 'ABOUT' links).



To be released @ FENS 2016 ...

Slide credit: Jeff Muller



Human Brain Project

10th FENS Forum of Neuroscience

July 2-6, 2016 | Copenhagen, Denmark

TECHNICAL WORKSHOPS

Saturday, July 2, 2016, 10:30 - 13:30

W04 - Introduction to the Human Brain Project Collaboratory.

Chaired by: E. Muller (Geneva, Switzerland); A. Davison (Gif sur Yvette, France)

"This workshop will provide an introduction to the Human Brain Project web-based platforms, ""The HBP Collaboratory""", with a focus on neuroscientific use-cases available as of the initial public release of spring 2016, and first success stories. These include neuroinformatics and data integration, single neuron and neural tissue modelling and analysis, integration with 3D brain atlases, and the joint Allen Institute for Brain Science and HBP community data and model release of March 2016."

Muller J. (Geneva, Switzerland)

The HBP "Collaboratory".

Zwahlen C. (Geneva, Switzerland)

The Neuroinformatics platform: Contributing and querying data.

Ramaswamy S. (Geneva, Switzerland)

Interactive resources for e-Neuroscience.

Grün S. (Juelich, Germany)

Analysis of large-scale recordings of neural activity in vivo and in silico.

D'Angelo E. (Pavia, Italy)

Modeling cerebellar neurons and microcircuits using the HBP Collaboratory.

Anastassiou C. (Seattle, USA)

Collaborative transatlantic efforts to create faithful single-neuron representations for the community.

HBP – Brain Simulation Platform - Relationship to other initiatives

- Focused on platforms
- Model building pipelines and workflows from Experimental data -> Neuroinformatics -> data-driven models
- Model validation suites bound to Neuroinformatics platform
- Not a data-generation project
 - Strategic data for atlas-based brain building in mouse and human
- Simplification strategies to engage users (mostly hypothesis-driven)
- Foundation software
 - large-scale simulation, analysis and visualization using supercomputers
- Closed-loop (with Neuromorphic , SpiNNaker) and Neurorobotics
- Will leverage/integrate/federate/develop many community developed tools
- Round-tripping with OSB via NeuroML
- Transitioning from Cathedral to Bazaar model

OSB

OpenWorm (Bazaar)

Geppetto (Bazaar)

Allen Brain Institute - Mind Scope

HBP Building Workflows

Morphologies &
Electrical models

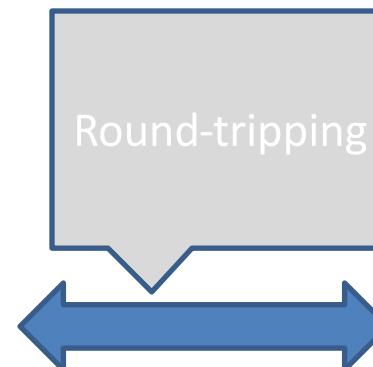
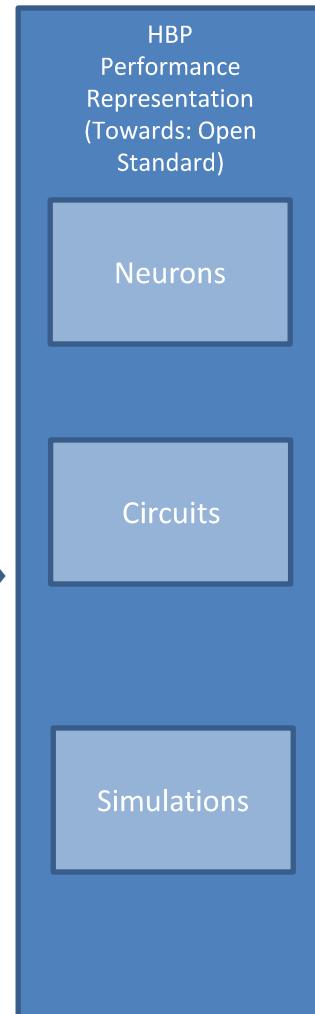


Circuits & Synapses



Simplifications

Simulations
STEPS, NEURON, NEST



Other HBP
supported
representation
formats ... ?

Old argument

We can't collaborate because we
don't have the tools!



SO WHAT HELPS PEOPLE COLLABORATE?

Believing
ANY idea
COULD BE
THE idea



Show how
IT SAVES
and



STIMU-
LATION
ARDI-
SATION

GIVING EVERYONE
ON
RO

NO TOOLS



REQUIRED

Mutual Respect

Demonstrate
AND
Celebrate
THE RESULTS OF
TEAMWORK

GOOD SEARCHABLE
KNOWLEDGE
at your FINGERTIPS

2 Minds
ARE ALWAYS
BETTER

HBP Hippocamp CA1: Collaborative and Integrative Modeling of Hippocampal Area CA1

31st March-1st April 2015, London, United Kingdom



Aim #1

- Engage community
- Highlight existing:
 - modeling efforts
 - strategic datasets

Aim #2

- Define and bootstrap an inclusive community-driven model and data-integration process
- Open pre-competitive reference models of hippocampus CA1
- Documented, validated, and alive

<http://neurallensensemble.org/meetings/HippocampCA1>

Why now?

- HBP roadmap to deliver models of neocortex, hippocampus, cerebellum of similar detail to BBP cortical column model (April 2016)
- CA1 Model by Sept 2015
- **HBP mandate for open, collaborative modeling**
- **Be inclusive, non-competitive with work of community of experts**
- **Avoid yet another model**
- **Models by community for community**

Next steps



- Openness consensus ✓
- Discuss relationship between HBP collaboratory and OSB
- Public Discussion Forum (June 1st) ✓
- **Hippocamp opinion paper**
 - Sources of fragmentation
 - Possible solutions
 - Review state-of-the-art
 - Community roadmap
- **Form CA1 Neuron Models Sub-group**
 - eFEL <https://github.com/BlueBrain/eFEL> ✓
 - http://bluebrain.github.io/eFEL/deap_optimisation.html ✓
- First release CA1
- Demonstrate: Simplification and hypothesis-driven use cases
- **FENS 2016 – Technical workshop - Copenhagen**
“Introduction to HBP Collaboratory”

<https://forum.humanbrainproject.eu/>

Screenshot of the Human Brain Project Forum website (https://forum.humanbrainproject.eu/). The page shows a list of topics, with the first topic being pinned.

The browser address bar shows the URL https://forum.humanbrainproject.eu. The search bar contains the text "human brain project github".

The forum header includes the Human Brain Project logo, a "Beta" badge, "Sign Up", "Log In", and a search icon.

The topic list includes:

- Welcome to Discourse**
The first paragraph of this pinned topic will be visible as a welcome message to all new visitors on your homepage. It's important! Edit this into a brief description of your community: Who is it for? What can they ...
[read more](#)
- eFeature Extraction Library (eFEL) is now open source and available on GitHub
- Human Brain Project Community Forum

A message at the bottom states: **There are no more latest topics.**

What is a community model?

HBP Collaboratory: Whole Community Modeling

An end-to-end approach which integrates the whole community, and will have a lasting impact on addressing fragmentation

Driving use case:

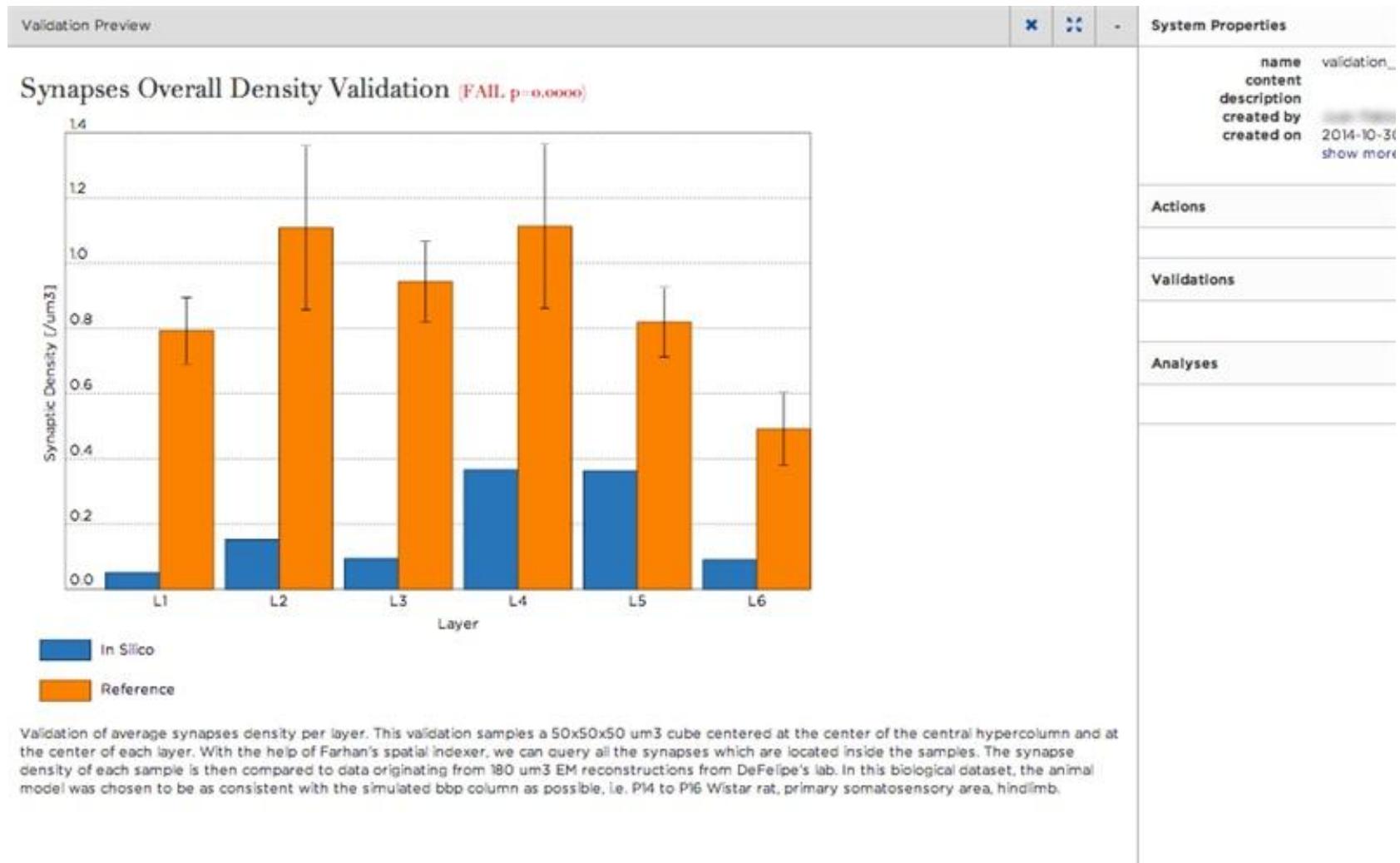
Experimentalists -> Neuroinformatics ->
Data-driven models -> Extensive validation ->
Principled simplification & validation ->
Hypothesis-driven science ->
Neuromorphic application

Contrast: Whole community building models

HBP Collaboratory: Whole Community Modeling

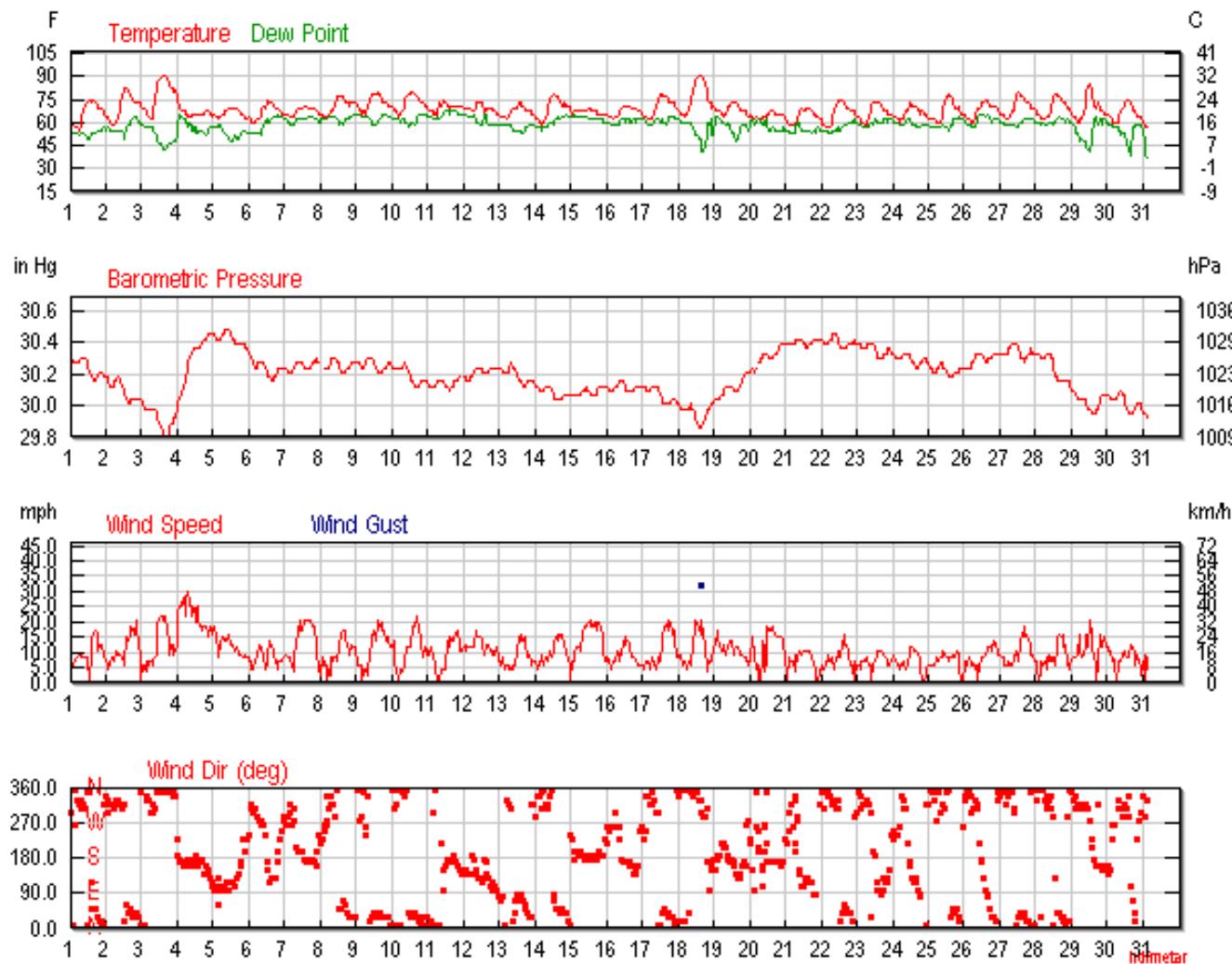
- Open data-driven model building tools beyond BBP
- Parameterization and validation data registered in (community or HBP) Neuroinf. platforms
- Transparency of model parameterization
 - Integration with Neuroinformatics
 - Provenance: Auditable by 3rd party
- Automated anatomy and physiology validation suite ala Test-Driven-Development (TDD)
 - User contributed validations
 - against data pulled from Neuroinformatics
 - Whole suite click button runnable by everyone
 - Produces human readable report, statistical tests against data, pass-fail
 - History tracked across releases

“Continuous Integration” for models



History tracking of validations ...

Models are finicky like the weather ...



~6 month release cycle?



- Freeze and release e.g. every 6 months
 - “It’s ready” ↗ archival checkpoint
 - End-to-end publication quality build
 - Achievable refinement “chunks”
 - Resource for hypothesis-driven science
 - Automatically generated simplifications
- Cost of release growing
 - As more data is integrated
 - Library of simplifications (and their validations)
 - Growing community contributed validation suite
 - Simulations as validations ...

HBP Collaboratory: Whole Community Modeling

- Functionality for regular releases
 - History of validations to track regression
- Community facilities to discuss and plan refinement roadmaps
 - ticket/task system, wikis, agile estimation tools, a discussion forum, comment streams
- Principle-based simplification
 - >same quality metrics as reference model
 - assess impact of simplification.
- Loss-less export of simplified models to neuromorphic
 - Compatible representation, e.g. PyNN.

Level of detail: A rift in neuroscience

1. Simplify the details

- minimal model for hypothesis-driven science
- Adhoc simplification
- Minimal for which question?

vs

2. Consider all known

- data-driven is data-ready
- Hypothesis-free integration of facts
- Algorithms fill in gaps from sparse data
- Fewer free parameters!
- Avoid wasting time hand tuning parameters for a given model “island”

Level of detail - False Dichotomy

- Integrate new data in models @ level of data
 - *Data-driven is data-ready*
 - Algorithms fill in gaps from sparse data
 - **Fewer free parameters!**
 - Avoid wasting time hand tuning parameters for a given “hypothesis island”
- Necessary but not sufficient:
 - Automated & principled & repeatable & quantifiable simplification
 - ... of data-driven reference
 - ... to minimal model for given hypothesis

$$f(x) =$$

$$f(a) + f'(a)(x-a) + \frac{f''(a)}{2!}(x-a)^2 + \frac{f^{(3)}(a)}{3!}(x-a)^3 + \dots + \frac{f^{(n)}(a)}{n!}(x-a)^n + \dots$$

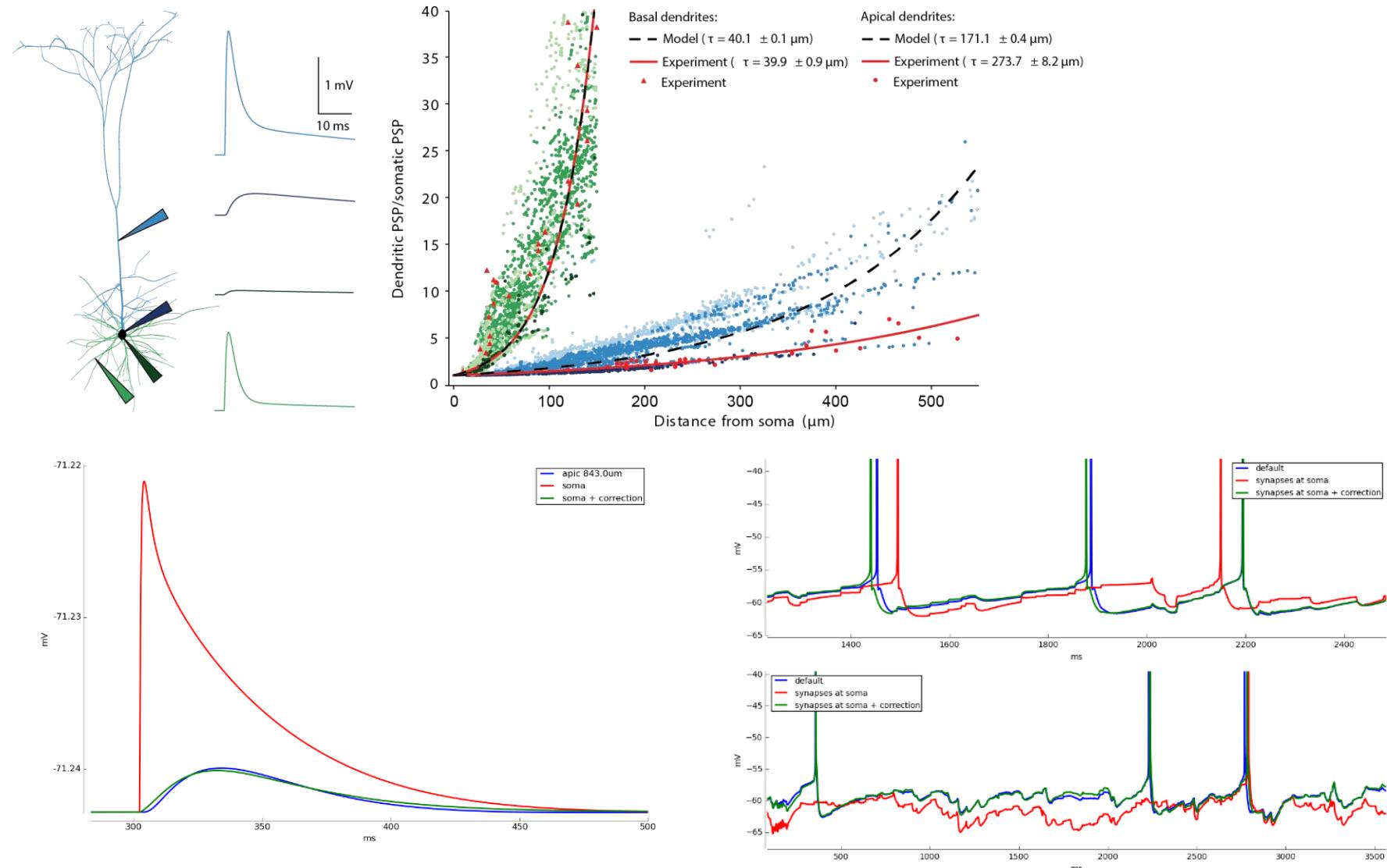
We know a lot, but understand very little ...

Take best of both worlds ...

Road to achieve community convergence?

- **Data-driven is data-ready -> resource**
- Latest biological facts taken into account at regular release intervals
- Growing automated validation test suite
- Hypothesis-driven **quantitative automated reproducible simplification**
- Families of simplified models (I&F, mean-field, simplified morphology, ...) are generated automatically from release

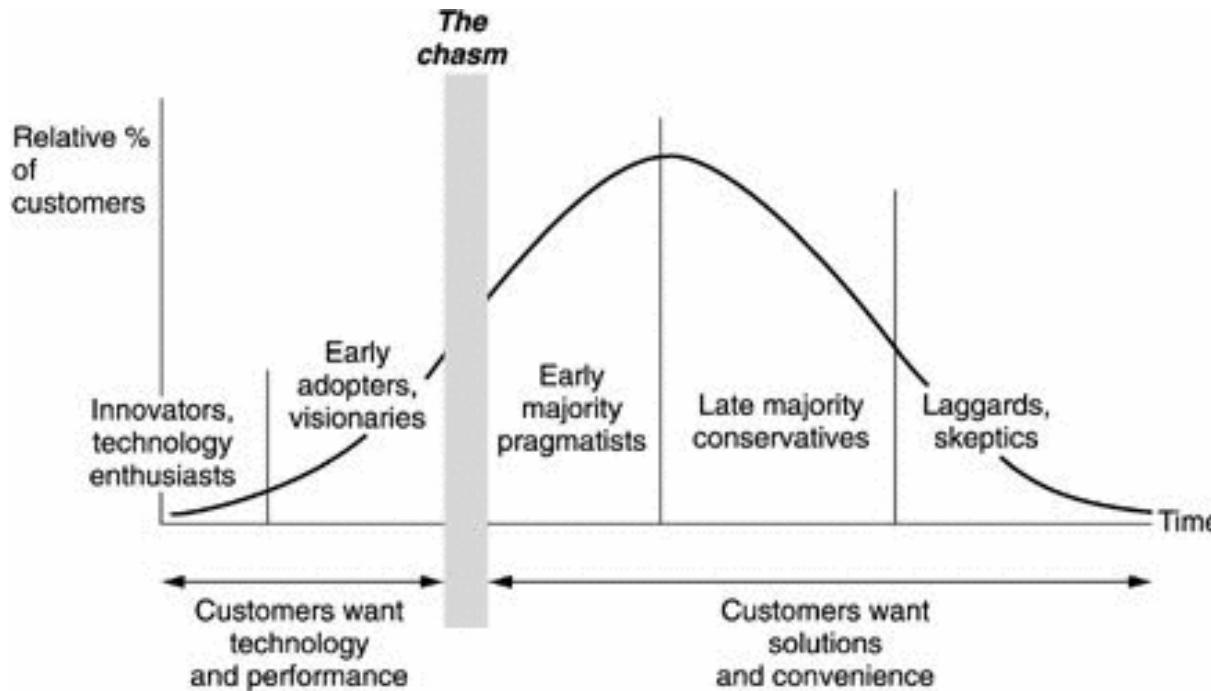
Simplification example: Collapsing synapses onto the soma (C. Roessert)



Working with HBP partners to fit point-neuron models suitable for NEST
(Circuit -> NMC portal; example simplification -> github)

Slide credit: Christian Roessert

Community vs Scaffold: The adoption curve



- **We seek early adopters! Establish a badwagon**
- **Work out key problems**
- **Target capabilities towards value proposition**
 - Prioritize implementing success stories
 - Don't focus on convenience now
- **Early adoptors: Scaffold models ... need to be on HBP payroll**
 - Co-designers of the pipeline: Experimental data -> Neuroninf -> Validated data-driven model
- **Early majority: Community models ... can't be on HBP payroll**
 - Key use case is: Whole-Community "Experiment to Neuromorphic"

Summary

HBP Collaboratory: Whole Community Modeling

An end-to-end approach which integrates the whole community, and will have a lasting impact on addressing fragmentation

Driving use case:

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Contrast: Whole community building models

Early adoptors: Scaffold models

- Co-designers of the pipeline: Experimental data -> Neuroninf -> Validated data-driven model
- For specific domains of expertise (Hippocampus, Cerebellum, ...)
- On HBP payroll to trail blaze key problems
- Establish a badwagon to jump on.
- Second phase: Drive boot-strapping community process

Early majority: Community models

- **Leverage Scaffold model workflows & pipelines**
 - Parameterization and validation data registered in (community or HBP) Neuroinf. platforms
 - **Transparency of model parameterization**
- **Benchmark/validation suite with community contribution**
 - Quantitative history
 - **Feedback on regression due to refinement**
 - **Low-barrier for community to become stakeholder**
- **Community facilities to coordinate roadmaps**
- **Support community use-cases:**
 - Portal: Simulation launch to HPC infra, collab. Analysis
 - **Simplification & export**
 - **Success stories drive adoption**

Acknowledgments



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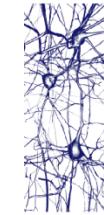


The BBP team

Hippocamp Participants



ÉCOLE POLYTECHNIQUE
FÉDÉRALE DE LAUSANNE



Blue
Brain
Project

SP6

Members (WP and Task Leaders)

Data-driven reconstruction of brain models

Henry Markram

Idan Segev
Marc-Oliver Gewaltig
Felix Schürmann



Brain Simulation Platform: integration and operations

Henry Markram

Jeffrey Muller

Brain Simulation Platform: user support and community building

Felix Schürmann

Molecular dynamics simulation

Paolo Carloni

Richard Lavery
Rebecca Wade



Brain simulation engines

Felix Schürmann

Erik De Schutter
Julian Shillcock
Michael Hines
Markus Diesmann
Fabien Delalondre



Brain Simulation Platform: scientific coordination

Felix Schürmann



Initial brain models

Jeanette Hellgren Kortaleski

Antoine Triller
Pierre Magistretti
Alex Thomson
Eilif Muller
Egidio D'Angelo
Sten Grillner

BBP Platform Team



Human Brain Project

HBP Platform Teams

- SP5 – Neuroinformatics
- SP6 – Brain Simulation
- SP7 – HPC
- SP8 - Medical Informatics
- SP9 – Neuromorphic
- SP10 - Neurorobotics



Human Brain Project