CNS 2017

26th Annual Computational Neuroscience Meeting

July 15-20 2017 University of Antwerp, Belgium



The Programme Book



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We are grateful to the following organizations for their support without which none of this would be possible:









Overview

Organization for Computational Neurosciences (OCNS)

2017 Board of Directors

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- Tatyana Sharpee (Salk Institute, San Diego, USA).
- Tatjana Tchumatchenko (Max Planck Institute for Brain Research, Frankfurt/Main, Germany).
- Krasimira Tsaneva-Atanasova (University of Exeter, UK).
- Wim van Drongelen (University of Chicago, USA).
- Christina M Weaver (Franklin & Marshall College, USA).
- Si Wu (Beijing Normal University, China).

2017 Local Organizers

- Michele Giugliano (University of Antwerp, Belgium).
- Daniele Marinazzo (University of Ghent, Belgium).

Fundraising

OCNS, Inc is a US non-profit, 501(c)(3) serving organization supporting the Computational Neuroscience community internationally. We seek sponsorship from corporate and philantropic organizations for support of student travel and registration to the annual meeting, student awards and hosting of topical workshops. We can also host booth presentations from companies and book houses. For further information on how you can contribute please email http://sponsorship@cnsorg.org.

Timetable

	Tutorials Main meeting					Workshops			
l Î	Sat, 15 July	Sun, 1	6 July	Mon, 17 July		Tue, 18 July		Wed, 19 July	Thu, 20 July
8:00	Registration	Regist		Registration		Registration		Registration	Registration
9:00		Announ	uncements Announcements		Announcements				
9:10 9:20	TUTORIALS								
9:30	morning session 1	Keyn		Keyno		Keyno		WORKSHOPS	WORKSHOPS
9:50		P. Poirazi E. De Schutte		chutter	K. Fri.	ston	morning session	morning session	
10:00									
10:10	Brook	Break Break		Dec	eak	Bre	ak		
10:20	Dreak			DIE	edK	ые	dK		
10:40			01		09		017	Break	Break
10:50			01		09		017		
11:00	TUTODIALC		02		010		018		
11:10 11:20	TUTORIALS morning session 2	Oral Session I		Oral Session IV		Oral Session VII			
11:30	morning session 2	363310111	03	Je331011 TV	011	Jession VII	019		
11:40			04		012		020	WORKSHOPS	WORKSHOPS
11:50			04		012		020	morning session	morning session
12:00									
12:10 12:20									
12:30	Lunch Break	Lunch	Break	Lunch	Break	Lunch	Break		
13:10									
13:20								Lunch Break	Lunch Break
13:30			05						
13:40 13:50					Featured Oral 2				
14:00	TUTORIALS	Oral	06	Oral	Olul 2	OCNS Memb	er Meeting		
14:10	afternoon session 1	Session II		Session V	013				
14:20	1		Featured		013				
14:30 14:40			Oral 1		014		021	WORKSHOPS	WORKSHOPS
14:40						Oral		afternoon session	afternoon session
15:00			Bre	Break Session VIII Featured					
15:10							Oral 3		
15:20 15:30		Oral	07	Oral	015				
15:40	TUTORIALS	Session III		Session VI		Bre	ak	Break	Break
15:50	afternoon session		08		016				
16:00	2								
16:10									
16:20									
17:00 17:10	Welcome and Announcements							WORKSHOPS	WORKSHOPS
17:20	ouncements							afternoon session	afternoon session
17:30			ession I	Poster S		Poster Se			
17:40	Keynote 1	P1-F	2104	P105-	-P209	P210	-314		
17:50 18:00	S. Denham								
18:10									
18:20									
18:30									
18:50									
19:00 19:10	Welcome					Travel to	Banquet		
19:20	Reception								
19:30									
19:50				CNS I	Party				
20:00						BANC	UET!		
20:20									
21:00									

General Information

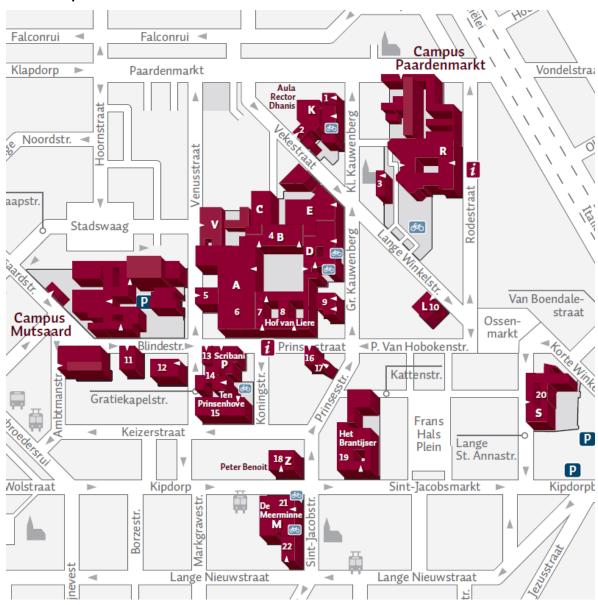
Meeting venue

City Campus of Antwerp University Kleine Kauwenberg 14 and Grote Kauwenberg 2 Antwerp Belgium



The conference will be held in the City Campus of Antwerp University, located in the historical center of Antwerp. The campus is well served by public transport. The main train station is within walking distance, as well as several bus stops.

Plan of the campus:



Meeting room locations:

See summary on back cover or single maps included in the corresponding sections (Tutorials, Main Meeting, Workshops).

Getting to the conference venue

From Brussels-Zaventem Airport:

Δ

Take the train to conveniently travel from Brussels Airport to Antwerp Central Station. Brussels Airport-Zaventem station is located on level -1 of the terminal, at a small distance from the arrivals hall (2nd floor) and the departures hall (3rd floor). The trip lasts 32 minutes, and the fare is 11.60 EUR. Info: http://www.brusselsairport.be/en/passngr/to-from-brussels-airport/train

B.

There is also a Antwerp-Brussels Airport express. The stop "Kon. Astridplein" (near the central station) is always served, the stop "Crowne Plaza" is only served on request. First departure 3am, last departure midnight, travel time 45 minutes. Tickets: 10 EUR. Info http://www.airportexpress.be/page?&orl=2&ssn=&lng=2&pge=4

From Amsterdam Schiphol Airport:

Take the train to conveniently travel from Amsterdam Schiphol Airport to Antwerp Central Station. You can use the Thalys fast train, trip duration circa 60 min, tickets around 50 EUR. Info: https://www.thalys.com/be/en/

Information for poster presentations

The poster area is located in building E (SportHall) as shown in the campus map. Poster boards will be numbered. Fixing adhesive material will be available. Poster orientation is portrait and maximum net dimensions are 180 cm x 95 cm (A0 is smaller than 120 x 85 cm/ or 47 x 33 in).

Poster Sessions will be held on July 16, 17, and 18 from 4:00 pm to 7:00 pm Poster set up:

Sunday, July 16, 2017: From 8 am to 4 pm Monday, July 17, 2017: From 8 am to 4 pm Tuesday, July 18, 2017: From 8 am to 4 pm

Poster should be removed:

Sunday, Monday, July 16 and 17, 2017: Next day morning latest

Tuesday, July 18, 2017: Before 7 pm

Posters that are not removed by the stated time will be discarded. The organisers are not responsible for loss or damage to posters not removed by the given times.

Registration and locations

Registration will be held at the hall of the Auditorium K.001 Aula Rector Dhanis. The entrance is at Kleine Kauwenberg 14.

Registration hours:

July 15: From 8 am to 6:30 pm July 16: From 8 am to 7 pm July 17: From 8 am to 7 pm July 18: From 8 am to 6:30 pm July 19: From 8:30 am to 6:30 pm July 20: From 8:30 am to 6:30 pm

Locations (see also floor plans in the Tutorials, Main Meeting and Workshop sections):

What	Where
Tutorials	Aulas B.001, B.002, B.003, B.004, K.101, K.102
Keynote Lectures and Oral Sessions:	Auditorium K.001 Aula Rector Dhanis
Workshops	Aulas C.101, C.102, K.101, K.102, K.103, K.201, K.202, K.203
OCNS board/program committee meetings	Aula E.201
Welcome Reception	Hall Auditorium K.001 Aula Rector Dhanis
Coffee Breaks	Hallway Building B through E

Local information

Good to Know

Detailed information is available on the official Belgium website at https://www.belgium.be/en/about belgium.

Official Language

The official language of the meeting is English. Interpreting is not provided.

Insurance

The organisers do not accept responsibility for individual medical, travel or personal insurance. All participants are strongly advised to take out their own personal insurance before travelling to Antwerp.

Currency & Banking

The Euro (EUR) is the official currency of Belgium. Exchange of foreign currency is available at the Airports, Central Station, and at most hotels, banks and exchange offices throughout the city. International credit cards are accepted for payments in hotels, restaurants and shops. Payment in cash (in EUR) is also available in some restaurants and shops, so please ask for details on-site if necessary. You can find the official exchange rates on the European Central Bank website at:

https://www.ecb.europa.eu/stats/policy_and_exchange_rates/euro_reference_exchange_rates.

Electricity

Belgium uses a 230 volt 50 Hz system. Sockets follow the standard also used in France, Denmark, Poland, Greece, Italy, Ireland and other countries and also have a grounding pin sticking out of the power socket, which is also known as a type E socket.

Shopping

Most shops in Antwerp are open from 9:30 am to 6:00 pm, from Monday to Saturday. However, some grocery stores may be open until 8:00 pm, from Monday to Sunday.

Time Zone

Belgium is on Central European Time (CET), which is Greenwich Mean Time (GMT) + 1 hour. Note that April to October is daylight saving time, i.e. GMT + 2 hours.

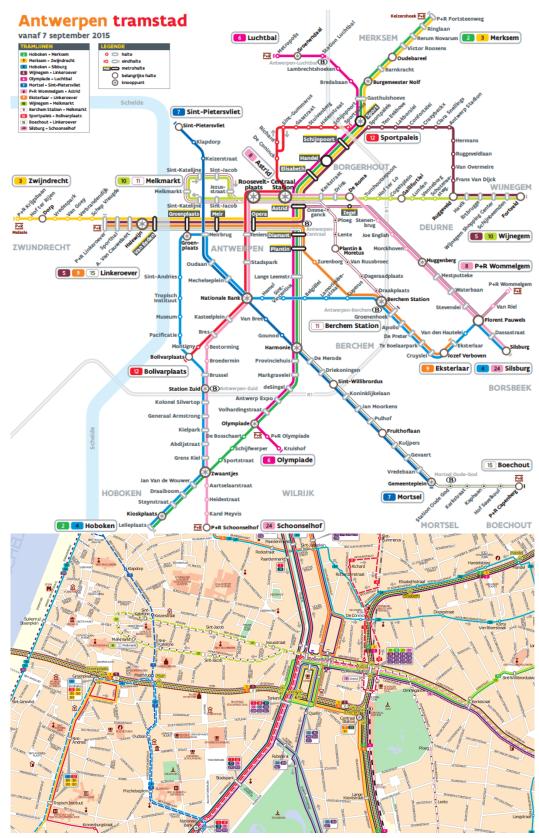
Tipping

Service is usually included in the bill in most bars and restaurants, but tips are welcome.

Tours

Tours are not a part of the meeting. If you wish to explore the city, check the possibilities in your hotel or choose the tour directly at: http://www.visitantwerpen.be/en

Transportation



https://static.delijn.be/Images/Antwerpen%20centrum%201op7500_tcm5-5301.pdf

Get around by public transportation

The public transportation company De Lijn (www.delijn.be/en/) operates a dense network of buses, trams, and underground tram connections in the city and its surroundings. If you plan on taking a bus or tram more than 5 times, then buy a 10-ride card (Lijnkaart) costing 15 EUR. They can be bought at self-service machine (no electronic payment means accepted however) or at fixed points in town (e.g. most supermarkets and any place that sells newspapers, just ask the cashier). Every time you enter a bus or tram, just put that card in one of the yellow ticket machines. A single ticket bought from the driver in the bus costs more (3.00 EUR per ride). For one fare, you can ride up to 1h within the city boundaries. The central bus station is the Franklin Rooseveltplaats, walking distance from the congress venue and also very close to the central railway station. Nearly all buses leave from there, or from the Antwerp-Central or Antwerp-Berchem train stations.

http://www.visitantwerpen.be/en/plan-your-visit/transport-antwerp/public-transport

On foot

Most things to see are within walking distance from the venue of the conference. These are near or within the Boulevards, the half-moon of avenues where there were once 16th century city-walls. This old town center, with a diameter of about 1.5 km can be walked, but there is excellent public transport.

By horse (!)

Horse tram (paardentram: (www.werkendtrekpaard.be/paardentram/) leaves from the Grote Markt every hour. It is an approximately 40 minutes / 1.5 mile touristic ride through the city.

Weather

The month of July is a typical summer month with daily high temperatures, which can occasionally exceed 30 degrees C. You can check for current weather conditions in Antwerp at www.meteovista.be

Important Telephone Numbers

112: General Emergency for Europe

100: Fire and Ambulance

101: Police

Free Wi-Fi

Free Wi-Fi is provided at the meeting venue. Each registered participant will receive their own unique password.

Taxi Services

Taxis are available, but they can be quite expensive. They await customers at specific locations around town (waving your hand will seldom work) like the Groenplaats (near the cathedral) or the central railway station. You can recognize these places by an orange TAXI sign. The prices are fixed in the taximeter. Beware that Antwerp and Brussels were named as the most congested cities in Europe and North America.

DTM Taxi - Transport - Moving: +32 03 366 66 66

Antwerp - Tax N.V.: +32 03 238 38 38

Gala Dinner

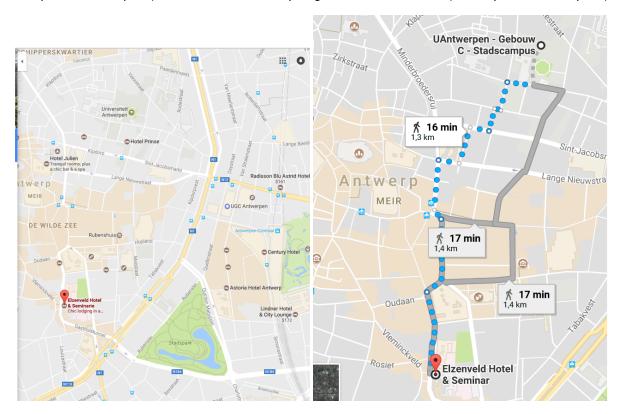
Date: Tuesday, July 18, 2017

Time: 7:30 PM

Venue: Elzenveld Seminar Centre, Lange Gasthuisstraat 45, 2000 Antwerpen http://www.elzenveld.be/en/hotel

Recommended dress code: Casual

How to get there: The easiest way to reach Elzenveld Seminar Centre from the meeting venue is on foot, it takes 18 min (1,5 km). By public transportation, it takes 10 min (i.e. Tramline n. 7 from Antwerpen Keizerstraat direction to Antwerpen Mechelseplein). Tramline n. 4 also stops right in front of the door (Antwerpen Mechelseplein).



Elzenveld Hotel and Conference Center:

The inner courtyard, chapel, former hospital wards, monastery and presbytery, all dating from the 15th to 17th century, now form an oasis of hospitality in Antwerp's city centre. The medieval section of the building has been beautifully restored and adapted to modern needs. Since time immemorial, bodies have been healed and spirits ministered to in this place. Aperitifs will be served in the Chapel, or in the garden if the weather allows, followed by a walking dinner in the Van Gessel Room.



CNS Party

Date: Monday, July 17, 2017 **Time:** 8:30 PM - 00:30 AM

Venue: Party Room HAVN CHURCH, Italielei 8, 2000 Antwerp (www.havnchurch.com)

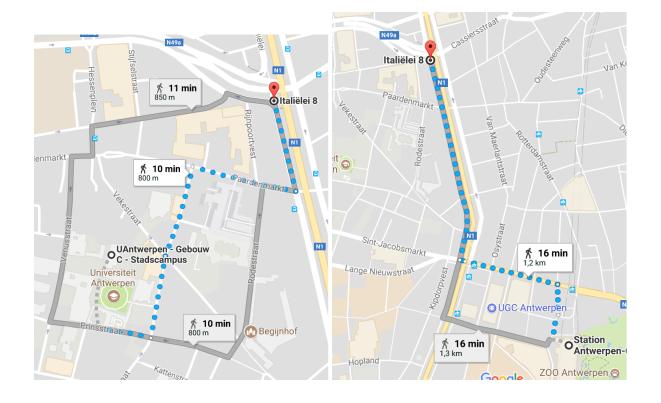
Recommended dress code: Casual



The party will take place in the HAVN CHURCH (http://www.havnchurch.com/). The building was originally used as a Catholic church, built in late Gothic style during the 16th century. It was seized by the Reformed in the 17th century. Today, it is no longer a church and it used as a ballroom, where night parties, banquets and weddings are celebrated.

Each participant will get a voucher for free drinks (beer, wine or soda) for the entire duration of the party. Cava will be served during the first hour and a half of the event. Note that neither (finger) food nor snacks will be served on site.

How to get there: From the meeting venue, it takes 10min (750m) on foot, while from the "Centraal" railway station it takes about 16 min on foot, or 12-15 min by public transportation (i.e. bus n. 720, 730, 770, 771, 776 from Rooseveltplaats Bus Station, direction Antwerpen Sint-Jansplein; or tram n. 12 from Centraal Station, direction Antwerpen Richard).



Restaurants

Selected restaurants close to the venue:

Boston Steak House Burger restaurant De Keyserlei 55, 2018 Antwerpen

Bourla Schouwburg Belgian cuisine Komedieplaats 19, 2000 Antwerpen http://bourlaschouwburg.nu/

Cafe Imperial Belgian-French cuisine Meir 50, 2000 Antwerpen www.cafe-imperial.be

Comocomo Spanish restaurant Kammenstraat 75, 2000 Antwerpen

De Pottekijker Belgian restaurant and grill Kaasrui 5, 2000 Antwerpen www.depottekijker.be

Ellis Gourmet Burger Burger Restaurant Sint-Aldegondiskaai 52, 2000 Antwerpen www.ellisgourmetburger.be

Gran Duca Italian restaurant De Keyserlei 28, 2018 Antwerpen https://www.granduca.be/

Gustav Grills cuisine Van Ertbornstraat 2, 2018 Antwerpen www.brasseriegustav.be

Gusto Belgian-French cuisine Steenhouwersvest 29, 2000 Antwerpen www.mygusto.be

Horta Grand Cafe & Art Nouveau Zaal European restaurant Hopland 2, 2000 Antwerpen www.grandcafehorta.be Korean Barbecue Korean food Statiestraat 8, 2018 Antwerpen

Roma Italian restaurant Statiestraat 19, 2018 Antwerpen

Rubens Inn Belgian restaurant Wapper 17, 2000 Antwerpen www.rubensinn.be

Super Natural Snakbar Minderbroedersrui 47, 2000 Antwerpen

Umi Sushi Asian cuisine Groenplaats 8, 2000 Antwerpen www.umi-sushi.be

Wagamama Japanese food De Keyserlei 13, 2018 Antwerpen

More Antwerp restaurant you can find at: $\underline{\text{http://www.cnsorg.org/cns-2017-local-info}}.$

Program

Tutorials

T1 Subcellular modeling

Room B.001, 15/07/2017, 09:00 - 17:00

<u>Andrew Gallimore</u>, Okinawa Institute of Science and Technology, Japan <u>Weiliang Chen</u>, Okinawa Institute of Science and Technology, Japan

T2 Detailed modeling of structure and function at the cellular level

Room B.002, 15/07/2017, 09:00 - 17:00

Benjamin Torben-Nielsen, Demiurge Technologies AG, Switzerland **Elisabetta lavarone**, Swiss Federal Institute of Technology in Lausanne, Switzerland

T3 Simulation of large-scale neural networks

Room B.003, 15/07/2017, 09:00 - 17:00

<u>Sacha J. van Albada</u>, Julich Research Centre and JARA, Germany Jonas Stapmanns, Julich Research Centre and JARA, Germany

T4 Modeling and analysis of extracellular potentials

Room B.004, 15/07/2017, 09:00 - 12:00

<u>Gaute T. Einevoll</u>, Norwegian University of Life Sciences & University of Oslo, Norway **Espen Hagen**, Dept. of Physics, University of Oslo, Norway

T5 Neuroscience data analysis

Room K.101, 15/07/2017, 13:30 - 17:00

<u>Arvind Kumar</u>, KTH Royal Institute of Technology, Sweden <u>Michael Denker</u>, Julich Research Centre, Germany

T6 Neuroinformatics Resources for Computational Modellers

Room B.004, 15/07/2017, 13:30 - 17:00

Padraig Gleeson, University College London, UK

Andrew P. Davison, French National Center for Scientific Research, France

Main Meeting

Saturday July 15

9:00 –	Registration (Hall Auditorium K.001 Aula Rector Dhanis)
9:00 – 16:30	Tutorials (Aulas B.001, B.002, B.003, B.004, K.101)
17:00 – 17:15	Welcome and announcements (Auditorium K.001 Aula Rector Dhanis)
17.15 – 18:15 K1	Keynote 1: Auditory scene analysis: support and challenges for predictive coding Sue Denham
18:30	Welcome reception (Hall Auditorium K.001 Aula Rector Dhanis)

Sunday July 16

	Culluly Cully 10
9:00 – 9:10	Announcements (Auditorium K.001 Aula Rector Dhanis)
9:10 – 10:10 K2	Keynote 2: Information coding with dendrites: lessons from computational models Panayiota Poirazi
10:10 – 10:40	Break
	Oral session I: Single-cell properties and modeling
10:40 – 11:00 O1	Impact of axon initial segment geometry on excitability Sarah Goethals, Romain Brette*
11:00 – 11:20 O2	Can integrate-and-fire models simulate robust neuromodulation? Tomas van Pottelbergh*, Rodolphe Sepulchre
11:20 – 11:40 O3	Sholl analysis predicted by dendrite spanning fields Alex Bird*, Hermann Cuntz
11:40 – 12:00 O4	Flexible Bayesian inference for complex models of single neurons Pedro Goncalves*, Jan-Matthis Luckmann, Giacomo Bassetto, Marcel Nonnenmacher, and Jakob Macke
12:00 – 13:30	Break for lunch
	Oral session II: Sensory processing: Vision and olfaction
13:30 – 13:50 O5	Learning to read out predictive information in early visual processing Audrey Sederberg*, Jason Maclean, and Stephanie Palmer

13:50 – 14:10 O6	Closed-loop estimation of retinal network sensitivity reveals signature of efficient coding Ulisse Ferrari*, Christophe Gardella, Olivier Marre, and Thierry Mora
14:10 – 14:50 F1	Featured Oral 1: Mixture processing in a biophysical model of the early olfactory system of honeybees Ho Ka Chan*, Thomas Nowotny
14:50 – 15:20	Break
	Oral session III: Synapses and plasticity
	- Car cooler in cyriapood and placticity
15:20 – 15:40 O7	Learning Quantal Parameters through Expectation-Maximization Emina Ibrahimovic*, Martin Müller, and Jean-Pascal Pfister
15:20 – 15:40 O7 15:40 – 16:00 O8	Learning Quantal Parameters through Expectation-Maximization

Monday July 17

9:00 – 9:10	Announcements (Auditorium K.001 Aula Rector Dhanis)
9:10 – 10:10 K3	Keynote 3: Molecular models of the early and late phases of bidirectional plasticity at cerebellar synapses Erik De Schutter
10:10 – 10:40	Break
	Oral session IV: Memory, decisions, and pathological activity
10:40 – 11:00 O9	Cortical correlations support optimal sequence memory Moritz Helias*, Jannis Schuecker, David Dahmen, and Sven Goedeke
11:00 – 11:20 O10	Rats decisions flexibly integrate sensory information and recent history of outcomes Alexandre Hyafil*, Ainhoa Hermoso Mendizabal, Pavel Ernesto Rueda-Orozco, Santiago Jaramillo, David Robbe, and Jaime de La Rocha
11:20 – 11:40 O11	Nicotinic modulation of hierarchal inhibitory circuit control over resting state ultra- slow fluctuations in the prefrontal cortex Marie Rooy*, Fani Koukouli, David Digregorio, Uwe Maskos, and Boris Gutkin
11:40 – 12:00 O12	The minimalistic mathematical model of the cerebral blood flow effects during cortical spreading depression Andrey Verisokin*, Darya V. Verveyko, and Dmitry Postnov
12:00 – 13:30	Break for lunch

Oral session V: Network structure and coherent activity

13:30 – 14:10	F2	Featured Oral 2: Heterogeneous layers stabilize propagation of a multiplexed spike signal in a feed- forward network Dongqi Han*, Sungho Hong	
14:10 – 14:30	O13	Necessity for coherence in motor control Willy Wong*, Omid Talakoub, Robert Chen, and Milos Popovic	
14:30 – 14:50	O14	Dissecting gamma phase and amplitude-specific information routing in V4 macaque during selective attention Dmitriy Lisitsyn*, Eric Drebitz, Iris Grothe, Sunita Mandon, Andreas Kreiter, and Udd Ernst	
14:50 – 15:20		Break	
		Oral session VI: Neural mass models	
15:20 – 15:40	O15	Structure-Function Relationships via Neural Field Theory Peter A Robinson*, Xuelong Zhao, Kevin Aquino, John Griffiths, Grishma Mehta-Pandejee, Natasha Gabay, James Maclaurin, and Somwrita Sarkar	
15:40 – 16:00	O16	Dynamic Operations of Hierarchically Interacting Canonical Microcircuits Tim Kunze*, Jens Haueisen, and Thomas R. Knösche	
16:00 – 19:00		Poster session II: Posters P105 – 209 (Building E)	
19:00		CNS Party (Party Room, Havn Church)	

Tuesday July 18

	ruesday bury 10
9:00 – 9:10	Announcements (Auditorium K.001 Aula Rector Dhanis)
9:10 – 10:10 K4	Keynote 4: I am therefore I think Karl Friston
10:10 – 10:40	Break
	Oral session VII: Large networks and large scale simulations
10:40 – 11:00 O17	Learning structure of 3D objects with cortical columns Subutai Ahmad*, Yuwei Cui, Marcus Lewis, and Jeff Hawkins
11:00 – 11:20 O18	Influence of network topology on spreading of epileptic seizure Simona Olmi*, Spase Petkoski, Fabrice Bartolomei, Maxime Guye, and Viktor Jirsa
11:20 – 11:40 O19	A model-based approach for detecting multiple change points in multivariate spike count data Hazem Toutounji*, Daniel Durstewitz

11:40 – 12:00 O20	Geppetto: an open source visualisation and simulation platform for neuroscience Matteo Cantarelli*, Adrian Quintana, Boris Marin, Matt Earnshaw, Padraig Gleeson, Robert Court, Robert A McDougal, R Angus Silver, Salvador Dura-Bernal, Stephen Larson, William W Lytton, and Giovanni Idili
12:00 – 13:30	Break for lunch
13:30 – 14:30	OCNS Member Meeting
	Oral session VIII: Grid cells and place cells
14:30 – 14:50 O21	Position is coherently represented during flickering instabilities of place-cell cog- nitive maps in the hippocampus Lorenzo Posani*, Simona Cocco, Karel Ježek, and Rémi Monasson
14:50 – 15:30 F3	Featured Oral 3:
	Modeling grid fields instead of modeling grid cells Sophie Rosay*, Tanja Wernle, and Alessandro Treves
15:30 – 16:00	Break
16:00 – 19:00	Poster session III: Posters P210 – P314 (Building E)
19:30	Gala Dinner (Elzenveld Seminar Centre)
	Wednesday July 19 and Thursday July 20
9:00 – 10:30	Workshop Morning Session 1 (Aulas C.101, C.102, K.101, K.102, K.103, K.201, K.202, K.203)
10:30 – 11:00	Break
11:00 – 12:30	Workshop Morning Session 2
12:30 – 14:00	Break for Lunch

Workshop Afternoon Session 1

Workshop Afternoon Session 2

Break

14:00 - 15:30

15:30 - 16:00

16:00 - 18:00

Workshops

W1 Neuronal Oscillations: Mechanisms and Functionality

Room K.202, Wed and Thur 9:00 to 18:00

Horacio G Rotstein, New Jersey Institute of Technology

Frances Skinner, Krembil Research Institute

Vassilis Cutsuridis, University of Lincoln

W2 Methods of Information Theory in Computational Neuroscience

Room C.101, Wed and Thur 9:00 to 18:00

Joseph T. Lizier, University of Sydney

Viola Priesemann, Max Planck Institute for Dynamics and Self-organisation

Justin Dauwels, Nanyang Technological University

Taro Toyoizumi, RIKEN Brain Science Institute

Alexander G Dimitrov, Washington State University

Lubomir Kostal, Academy of Sciences of the Czech Republic

Michael Wibral, Goethe University, Frankfurt

W3 Recent Methods and Analyses for Large-scale Neuronal Population Recordings

Room C.102, Wed and Thur 9:00 to 18:00

Michela Chiappalone, Istituto Italiano di Tecnologia, Genova

Valentina Pasquale, Istituto Italiano di Tecnologia, Genova

Pierre Yger, Institut de la Vision, INSERM, Paris

W4 New Advances in Theoretical Tools for the Study of Large-scale Neural Systems

Room K.101, Wed 9:00 to 18:00 and Thur 9:00 to 12:30

Simona Olmi, Institute of Complex Systems- CNR

David Angulo-Garcia, Aix-Marseille University

Benjamin Lindner, Humboldt University Berlin

W5 Theoretical Neuroscience in the Human Brain Project

Room K.201, Wed 9:00 to 18:00

Michele Giugliano, Universiteit Antwerpen

<u>Alain Destexhe</u>, Centre National de la Recherche Scientifique (CNRS)

Viktor Jirsa, Aix-Marseille University

W6 Computational and Experimental Advances in Cerebellum Research

Room K.102, Wed 9:00 to 18:00

Erik De Schutter, Okinawa Institute of Science and Technology

Yunliang Zang, Okinawa Institute of Science and Technology

W7 Principles and Applications of Extracellular Potentials

Room K.203, Wed 9:00 to 18:00

Michiel Remme, Humboldt University Berlin

Torbjoern Ness, Norwegian University of life Sciences

Gaute Einevoll, Norwegian University of life Sciences - University of Oslo

W8 Fingerprints and Applications of Brain Dynamics Estimated from Neuroimaging Data

Room K.103. Wed 9:00 to 18:00

Matthieu Gilson, University Pompeu Fabra

Tim van Hartevelt, Oxford

W9 Emerging Models in Scientific Communication and Discussion

Room K.201, Thur 9:00 to 18:00

Romain Brette, Institut de la Vision, Paris

W10 Reaction-diffusion Modeling for Neurobiology

Room K.203, Thur 9:00 to 18:00

Robert McDougal, Yale University

William W Lytton, SUNY Downstate

Avrama Blackwell, George Mason

W11 Recent Developments in Epilepsy Modeling

Room K.103, Thur 9:00 to 18:00

Wim van Drongelen, The University of Chicago

Stephan A. van Gils, University of Twente

W12 Neuroscience Gateway: Enabling Developers and Users to Utilize Open High Performance Computing Resources for Large Scale Simulations

Room K.101, Thur 14:00 to 18:00

Amit Majumdar, University of California San Diego, La Jolla

Subhashini Sivagnanam, University of California San Diego, La Jolla

Ted Carnevale, Yale University

W13 Cortical Function: Towards Understanding and Developing Integrative Theories

Room K.102, Thur 14:00 to 18:00

Hamish Meffin, The University of Melbourne

Anthony Burkitt, The University of Melbourne

W14 Postdoc Career Workshop

Room K.102, Thur 9:00 to 12:30

Joanna Jedrzejewska-Szmek, University of Warsaw

Abstracts

Tutorials

T1 Subcellular modeling

Room B.001, 15/07/2017, 09:00 - 17:00

Andrew Gallimore, Okinawa Institute of Science and Technology, Japan Weiliang Chen, Okinawa Institute of Science and Technology, Japan

Many important neural functions are controlled by complex networks of intracellular proteins and signalling molecules. A variety of modular signalling pathways connect and interact to form large networks possessing emergent properties irreducible to individual molecules or pathways. These include bistable and ultrasensitive switches, as well as feedback regulation, and synchronisation. These properties are essential for the induction and regulation of critical neural functions, such as long-term depression and potentiation. The complexity of these networks renders their analysis by inspection alone unfeasible, and we must turn to computational modelling to understand them.

The first half of this tutorial will focus on the structure and function of intracellular networks and deterministic methods for modelling and analysing them. We will use a number of important subcellular pathways to illustrate the key concepts and demonstrate the importance and utility of deterministic methods in their modelling and simulation. We will discuss both the biochemistry of these pathways and their mathematical representation. We will then discuss how these modular pathways connect and interact to form large networks. Important network motifs and their emergent properties will also be explained with specific examples given, as well as mathematical methods for their analysis. We will discuss a number of tools for simulating these differential equation models, but will use the open source software Copasi in the tutorial, owing to its ease of installation and use. Participants will have the opportunity to build and simulate their own signalling pathway model in Copasi. This part of the tutorial will serve as a good introduction to molecular systems modelling for those with little prior experience, and will assume little more than a grasp of basic differential equations and biochemistry.

The second half of the tutorial will focus on more advanced modelling approaches based on several state of the art software packages. We will explain how the time evolution of real molecular systems can diverge from a differential equation-based description due to concepts such as probabilistic interactions in small volumes and spatial heterogeneity. We will describe mathematical approaches to modelling stochastic effects and diffusion and introduce a number of software tools that are based on such descriptions. These include particle-tracking packages such as MCell and Smoldyn, and voxel-based packages such as NeuroRD and STEPS. We will then demonstrate the typical modeling practices with these applications, from model and geometry description to simulation execution and data gathering. Finally, we will briefly discuss recent advances and expected near-future directions of the field.

- [1] STEPS: http://steps.sourceforge.net
- [2] MCell: http://mcell.org/
- [3] Smoldyn: http://www.smoldyn.org/
- [4] NeuroRD: http://krasnow1.gmu.edu/CENlab/software.html
- [5] Antunes, G., and De Schutter, E. (2012). A Stochastic Signaling Network Mediates the Probabilistic Induction of Cerebellar Long-Term Depression. Journal of Neuroscience 32, 9288-9300.
- [6] Bhalla, U.S., and Iyengar, R. (1999). Emergent properties of networks of biological signaling pathways. Science 283, 381-387.
- [7] Eungdamrong, N.J., and Iyengar, R. (2004). Computational approaches for modeling regulatory cellular networks. Trends in Cell Biology 14, 661-669.
- [8] Gallimore, A.R., Aricescu, A.R., Yuzakl, M., and Calinescu, R. (2016). A Computational Model for the AMPA Receptor Phosphorylation Master Switch Regulating Cerebellar Long-Term Depression. Plos Computational Biology 12, 23.

[9] Kotaleski, J.H., and Blackwell, K.T. (2010). Modelling the systems biology approaches. Nature Reviews Neuroscie	molecular mechanisms of synaptic plasticity using ence 11, 239-251.

T2 Detailed modeling of structure and function at the cellular level

Room B.002, 15/07/2017, 09:00 - 17:00

Benjamin Torben-Nielsen, Demiurge Technologies AG, Switzerland **Elisabetta lavarone**, Swiss Federal Institute of Technology in Lausanne, Switzerland

In the morning session, we introduce the morphology of dendrites and axons, the specialised input and output arborisations of neurons. Their shape is pivotal for brain functioning for two reasons: First, overlap between dendrites and axons defines the micro-circuit. Second, the shape and membrane composition of dendrites define how inputs are transformed into relevant outputs. In this tutorial, we will start by explaining the importance of morphologies and how to quantify them (say, in order to distinguish healthy from pathological morphologies). We will touch on algorithmic synthesis of large numbers of unique neuronal morphologies for application in large-scale modelling efforts. We finish the morning session with a hands-on tutorial using btmorph [1] to analyse populations of neuronal morphologies.

In the afternoon session, we explain how neuronal dynamics takes place at the single neuron level and how dendrites turn input signals into an output. We briefly explain the conductance-based and compartmental-modelling paradigms to simulate the dynamics on neurons with detailed membrane composition and elaborate neuronal morphologies. We then proceed to show several free community resources to construct, simulate, share and analyse single neuron models. We will also introduce methods to quantify neurons electrophysiological properties. We end the afternoon session with a hands-on demonstration of how to construct and simulate detailed models of neurons using NEURON and python [2] and on how to constrain their free parameters with experimental data using BluePyOpt [7].

- [1] Torben-Nielsen B. An efficient and extendable Python library to analyze neuronal morphologies. Neuroinformatics 12:619-622, 2014.
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- [3] Torben-Nielsen B., Cuntz H. Introduction to dendritic morphology, The Computing Dendrite, Springer, 2014.
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- [6] Silver A. Neuronal arithmetic. Nature Reviews Neuroscience 11, 474-489, 2010.
- [7] Van Geit W., Gevaert M., Chindemi G., Rossert C., Courcol J. D., Muller E. B., Markram H. (2016). BluePy-Opt: Leveraging open source software and cloud infrastructure to optimise model parameters in neuroscience. Frontiers in Neuroinformatics, 10.
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- [10] Markram H., Muller E., Ramaswamy S., Reimann M. W., Abdellah M., Sanchez C. A., ..., Kahou G. A. A. (2015). Reconstruction and simulation of neocortical microcircuitry. Cell, 163(2), 456-492.

T3 Simulation of large-scale neural networks

Room B.003, 15/07/2017, 09:00 - 17:00

<u>Sacha J. van Albada</u>, Julich Research Centre and JARA, Germany Jonas Stapmanns, Julich Research Centre and JARA, Germany

This tutorial starts with an introduction to large-scale neuronal networks, giving examples of existing models and identifying some challenges these networks pose for modeling and simulation. This is followed by an introduction to the NEural Simulation Tool (NEST [1]), shedding light on its design principles, which address challenges for large-scale simulations. An overview of the features of NEST is provided, also touching upon advanced properties of neuronal networks like gap-junctions [2]. To familiarize participants with the basic usage of NEST, some simple networks are programmed in hands-on exercises. Next, the tutorial explains how NEST enables parallel simulations via both distributed and threaded computations. Threaded simulations are demonstrated on a cortical microcircuit model [3]. Finally, the tutorial provides an introduction to the NEST Modeling Language (NESTML [4]). In this final hands-on part of the tutorial, the participants learn how to create neuron models in NEST using NESTML.

The tutorial does not assume any prior knowledge of NEST. However, it is recommended that participants install NEST on their laptops beforehand [5]. Furthermore, it is recommended to have VirtualBox installed and to have at least 4 GB of free memory available.

- [1] Kunkel S, Morrison A, Weidel P, Eppler JM, Sinha A, Schenck W, Plesser HE. (2017). NEST 2.12.0. Zenodo. http://doi.org/10.5281/zenodo.259534
- [2] Hahne J, Helias M, Kunkel S, Igarashi J, Bolten M, Frommer A, Diesmann M (2015) A unified framework for spiking and gap-junction interactions in distributed neuronal network simulations Front. Neuroinform. 9:22
- [3] Potjans TC, Diesmann M. The Cell-Type Specific Cortical Microcircuit: Relating Structure and Activity in a Full-Scale Spiking Network Model. Cereb. Cortex. 2014;24(3):785-806. DOI: 10.1093/cercor/bhs358.
- [4] Plotnikov D., Rumpe B., Blundell I., Ippen T., Eppler J.M., Morrison A., 2016. NESTML: a modeling language for spiking neurons. arXiv preprint arXiv:1606.02882.
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T4 Modeling and analysis of extracellular potentials

Room B.004, 15/07/2017, 09:00 - 12:00

<u>Gaute T. Einevoll</u>, Norwegian University of Life Sciences & University of Oslo, Norway **Espen Hagen**, Dept. of Physics, University of Oslo, Norway

While extracellular electrical recordings have been one of the main workhorses in electrophysiology, the interpretation of such recordings is not trivial [1,2,3], as the measured signals result of both local and remote neuronal activity. The recorded extracellular potentials in general stem from a complicated sum of contributions from all transmembrane currents of the neurons in the vicinity of the electrode contact. The duration of spikes, the extracellular signatures of neuronal action potentials, is so short that the high-frequency part of the recorded signal, the multi-unit activity (MUA), often can be sorted into spiking contributions from the individual neurons surrounding the electrode [4]. No such simplifying feature aids us in the interpretation of the low-frequency part, the local field potential (LFP). To take a full advantage of the new generation of silicon-based multielectrodes recording from tens, hundreds or thousands of positions simultaneously, we thus need to develop new data analysis methods and models grounded in the biophysics of extracellular potentials [1,3,4]. This is the topic of the present tutorial.

- In the tutorial we will go through the biophysics of extracellular recordings in the brain,
- a scheme for biophysically detailed modeling of extracellular potentials and the application to modeling single spikes [5-7], MUAs [8] and LFPs, both from single neurons [9] and populations of neurons [8,10,11],
- LFPy (LFPy.github.io) [12], a versatile tool based on Python and the NEURON simulation environment [13] (www.neuron.yale.edu) for calculation of extracellular potentials around neurons, and
- new results from applying the biophysical forward-modeling scheme to predict LFPs from comprehensive point-neuron network models, in particular Potjans and Diesmann's model of the early sensory cortical microcircuit using hybridLFPy [14,15] will be presented.

- [1] KH Pettersen et al., Extracellular spikes and CSD, in Handbook of Neural Activity Measurement, Cambridge (2012)
- [2] G Buzsaki et al., Nat Rev Neurosci 13:407 (2012)
- [3] GT Einevoll et al., Nat Rev Neurosci 14:770 (2013)
- [4] GT Einevoll et al., Curr Op Neurobiol 22:11 (2012)
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- [6] J Gold et al., J Neurophysiol 95:3113 (2006)
- [7] KH Pettersen and GT Einevoll, Biophys J 94:784 (2008)
- [8] KH Pettersen et al., J Comp Neurosci 24:291 (2008)
- [9] H Lindén et al., J Comp Neurosci 29: 423 (2010)
- [10] H Lindén et al., Neuron 72:859 (2011)
- [11] S Łęski et al., PLoS Comp Biol 9:e1003137 (2013)
- [12] H Lindén et al., Front Neuroinf 7:41 (2014)
- [13] ML Hines et al., Front Neuroinf 3:1 (2009)
- [14] TC Potjans and M Diesmann, Cereb Cortex 24:785 (2014)
- [15] E Hagen et al., Cereb Cortex 26:4461 (2016)

T5 Neuroscience data analysis

Room K.101, 15/07/2017, 13:30 - 17:00

<u>Arvind Kumar</u>, KTH Royal Institute of Technology, Sweden <u>Michael Denker</u>, Julich Research Centre, Germany

In this tutorial we will explain the theory and practical issues associated with some of the most common tools to analyse spiking activity. Specifically, we will focus on the estimation of firing rate, spike train irregularity, pairwise and higher order correlations, trial-by-trial variability and co-variability, spectrum, dimensionality reduction and estimation (e.g. Gaussian factor analysis). The tutorial will be split in two parts.

In the first part we will provide the theoretical background behind the analysis methods and interpretation of the results. In the second part we will demonstrate how various tools from neuroinformatics, in particular the Python libraries Neo and Elephant, work together in building up robust and reproducible analysis workflows. In addition, we will discuss practical issues related to the analysis methods and interpretation of the results. All through the tutorial we will focus on the spiking activity but most of the methods can be generalized to study other neural data.

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- [3] Averbeck B, Latham PE, Pouget A (2006) Neural correlations, population coding and computation. Nature Reviews Neurosci. 7:358-366 (2006).

T6 Neuroinformatics Resources for Computational Modellers

Room B.004, 15/07/2017, 13:30 - 17:00

Padraig Gleeson, University College London, UK

Andrew P. Davison, French National Center for Scientific Research, France

Neuroinformatics resources are becoming an essential part of computational investigations in neuroscience. A movement towards making data and software freely available to the community means that more and more experimental datasets, general purpose analysis tools and infrastructure for computational modelling and simulation are available for computational neuroscientists to help build, constrain and validate their models. This tutorial will give an overview of the range of neuroinformatics resources currently available to the community. The first half will give a brief introduction to a number of these under the headings: Experimental datasets; Structured data from literature; Analysis tools; Simulation environments; Model sharing; Computing infrastructure; Open source initiatives. The second half of the tutorial will involve hands on exercises where multiple resource will be accessed, data transformed and analysed and new models executed. Note that this tutorial will focus on neuroinformatics resources for cell and network modelling, and not cover the wide range of neuroimaging or genetics databases.

Invited Presentations



Sue Denham School of Psychology, Faculty of Health and Human Sciences, Plymouth University, Plymouth, Devon, UK

K1 – Auditory scene analysis: support and challenges for predictive coding

Perception seems so simple. I look out of the window to see houses, trees, people walking past, the sky above, the grass below. I hear birds in the trees, cars going past, the distant sound of an alarm. The world is full of objects that make their presence known to me through my senses – what could be more simple? Yet the efficacy of perceptual experience hides a host of questions for which we do not yet have the answers. Information reaching our senses is generally incomplete, ambiguous, distributed in space and time and not neatly sorted according to its source, so a key function of our perceptual systems is to discover the likely causes of our sensations. Perception as inference or hypothesis testing, formalised in the predictive coding theory, offers an attractive framework for exploring these issues. From this perspective, regularities or patterns provide perceptual systems with some traction, allowing the formation of expectations and a basis for decomposing the world into discrete objects. But in the dynamic world which we inhabit, object representations must be similarly dynamic, and need to form and dissolve, dominate and yield, in a way that facilitates veridical perception. In this talk I will discuss auditory scene analysis in the context of predictive coding using experimental data, exemplar models, and the phenomenon of perceptual multistability.



Panayiota Poirazi Institute of Molecular Biology and Biotechnology (IMBB), Foundation for Research and Technology-Hellas (FORTH), Heraklion, Crete, Greece

K2 – Information coding with dendrites: lessons from computational models

The goal of this presentation is to provide a set of predictions generated by biophysical and/or abstract mathematical models regarding the role of dendrites in information processing, learning and memory across different brain regions. Towards this goal I will present modelling studies from our lab – along with supporting experimental evidence – that investigate how dendrites may be used to facilitate the learning and coding of both spatial and temporal information at the single cell, the microcircuit and the neuronal network level. I will briefly present early work on how the dendrites of individual CA1 pyramidal neurons may allow a single cell to act as a 2-stage neural network classifier [1], thus massively increasing the storage capacity of the neural tissue [2]. I will then discuss how such dendritic nonlinearities may enable stimulus specificity in individual PFC pyramidal neurons during working memory [3] and underlie the emergence of sustained activity at the single cell and the microcircuit level [3,4]. The role of dendrites in memory phenomena will be assessed using circuit models of the Dentate Gyrus implementing pattern separation [5,6] as well as hippocampal models capable of learning associative memories and linking them across time [7]. This presentation aims to highlight how dendrites are likely to serve as key players in different memory functions.

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- [2] Poirazi P., Mel B.W. (2001) Impact of Active Dendritic Processing and Structural Plasticity on Learning and Memory. Neuron 29: 779-796.
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Erik De Schutter Okinawa Institute of Science and Technology Graduate University, Okinawa, Japan

K3 – Molecular models of the early and late phases of bidirectional plasticity at cerebellar synapses

Synaptic plasticity at the parallel fiber to Purkinje cell synapse has been studied extensively, both experimentally and computationally. The initial focus was on long-term depression (LTD) evoked by concurrent parallel fiber and climbing fiber activation, but more recently experimental studies have emphasized the behavioral importance of long-term potentiation (LTP) triggered by exclusive parallel fiber activation. Expression of these forms of plasticity is based on changes in the number of AMPA receptors in the postsynaptic density (PSD), LTD leading to a decrease and LTP to an increase. As such, this plasticity is bidirectional and can be described as the outcome of a competition by opposing processes. Through studies of hippocampal plasticity we have come to understand the importance of all aspects of the AMPA receptor cycle in bidirectional synaptic plasticity, with LTD increasing both diffusion out of the PSD and endocytosis of receptors and LTP favoring insertion of receptors that diffuse to the PSD. Moreover, the endosomal cycle is quite important because most endocytosed AMPA receptors are rapidly recycled to the postsynaptic membrane. Calcium influx is always the first step in synaptic plasticity, but this influx is brief compared to the tens of minutes required to reach the maximum change in synaptic strength. For cerebellar LTD it is well established that the calcium signal activates a MAP-kinase based positive feedback loop that is essential for the early phase of LTD. We have built a completely new molecular model of bidirectional cerebellar plasticity that replicates experimental findings, including the dual role of nitric oxide in LTP and LTD. LTD requires activation of the MAP-kinase based positive feedback loop and this activation is controlled by CaM kinase. An emergent property of the model is an automatic shutdown of the positive feedback loop, corresponding to the end of the early phase. In a second, simpler model, we have explored how the early phase can transition into a stable late phase by simple manipulations of the endosomal cycle. Unfortunately, experimental data on these processes is less complete, particularly about possible spatial restriction to single spines.



Karl Friston Wellcome Trust Centre for Neuroimaging, Institute of Neurology University College London, London, UK

K4 - I am therefore I think

This overview of the free energy principle offers an account of embodied exchange with the world that associates neuronal operations with actively inferring the causes of our sensations. Its agenda is to link formal (mathematical) descriptions of dynamical systems to a description of perception in terms of beliefs and goals. The argument has two parts: the first calls on the lawful dynamics of any (weakly mixing) ergodic system – from a single cell organism to a human brain. These lawful dynamics suggest that (internal) states can be interpreted as modelling or predicting the (external) causes of sensory fluctuations. In other words, if a system exists, its internal states must encode probabilistic beliefs about external states. Heuristically, this means that if I exist (am) then I must have beliefs (think). The second part of the argument is that the only tenable beliefs I can entertain about myself are that I exist. This may seem rather obvious; however, it transpires that this is equivalent to believing that the world – and the way it is sampled – will resolve uncertainty about the causes of sensations. We will consider the implications for functional anatomy, in terms of predictive coding and hierarchical architectures, and conclude by looking at the epistemic behaviour that emerges using simulations of active inference.

Contributed Talks

F1 Mixture processing in a biophysical model of the early olfactory system of honeybees

Ho Ka Chan*, Thomas Nowotny

School of Engineering and Informatics, University of Sussex, Brighton, UK

In their natural environment, Insects often encounter complex mixtures of odors in their natural environment. It is an important open question whether and how the processing of complex mixtures of multi-component odors differs from that of simpler mixtures or single components. To approach this question, we built a full-size model of the early olfactory system of honeybees, which predicts responses to both single odorants and mixtures. The model is designed so that olfactory response patterns conform to the statistics derived from experimental data [1] for a variety of its properties. It also takes into account several biophysical processes at a minimal level, including processes of chemical binding and activation in receptors, and spike generation and transmission in the antennal lobe network. We verify that key findings from other experimental data not used in building the model [2-4] are reproduced in it. In particular, we replicate the strong correlation among receptor neurons and the weaker correlation among projection neurons observed in experimental data [1,2] and show that this decorrelation is predominantly due to inhibition by interneurons. By simulation and mathematical analysis of our model, we demonstrate that the chemical processes of receptor binding and activation already lead to significant differences between the responses to mixtures and those to single component stimuli. On average, the response latency of olfactory receptor neurons at low stimulus concentrations is reduced (See Figure 1A) and the response patterns become less variable across concentrations (See Figure 1B) as the number of odor components in the stimulus increases. These effects are preserved in the projection neurons. Our results provide hints that the early olfactory system in insects may be particularly efficient in processing mixtures, which corresponds well to the observation that chemical signaling in nature predominantly utilizes mixtures.

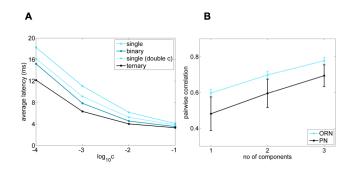


Figure 1: A. The average response latency decreases with the number of components in the odor stimulus. The effect is most significant when the stimulus concentration is low. B. The pairwise correlation, averaged over all ORNs, between the response patterns at low and high concentration increases with the number of components in the odor stimulus.

Acknowledgements

This work was supported by HFSP, RGP0053/2015 and EPSRC, EP/J019690/1.

- [1] Galizia CG, Sachse S, Rappert A, Menzel R: **The glomerular code for odor representation is species specific in the honeybee Apis mellifera**. *Nat Neurosci* 1999 May, 2(5):473-478.
- [2] Ditzen M: Odor concentration and identity coding in the antennal lobe of the honeybee Apis mellifera [PhD thesis]. *Berlin: Freie Universitat Berlin* 2005.
- [3] Szyszka P, Gerkin RC, Galizia CG, Smith BH: **High-speed odor transduction and pulse tracking by insect olfactory receptor neurons**. *Proc Natl Acad Sci U S A* 2014,111(47):16925-16930.
- [4] Deisig N, Giurfa C, Sandoz JC: **Antennal lobe processing increases separability of odor mixture representations in the honeybee**. *J Neruophysiol* 2010,103:2185-21.

F2 Heterogeneous layers stabilize propagation of a multiplexed spike signal in a feedforward network

Dongqi Han¹*, Sungho Hong²

Feedforward networks are ubiquitous structures in neural systems and have been studied in many contexts such as models for signal transmission [1,2], architectures for rich information processing [3], etc. However, most studies have ignored an important property commonly observed in real feedforward networks: neurons in one layer have contrasting characteristics from those in other layers. For example, the cerebellar granule cells are tiny and relatively simplistic neurons while their postsynaptic targets, the Purkinje cells, are much bigger, complex, and therefore have very different intrinsic properties. What would be the role of such layer-to-layer differences in neural circuits?

Here we address this question by simulation of a model feedforward network, inspired by a recent experimental study on the Drosophila olfactory system [4]. In this model, all the adjacent layers have Morris-Lecar neurons with different excitability types from each other and therefore different computational functions. If one layer has cells with class I excitability, which behave like integrators of inputs, neurons in the adjacent layer are of class III, which act as coincidence detectors [5], and vice versa.

We found that spikes from one layer evoked a response in next layer neurons better when they had the same excitability type. However, in a deep feedforward network, this caused gradual accumulation of signal distortion, leading to the undesirable responses in deep layers that all the neurons either fired synchronously or became silent, as seen in classical studies (e.g., [1]). On the other hand, the network with heterogeneous layers demonstrated a novel signal transformation property as observed in [4] (Fig. 1A), and showed stable propagation of a signal into deep layers with a preserved temporal fidelity and spike count (Fig. 1B, C). We analyzed the result by using a phase space method in [1] and showed how mixing different coding schemes enables this feature (Fig. 1C). We conclude that heterogeneous layers in feedforward neural networks can be a mechanism for optimal information transfer.

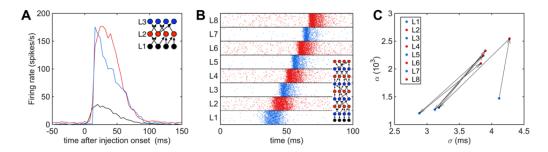


Figure 1: A. Firing rate of input (black, L1), integrator (red, L2), and coincidence detector (blue, L3) neurons. Note that a peak of L3 firing precedes that of L2, as observed in [4]. B. Stable spike propagation in a network with many heterogeneous layers. C. Layer-to-layer change in the SD of spike times and spike count of B.

- [1] Diesmann M, Gewaltig M-O, Aertsen A: **Stable propagation of synchronous spiking in cortical neural networks**. *Nature* 1999, 402:529-33.
- [2] Kumar A, Rotter S, Aertsen A: **Spiking activity propagation in neuronal networks: reconciling different perspectives on neural coding**. *Nat Rev Neurosci* 2010, 11:615-27.
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- [4] Jeanne JM, Wilson RI: Convergence, Divergence, and Reconvergence in a Feedforward Network Improves Neural Speed and Accuracy. *Neuron* 2015, 88:1014-26.
- [5] Ratte S, Hong S, De Schutter E, Prescott SA: Impact of neuronal properties on network coding: roles of spike initiation dynamics and robust synchrony transfer. *Neuron* 2013, 78:758-72.

¹ Graduate School, Okinawa Institute of Science and Technology, Okinawa, Japan

²Computational Neuroscience Unit, Okinawa Institute of Science and Technology, Okinawa, Japan

F3 Modeling grid fields instead of modeling grid cells

Sophie Rosay¹*, Tanja Wernle², and Alessandro Treves¹

Grid cells are neurons found in the rodent medial entorhinal cortex. They take their name from their astonishing firing patterns: grid cells are active specifically in certain regions of physical space, called grid fields, that form a triangular grid tesselating the space explored by the animal. While experiments have been investigating the geometrical properties of grid fields [1], computationalists have tried to explain them by neural network models. However, these models still fail to account for some of the experimental results, in particular how two distinct grid patterns are integrated when two compartments are merged into one (Wernle et al, in preparation).

We take a different approach: instead of modelling grid cells, we directly model single grid fields as point objects interacting with each other and with the environment's borders (see Figure 1A). This description is motivated by the way grid patterns are naturally considered as geometrical objects. We thus consider a system of interacting objects evolving as colloidal particles on a substrate [2]. First we consider only grid fields from one grid cell, then we add coupling between several cells. We simulate the system with varying forms and intensity of the interactions. The simplicity of the model allows us to test many such possibilities and their outcome in several 'experimental' setups.

We show that under certain conditions the model does reproduce the behavior of experimental grid fields (see Figure 1B). These conditions imply repulsion between fields, the involvement of a large number of fields, interaction between grid cells and with the walls. We are able to reproduce observed data from experiments in merged environments. We can also make predictions for setups not tested experimentally yet. The question that then naturally arises is how to connect our description at the level of grid fields to models at the level of grid cells. We show how our grid-field model puts constraints on models of the underlying grid cells. Conversely, we discuss how existing grid-cell models can be described at the level of grid fields.

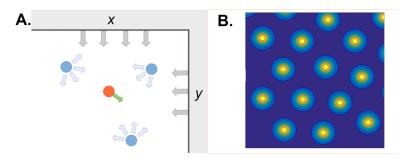


Figure 1: General idea of grid field modeling. A. Schematic representation of the model. A given grid field (red) feels the influence of other grid fields (blue) as well as the borders (grey) plus a viscosity force (green). B. Example of a resulting pattern in a square box, converted into a firing rate map.

In conclusion, tackling the issue of grid patterns from a grid-field perspective provides new insights on their formation. Beyond grid cells, our work raises the question of the ultimate purpose of a model and the subtle interplay between description and explanation.

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¹ Cognitive Neuroscience Area, Scuola Internazionale Superiore di Studi Avanzati, Trieste, Italy

²Kavli Institute for Systems Neuroscience, NTNU, Trondheim, Norway

O1 Impact of axon initial segment geometry on excitability

Sarah Goethals, Romain Brette*

Institut de la Vision, Sorbonne Universités, UPMC Univ Paris 06, INSERM, CNRS, Paris, France

In most vertebrate neurons, action potentials are triggered at the distal end of the axon initial segment (AIS). They are then transmitted to the soma where they are regenerated and further propagated in the dendritic tree. The AIS position and length can be altered by changes in electrical activity, suggesting a strong link between AIS geometry and excitability. We studied theoretically the influence of AIS geometry on the somatic threshold for AP initiation. For this purpose, we solved the cable equation with appropriate boundary conditions in a cylindrical axon model. Our theoretical analysis shows that the somatic threshold depends logarithmically on the surfacic sodium conductance density and that increasing either the AIS length or the AIS start position lowers the threshold. We confirmed our prediction with numerical simulations in a more detailed neuron model. Our analysis suggests that either a longer or a more distal AIS increases excitability, which supports a current hypothesis that the AIS is preferably isolated from the large capacitance of the soma. Secondly, we examined how the AIS geometry influences the peak axonal current that is transmitted to the soma at spike initiation. Again we used cable analysis to study this current in a two-cylinder model that represents the main geometrical features of a thick-tufted layer 5 pyramidal neuron. Our analysis shows that in order to obtain somatic spikes with a given speed, the AIS position should be proportional to the diameter of the apical dendrite raised to the -3/2. We confirmed this theoretical result with numerical simulations of a more detailed model. In addition, correlation analysis of layer 5 pyramidal neurons morphology confirms this theoretical prediction [1]. Our previous analyses suggest that the AIS geometry is finely tuned for successful spike transmission to the soma. More generally, neural systems tend to be efficient in their use of resources [2], which suggests that AIS geometry might also be optimized for minimal energy consumption. As the energy consumption at subthreshold voltages is proportional to the number of channels, we asked whether there exists an AIS geometry that minimizes the total number of sodium channels. For this purpose, we used variational techniques to calculate the AIS geometry that minimizes total Na conductance, for a given spike threshold and axonal current.

Conclusion

Our theoretical analysis shows that AIS geometry has a strong impact on several aspects of excitability including energy consumption, which suggests that the AIS morphology is functionally tuned and possibly optimized.

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O2 Can integrate-and-fire models simulate robust neuromodulation?

Tomas van Pottelbergh*, Rodolphe Sepulchre

Department of Engineering, University of Cambridge, Cambridge, Cambridgeshire, UK

By controlling the state of neuronal populations, neuromodulators ultimately affect behaviour. A key neuromodulation mechanism is the alteration of neuronal excitability via the modulation of ion channel expression. This type of neuromodulation is normally studied via conductance-based models, but those models are computationally challenging for large-scale network simulations needed in population studies. Integrate-and-fire models provide a computationally advantageous alternative, but such models are only partially successful in robustly capturing modulation between firing patterns.

In this work, we propose a modelling framework that extracts the qualitative properties of neuromodulation to produce neuromodulable and computationally efficient neuron models. Our framework is based on dynamic I-V curves, i.e. instantaneous I-V curves in a certain timescale [1]. These dynamic I-V curves make the connection between qualitative conductance-based models and integrate-and-fire models: how a change in ion channel conductance can be related to a change of dynamic I-V curves and subsequently to a parameter change in the reduced integrate-and-fire model. We focus on the modulation between tonic firing and bursting as an example. We argue that this modulation crucially relies on the co-regulation of two points of high sensitivity (i.e. excitability) in two distinct timescales. The points of high sensitivity are local extrema in the I-V curves and correspond to an exact balance of positive and negative feedback. Those signatures have a direct correlate in the fast-slow phase portraits: a hysteretic V-nullcline in the presence of one (fast) balance, and a mirrored hysteresis in the presence of both a fast and a slow balance [2]. The classical quadratic integrate-and-fire model captures the fast balance, but does not account for the slow one.

The simple idea underlying the proposed multi-quadratic integrate-and-fire model (MQIF) is to allow for several distinct balance points in an integrate-and-fire model. An integrate-and-fire model with two balance points is shown to robustly capture the neuromodulation between spiking and bursting, opening novel computational avenues for large-scale simulation of neuromodulated populations. The robustness and modulation properties of this integrate-and-fire model are in sharp contrast to those of existing (generalised) integrate-and-fire models, which lack the slow excitability.

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O3 Sholl analysis predicted by dendrite spanning fields

Alex Bird1,2*, Hermann Cuntz1,2

Sholl analysis has been an important technique in dendritic anatomy for over sixty years [1]. In counting the number of dendritic branches at a given distance from the soma, the Sholl intersection profile is often taken as a crucial measure of dendritic complexity; it has been used in a broad range of applications, from estimating the expected number of possible synapses [2], to evaluating the changes in structure induced by pathologies [3].

We have shown that Sholl intersection profiles can be predicted by two more basic measures: the domain spanned by the dendritic arbor and the angular distribution of how far dendritic segments deviate from a direct path to the soma (Figure 1C). The first measure is principally determined by axon location and hence microcircuit structure [4]; the second arises from optimal wiring [5]. These two measures allow Sholl analysis to be given a more functional interpretation across all of its applications.

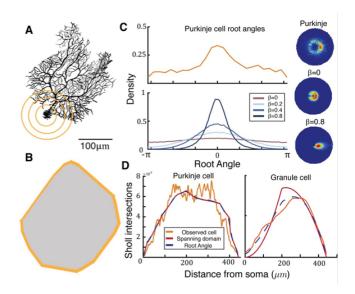


Figure 1: The dendrite spanning field predicts the Sholl intersection profile of a Purkinje cell. A. Rat Purkinkje cell [6] and schematic of Sholl analysis: the number of times the dendrite intersects with a gold arc gives the value of the Sholl intersection profile at that radius. B. The spanning field of the above cell. C. Root angle distributions and (inset) joint angular and euclidean connection probabilities for the Purkinje cell (top) and artificial dendrites with different balances between wiring and delay costs (bottom). D. Sholl intersection profiles for rat Purkinje cell (left) and mouse dentate gyrus granule cell [7] (right): observed (gold), predicted by just the spanning field (red), and predicted by the spanning field and root angles (blue dashed).

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¹ Ernst Strüngmann Institute for Neuroscience, Max Planck Society, Frankfurt, Germany

² Frankfurt Institute for Advanced Studies, Frankfurt, Germany

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O4 Flexible Bayesian inference for complex models of single neurons

Pedro Goncalves^{1*}, Jan-Matthis Luckmann¹, Giacomo Bassetto¹, Marcel Nonnenmacher¹, and Jakob Macke^{1,2,3}

Characterizing the input-output transformations of single neurons is critical for understanding neural computation. Single-neuron models have been extensively studied, ranging from simple phenomenological models to complex multi-compartment neurons. However, linking mechanistic models of single-neurons to empirical observations of neural activity has been challenging. Statistical inference is only possible for a few neuron models (e.g. GLMs), and no generally applicable, effective statistical inference algorithms are available: As a consequence, comparisons between models and data are either qualitative or rely on manual parameter tweaking, parameter-fitting using heuristics or brute-force search [1]. Furthermore, parameter-fitting approaches typically return a single best-fitting estimate, but do not characterize the entire space of models that would be consistent with data (the posterior distribution).

We overcome this limitation by presenting a general method to infer the posterior distribution over model parameters given observed data on complex single-neuron models. Our approach can be applied in a 'black box' manner to a wide range of single-neuron models without requiring model-specific modifications. In particular, it extends to models without explicit likelihoods (e.g. most single-neuron models). We achieve this goal by building on recent advances in likelihood-free Bayesian inference [2]: the key idea is to simulate multiple data-sets from different parameters, and then to train a probabilistic neural network which approximates the mapping from data to posterior distribution.

We illustrate this approach using single- and multi-compartment models of single neurons: On simulated data, estimated posterior distributions recover ground-truth parameters, and reveal the manifold of parameters for which the model exhibits the same behaviour. On in-vitro recordings of membrane voltages, we recover multivariate posteriors over biophysical parameters, and voltage traces accurately match empirical data. Our approach will enable neuroscientists to perform Bayesian inference on complex neuron models without having to design model-specific algorithms, closing the gap between biophysical and statistical approaches to single-neuron modelling.

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¹Neural Systems Analysis, research center caesar, an associate of the Max Planck Society, Bonn, Germany

²Max Planck Institute for Biological Cybernetics, Tübingen, Germany

³Bernstein Center for Computational Neuroscience, Tübingen, Germany

O5 Learning to read out predictive information in early visual processing

Audrey Sederberg^{1,2,3*}, Jason Maclean^{2,4}, and Stephanie Palmer^{3,4}

To generate appropriate behavior, the brain must predict the future state of the world from past sensory information. Taking the salamander visual system as an example, at the minimum such predictions need to compensate for the 50-80 ms processing time of the retina [1] as well as the time for a motor response to be generated. Making these predictions requires leveraging the spatiotemporal structure of the natural world, a computation that is performed efficiently at the first stage of visual processing, in populations of retinal ganglion cells [2]. Neurons downstream of the retina infer predictions about object motion from the firing of their inputs, but to do so, downstream cells must learn to read out predictive information from the retinal activity.

More concretely, stimulus predictive information in a sensory population is defined as the mutual information of particular patterns of spiking across retinal ganglion cells (RGCs) with the future stimulus [2]. We consider the output of a downstream model neuron that receives weighted inputs from several RGCs. By constructing moving-bar dynamics that contain both predictable and stochastic motion, we can visualize the predictive information in readout spiking as the difference between the prior stimulus distribution (Figure 1A, gray) and the spike-triggered stimulus distribution: the larger the difference, the more informative the spike. In this example, the readout neuron was informative of the future stimulus (Figure 1A), but most other readouts were not (not shown). Even for an experimenter with a well-controlled sensory input, finding this optimal readout in the space of all possible readouts is difficult. In more realistic circumstances, the organism must make predictions about the future state of complex natural stimuli from the retinal spiking activity alone (Figure 1B).

Here we address whether biologically plausible learning rules can find readout weights that transform RGC input into predictive downstream output. Input activity from the RGCs was previously recorded in the salamander retina in response to a natural movie (Figure 1B). We first show that internal predictive information, the information that the readout has about its own future input, is correlated with stimulus predictive information, so that becoming more predictive of its inputs drives the readout neuron to encode more information about the future stimulus. Starting from a set of random weights connecting the RGCs to the readout neuron, we allow the weights to evolve via spike timing-dependent plasticity. Across many groups of RGCs, we find that learned readouts convey 80% of the possible predictive information for groups of four cells, but only 30% for groups of ten cells. This decrease reflects a compressibility limit of predictive information and suggests an optimal pooling size for cells downstream from retina.

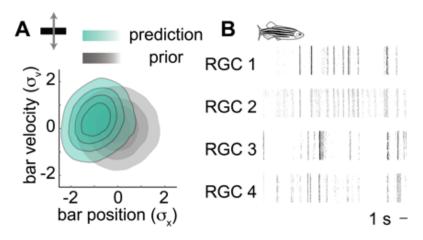


Figure 1: A. Spike-triggered average of future stimulus position and velocity for a particular readout of population spiking activity with high predictive information. B. Raster plots for simultaneously recorded retinal ganglion cells (RGCs) in response to a naturalistic movie featuring swimming fish. These are used to drive learning simulations.

Acknowledgements

¹Biomedical Engineering Department, Georgia Institute of Technology, Atlanta, GA, USA

²Department of Neurobiology, University of Chicago, Chicago, IL, USA

³Department of Organismal Biology and Anatomy, University of Chicago, Chicago, IL, USA

⁴Committee on Computational Neuroscience, University of Chicago, Chicago, IL, USA

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O6 Closed-loop estimation of retinal network sensitivity reveals signature of efficient coding

Ulisse Ferrari¹*, Christophe Gardella^{1,2}, Olivier Marre¹, and Thierry Mora²

According to the theory of efficient coding, sensory systems are adapted to represent natural scenes with high fidelity and at minimal metabolic cost. Testing this hypothesis for sensory structures performing non-linear computations on high dimensional stimuli is still an open challenge. Here we develop a method to characterize the sensitivity of the retinal network to perturbations of a stimulus. Using closed-loop experiments, we explore selectively the space of possible perturbations around a given stimulus. We then show that the response of the retinal population to these small perturbations can be described by a local linear model. Using this model, we computed the sensitivity of the neural response to arbitrary temporal perturbations of the stimulus, and found a peak in the sensitivity as a function of the frequency of the perturbations. Based on a minimal theory of sensory processing, we argue that this peak is set to maximize information transmission. Our approach is relevant to testing the efficient coding hypothesis locally in any context where no reliable encoding model is known.

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¹Institut de la Vision, Sorbonne Universités, INSERM, Paris, France

²Laboratoire de Physique Statistique, Ecole Normale Supérieure, Paris, France

O7 Learning Quantal Parameters through Expectation-Maximization

Emina Ibrahimovic^{1,2*}, Martin Müller¹, and Jean-Pascal Pfister²

Large synapses, i.e the neuromuscular junction (NMJ) or the calyx of Held, have been invaluable model synapses that have significantly advanced the field of synaptic transmission. No generative model approach can faithfully retrieve quantal parameters from synapses with a large number of release sites (N). Here we propose an expectation maximization (EM) method that is based on particle smoothing (PS) to extract quantal parameters from large N synapses. In contrast to an existing EM-based approach [1], using Baum-Welch (BW) which scales with a complexity of N^4 and hence cannot retrace quantal parameters of synapses with hundreds of release sites, our method is independent of N and therefore suitable for large synapses. First, our model was validated on synthetic data. As shown in figure 1, all parameters $\Theta = N, p, q, \sigma, \tau$ were faithfully retrieved. Next, we applied the model to the Drosophila NMJ, which is predicted to harbor hundreds of release sites. The model predicted quantal parameters that are in line with parameters predicted by two empirical approaches (variance-mean analysis and cumulative amplitude analysis of postsynaptic currents). In contrast to these two techniques, which require data recorded under conditions of high release probability (p), our method is independent of p or stimulation protocol. Given the genetic tractability of this synapse, our theoretical approach is expected to help linking quantal parameters to molecular function.

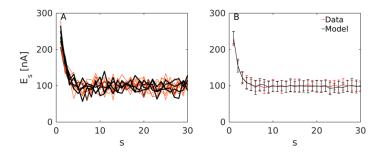


Figure 1: A. Postsynaptic currents at the *Drosophila* NMJ, 50 trains of 30 presynaptic action potentials at 60 Hz compared with model generated data from the fitted parameters at each stimulation step s; number of release sites N=710, release probability p=0.44, quantal content q=0.74 nA , background noise $\sigma=13.22$ nA and refilling time constant $\tau=59$ ms. B. Comparison of the mean and standard deviation.

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¹Institute of Molecular Life Sciences, University of Zürich, Switzerland

²Institute of Neuroinformatics, University of Zürich/ETH Zürich, Switzerland

O8 Emergence of disparity selective neurons through spike-based learning from naturalistic stereoscopic datasets

Tushar Chauhan^{1,2*}, Timothee Masquelier^{1,2}, Alexandre Montlibert^{1,2}, and Benoit Cottereau^{1,2}

Models such as sparse coding [1] have shown that natural scene statistics can be used to predict basis units with Gabor-like receptive fields close to those observed in V1 simple-cells. Since the inputs to these models are natural images captured using a single camera, their outputs are monocular. Recently, attempts have also been made to exploit statistics of stereo-images of natural scenes [2]. The resulting bases units show binocular, Gabor-like receptive fields with population statistics close to those observed in V1. Although these models are able to replicate certain aspects of V1 binocular populations, they are either supervised, or mimic the result of learning from natural datasets, but not the process.

We propose a novel method of deriving monocular and binocular units through unsupervised learning from natural stereoscopic datasets using spike-timing-dependent-plasticity (STDP). Using the Hunter-Hibbard dataset [2], we first employed ON/OFF-center difference-of-Gaussian convolutions to mimic LGN responses (Figure 1.A). The responses were thresholded and converted to spike-latencies using a monotonically decreasing function. This ensured that the most activated units fired first, while the less active units fired late, or not at all. We then trained an STDP based neural network using 1x1 degree spatial pools from the aforementioned LGN layer. The network was composed of integrate-and-fire neurons and incorporated a lateral inhibition scheme. Finally, we characterized the receptive field in each eye by fitting Gabor functions. Our results (Figure 1.B) showed that most units developed Gabor-like receptive fields similar to those observed in V1 simple cells, with a continuum of ocular dominance from pure monocularity to perfect binocularity. In line with single-unit recordings in primates, disparity selectivity was principally observed along the horizontal dimension, where it ranges between -0.5 and 0.5°. Neurons also showed selectivity to vertical disparity, although it was less pronounced. When tested with phase-shifted sine gratings, the units also showed disparity-tuning curves similar to those observed in the cat visual system.

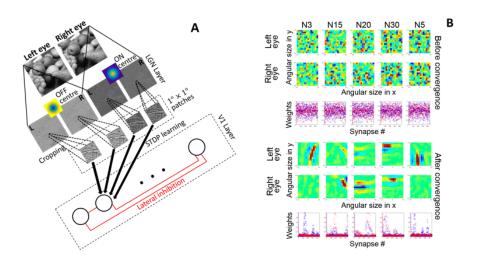


Figure 1: A. Schematic of the processing pipeline. B. Five representative neurons (one per column) before and after convergence. Rows 1,2: Left and right eye receptive fields before convergence; Rows 4,5: The corresponding receptive fields after convergence; Rows 3,6: Weights before and after convergence.

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¹ Université de Toulouse, Centre de Recherche Cerveau et Cognition, Toulouse, France

²Centre National de la Recherche Scientifique, Toulouse Cedex, France

O9 Cortical correlations support optimal sequence memory

Moritz Helias^{1,2*}, Jannis Schuecker¹, David Dahmen¹, and Sven Goedeke¹

The brain processes time-varying input, but is it not known if its dynamical state is optimal for this task. Indeed, recurrent and randomly coupled networks of rate neurons display a rich internal dynamics near the transition to chaos [1], which has been associated with optimal information processing capabilities [2, 3, 4]. In particular, the dynamics becomes arbitrarily slow at the onset of chaos similar to critical slowing down. The interplay between time-dependent input signals, network dynamics, and the resulting consequences for information processing are, however, yet poorly understood.

We here present a completely solvable model that allows us to investigate the effect of time-varying input on the transition to chaos. We analytically obtain the phase diagram spanned by the coupling strength and the input amplitude: External drive shifts the transition to chaos to significantly larger coupling strengths than predicted by linear stability analysis. The intermediate regime is absent in time-discrete networks [5] and only exists in their more realistic time-continuous counterparts. This novel dynamical regime combines locally expansive dynamics with asymptotic stability. We investigate sequential memory [5] and analytically show that memory capacity is optimal within the novel regime. Because it is unclear if cortex operates in such a computationally beneficial regime, we develop a finite-size mean-field theory which relates the statistics of measured covariances to the statistics of connections, in particular the spectral radius of the connectivity matrix. The theory shows that the large dispersion of spike count covariances across pairs of neurons, observed in massively parallel recordings, is an indicator that cortex indeed operates close to the breakdown of linear stability (see Figure 1).

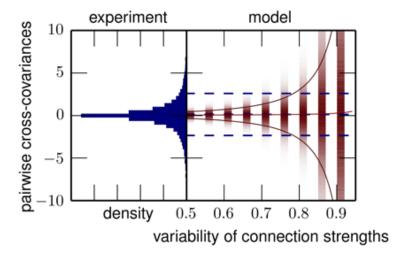


Figure 1: Distribution of spike count cross-covariances across neurons in macaque motor cortex. The low mean and large standard deviation (blue dashed horizontal lines) of experimentally observed cross-covariances between spike counts (left) are explained by a model network (right, shading indicates density of histogram) with a large spectral radius ($R \approx 0.9$) of the connectivity matrix. Red curves: analytical prediction for mean and standard deviation. Data from 155 neurons mostly located in layer 5 of macaque motor cortex (M1). Data courtesy of A. Riehle and T. Brochier.

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¹Institute of Neuroscience and Medicine (INM-6) & Institute for Advanced Simulation (IAS-6), Jülich Research Centre, Germany

²Department of Physics, Faculty 1, RWTH Aachen University, Germany

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O10 Rats decisions flexibly integrate sensory information and recent history of outcomes

Alexandre Hyafil^{1,2*}, Ainhoa Hermoso Mendizabal¹, Pavel Ernesto Rueda-Orozco³, Santiago Jaramillo⁴, David Robbe⁵, and Jaime de La Rocha¹

Animal decisions not only reflect current sensory information but are also shaped by recent experience. There is however little understanding about the determinants of these history-dependent decision biases. We used rats in a novel two-alternative forced choice auditory discrimination task, in which the probability to repeat the previous stimulus category was varied in blocks of trials. Rats adapted to this environment by developing a strategy that capitalized on the serial correlations of the stimulus sequence: a bias towards repeating the same response built up after correct repetitions, and conversely an alternation bias developed after correct alternations. Strikingly, both repetition and alternation biases disappeared after an incorrect trial, irrespective of the number of previous correct trials performed previously.

A GLM analysis revealed that rats decisions in each trial relied on: (1) the current sensory stimulus; (2) a lateral bias towards (away from) the side of recently rewarded (unrewarded) responses on the last 5-10 trials, i.e. win-stay-lose-switch strategy; (3) a novel and strong transition bias that reinforced recent correct transitions (repetitions vs. alternations). Intriguingly the transition bias had no impact on choice after error trials. Subsequent analysis showed that the value of the bias was not reset but simply ignored after an error, and it was recovered after the first subsequent correct trial. Thus, the weight of the history-dependent transition bias could be flexibly and transiently put aside after error choices when possibly the reliability of the internal model was questioned. This nonlinear effect could not be captured by the GLM fitted to both correct and incorrect trials and was not present on the lateral bias, i.e. it was specific of the transition bias. We thus built a latent generative model of rats decisions, whereby lateral and transitions biases are updated at each trial, while the influence of the latter on current decisions is gated by a reward-dependent confidence signal. When fitted to the data, the model accounted quantitatively for all described behavioral effects: in particular, the absence of a transition bias after incorrect choices was due to a reset of the confidence signal. Because the value of the transition bias did not reset after errors but it kept the information about whether the animal would repeat or alternate, a single correct trial was sufficient to increase the confidence and recover the accumulated choice bias. Overall, we show that history-dependent biases in rodent perceptual choices reflect consistent strategic adaptations to behavioural outcomes.

¹ Idibaps, Barcelona, Spain

²Center for Brain and Cognition, Barcelona, Spain

³Universidad Nacional Autónoma de Mexico, Mexico

⁴Institute of Neuroscience, University of Oregon, Oregon, USA

⁵Inmed, Marseille, France

O11 Nicotinic modulation of hierarchal inhibitory circuit control over resting state ultra-slow fluctuations in the prefrontal cortex: modeling of genetic modification and schizophrenia-related pathology

Marie Rooy¹[⋆], Fani Koukouli^{2,3}, David Digregorio^{3,4}, Uwe Maskos^{2,3}, and Boris Gutkin^{1,5}

The prefrontal cortex (PFC), key for higher order cognitive processes, exhibits spontaneous activity that is altered in schizophrenia [1]. Cortical acetylcholine (ACh) release modulates PFC activity via nicotinic acetylcholine receptors (nAChRs) [2] specifically expressed within a hierarchical circuit of inhibitory neurons within layer II/III [3]. Parvalbumin (PV) interneurons, expressing α 7 nAChRs subunits [2], target pyramidal cells axosomaticaly, exerting divisive effects on their activity. Somatostatin (SOM) interneurons, expressing both $\alpha 7$ and $\beta 2$ nAChRs subunits [2], target the dendrites of pyramidal cells, exerting substractive inhibition [4]. The α 5 nAChRs subunits are expressed only by vasoactive intestinal polypeptide (VIP) interneurons, that preferentially inhibit the SOM cells. In vivo two-photon imaging showed that neural activity of PFC in mice is characterized by synchronous ultra-slow fluctuations, with alternating periods of high and low activity [5]. Genetic deletion of specific nAChRs subunits disrupted these ultra-slow fluctuations, leading to changes in synchrony and duration of activity states. Furthermore, mice expressing a human polymorphism in the α 5 nAChRs subunits (α 5SNP) associated with high risk for nicotine addiction and schizophrenia [6, 7], show reduced spontaneous activity in the PFC that is reversed by nicotine [3]. Using a circuit modeling approach, we studied the roles of distinct GABAergic interneurons in the generation of synchronous ultra-slow fluctuations. In order to study the effects of substractive vs. divisive inhibition on bistable dynamics in the pyramidal neuron, by the SOM and PV interneuron populations respectively, we used population firing rate modelling incorporating both mechanisms [8], and simulated the effects of nAChRs knock outs. With our model, we could fully account for the changes seen in resting state dynamics under the genetic modifications. We further predict that SOM interneurons play dominant role in the changes of activity-state structure seen in mutant mice, and in the restaration of activity to basal levels recorded in α 5SNP mice under nicotine application.

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¹Group for Neural Theory, Laboratoire de Neurosciences Cognitives, INSERM Unité 969, Département d'Études Cognitive, École Normale Supérieure, Paris, France

²Institut Pasteur, Neurobiologie intégrative des systèmes cholinergiques, Paris, France

³CNRS UMR 3571, Paris, France

⁴Institut Pasteur, Dynamic Neuronal Imaging, Paris, France

⁵Centre for Cognition and Decision Making, National Research University Higher School of Economics, Moscow, Russia

O12 The minimalistic mathematical model of the cerebral blood flow effects during cortical spreading depression

Andrey Verisokin¹*, Darya V. Verveyko¹, and Dmitry Postnov²

Cortical spreading depression (CSD) is one of the most common abnormalities in biophysical brain functioning. We have proposed a minimalistic model that reproduces the main dynamical features of cortical spreading depression dynamics and takes into account CSD and cerebral blood flow (CBF) coupling. Despite the fact that there are many mathematical models describing the CSD, most of them do not take into consideration the role of redistribution of CBF. In contrast to previous modelling attempt [1] which was chosen as the template, we focus on the role of CBF redistribution during the formation and propagation of wave front. The flowchart of the developed model is shown in Figure 1. The model includes six dynamical variables: activator v and inhibitor w, extracellular potassium z, blood vessel radius r and upstream blood pressure p, and available neuron energy u (see Figure 1).

The main model features: (1) we have modified and extended the components of basic model [1] that stand for the energy balance; (2) the proposed model counts the relation between the extracellular potassium concentration and the radius of the nearby located blood vessel: we take into account the effect of spatial coupling (functional hyperemia) by means of weighted summation of vasodilatory driving force over some distance from neuron; (3) we propose a lumped description for hemodynamic spatial coupling, being the direct result of blood flow redistribution between different areas fed from the single upstream arterial vessel.

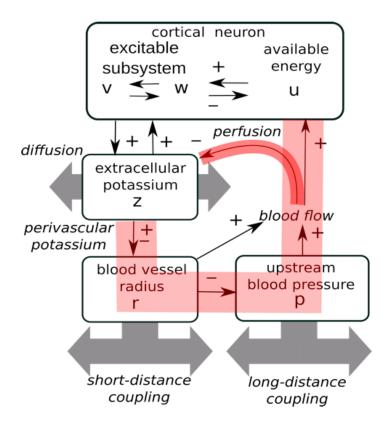


Figure 1: The schematic representation of flowchart of the developed model.

Based on the results of the numerical simulation we can conclude that the proposed model: (1) shows qualitatively reasonable results comparing with the experimental data: the uncorrelated noise-induced firing at rest; the persistent neuronal depolarization during the active phase of CSD; the depressed state afterwards, when model medium temporary losses the excitability and does not response on noisy stimuli; (2) reproduces main spatial patterns known for cortical spreading depression, migraine waves and spreading depolarization events observed in stroke and brain injuries; (3) predicts the formation of stationary dissipative Turing-like structures, formed due

¹Department of Theoretical Physics, Kursk State University, Kursk, Russian Federation

²Department of Physics, Saratov State National Research University, Saratov, Russian Federation

to the substantially different type of spatial relation – tissue perfusion. The role of perfusion in the formation of the structures was elucidated.

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O13 Necessity for coherence in motor control

Willy Wong^{1,2*}, Omid Talakoub³, Robert Chen^{4,5}, and Milos Popovic^{2,6}

The basic premise of this study is that the coherence of neural activity is required for the coordination of motor control. Motor control involves a number of brain centres, most notably the cortex, the basal ganglia, thalamus and cerebellum. How are messages coordinated between the different centres during complex movements is an open question. Following [1], we examined the hypothesis that different neural sub-populations follow a communication via coherence hypothesis. That is, for two neural sub-populations to communicate their activity must be coherent and therefore exhibit some form of mathematical synchronization.

As detailed in [2], we performed deep-brain measurements on patients undergoing treatment for Parkinson's disease (n=6) and dystonia (n=7). During a brief period after implantation of the electrodes, we were able to record the activity from either the sub-thalamic nucleus (STN) or the globus pallidus interna (GPi). These recordings constitute local field potential (LFP) recordings. Patients were asked to perform one cycle of wrist movement lasting approximately one second in duration. The movements were executed either as externally cued or through self-initiation. Simultaneous to local field recordings measured at either the STN or GPi, electroencephalographic signals (EEG) were recorded over the motor cortex. For LFP, recordings were processed by subtracting the activity from adjacent electrodes. EEG was recorded in a bipolar montage (either C3-Cz or C4-Cz). We believe that the activity we record is local in origin and not due to volume conduction, or due to the use of a common reference. Our results show that during movement, and only during movement, is there significant coupling between changes in the power of the activity with changes in coherence between the basal ganglia and the motor cortex. The changes can happen such that for beta band activity (20-30 Hz) both power/coherence is high pre and post movement, but low during course of movement. For gamma activity (30+ Hz), we observe the opposite: only during movement do we observe a coupling of increased power with increased levels of coherence either between GPi-cortex or STN-cortex. The coupling of power with coherence is not artifactual.

To better understanding the origins of these findings, we need to develop suitable mathematical models of coupled neural ensembles. We have been extending the Kuramoto model of coupled oscillators for application to this problem. Two distinct neural ensembles (in the basal ganglia and in the cortex) have neurons that are each interconnected. Moreover, the two ensembles are further connected to each other through additional links. What we can show is that an increase in power in either ensembles will lead to increased amplitude/phase coherence between the two ensembles just as found experimentally. This thus provides a first model of motor coordination between cortex and basal ganglia. Establishing the necessity for coherence in motor coordination suggests new strategies for neuromodulation similar to how functional electrical stimulation works to restore peripheral motor function.

Acknowledgements

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¹ Dept. of Electrical and Computer Engineering, University of Toronto, Toronto, ON, Canada

²Institute of Biomaterials and Biomedical Engineering, University of Toronto, Toronto, ON, Canada

³Dept. of Biology, York University, Toronto, ON, Canada

⁴Krembil Research Institute - University Health Network, Canada

⁵Division of Neurology, Faculty of Medicine, University of Toronto, Canada

⁶Toronto Rehabilitation Institute - University Health Network, Canada

O14 Dissecting gamma phase and amplitude-specific information routing in V4 of macaque during selective attention

Dmitriy Lisitsyn¹*, Eric Drebitz², Iris Grothe³, Sunita Mandon², Andreas Kreiter², and Udo A Ernst¹

Communication through coherence (CTC) postulates that stimulus information transmission is enhanced between oscillating neural populations in a favorable phase relationship, and suppressed otherwise. For example, in the case of visual cortical gamma-band synchronization during selective attention, V1 spikes arriving to V4 during its excitability peaks should be much more likely to elicit further spikes, resulting in effective signal gating; V1 spikes arriving during excitability troughs should fail or at least be less effective in evoking further activity. Further, it has been observed that average gamma power increases with attention, however, this increase appears to occur in bursts, rather than a constant oscillation. If the CTC hypothesis holds, one should expect descriptive gamma phase and amplitude dependent modulations in stimulus information routing in V4.

To explore this idea, we analyzed neural data from a previous study [1], recorded from V4 superficial layers in macaques performing a visual spatial attention task. The task required the animals to attend one of two dynamic stimuli over an extended time period. Crucially, each stimulus was superimposed with its own fluctuating luminance signature, irrelevant to the behavioral task. This allowed us to quantify the information content I of each stimuli conveyed by the physiological signal, by computing spectral coherence between each stimuli's luminance signal and V4 activity. To assess modulation effects at multiple population scales, we analyzed both LFP and spiking activity. Using gamma-band activity extracted from LFP, we dissected both LFP and spiking neural activity into phase/amplitude-specific components. We then computed the information contribution of each stimulus to these components, giving us the opportunity to assess phase/amplitude signal gating effects.

The results show that information routing is modulated by the gamma phase for both LFP and spiking activity. In LFP, we found the information routing at excitability peaks I_{peak} is significantly higher than at excitability troughs I_{trough} for both attended and non-attended stimuli (Figure 1A). We did not see this effect for spikes, which still show significant gamma phase dependence but without a preference for a specific phase across recording sessions. Comparing the stimuli content during high gamma activity $I_{high\gamma}$ against low gamma activity $I_{low\gamma}$, we found that the spiking activity exhibits significant gating increase for the attended stimulus and decrease for the non-attended stimulus (Figure 1B), however, we do not find this effect in the LFP. In summary, our study confirms basic predictions on the nature of selective information processing, namely its modulation in dependence on phase and amplitude of LFP gamma activity. Surprisingly, consistent phase modulation was only found in LFPs, while consistent amplitude modulation was only seen in spiking activity, indicating that the mechanisms implementing CTC are not yet fully understood. In particular, our results strongly motivate a refinement of current CTC models, requiring an approach encompassing different levels of complexity capable of reproducing local spiking and global population activity from different laminar sources.

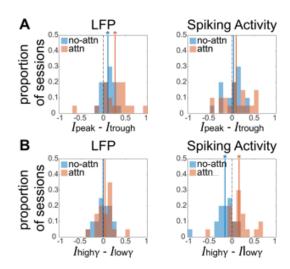


Figure 1: A. Peak vs trough γ phase info routing. B. High vs low γ amplitude info routing.

¹Computational Neuroscience Lab, Institute for Theoretical Physics, Bremen University, Germany

²Institute for Theoretical Neurobiology, Brain Research Institute, Bremen University, Germany

³Ernst Strüngmann Institute (ESI) for Neuroscience, Frankfurt, Germany

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O15 Structure-Function Relationships via Neural Field Theory

Peter A Robinson^{1,2*}, Xuelong Zhao^{1,2}, Kevin Aquino^{1,2,3}, John Griffiths^{1,2,4}, Grishma Mehta-Pandejee^{1,2}, Natasha Gabay^{1,2}, James Maclaurin^{1,2}, and Somwrita Sarkar^{1,2,5}

Patterns of brain activity are observed to be highly conserved across states of arousal, and between task and non-task conditions. This strongly suggests that these are natural modes (eigenmodes) of the brain, which are excited in different ways under different circumstances. Neural field theory (NFT), which averages over brain microstructure, is ideally suited to deriving brain eigenmodes and interpreting them in terms of underlying physiology. It also provides means of systematically interrelating structure and function via these eigenmodes.

Here, NFT is used to predict the eigenmodes of the continuous cortical surface, including interhemispheric connections. For comparison, eigenmodes of a discrete cortical connection matrix are calculated by standard matrix procedures. Mode energies and symmetry properties are used to constrain interhemispheric conductivities and physiological properties of the cortex. Eigenmodes are then used to derive underlying effective and functional connectivities from system transfer functions and two-point correlations of background activity, respectively.

Neural field eigenmodes are shown to occur in a hierarchy closely related to that of the eigenmodes of a sphere, with added symmetries induced by bihemispheric structure. A close correspondence is also found with the eigenmodes of an anatomical connection matrix, confirming the validity of the neural field approach. The results demonstrate that the brain is in a near-critical state, consistent with estimates from electroencephalographic spectra. It is found that each hemisphere receives near-balanced inputs, with approximately 15 percent of net inputs coming from the contralateral hemisphere, 73 percent from the ipsilateral one, and 12 percent from the environment, meaning that it is in a highly instrospective state. Most activity is predicted to be in symmetric modes, in accord with experiment.

NFT allows structure and activity to be unequivocally interrelated, including the correlations used to define functional connectivity matrices. Eigenmode decomposition of these matrices enables underlying effective connectivities to be systematically derived from functional connectivities, and vice versa, and related to resulting activity patterns. This means that relatively easily observed correlations can be used to infer both average structure and the strengths of effective connectivities that it supports in a noninvasive manner.

In summary, physiologically-based NFT thus explains and unifies multiple phenomena relating to structure, function, and activity via eigenmodes. This allows analysis of activity and structure in terms of the natural dynamic modes of the system, rather than ones that are defined via statistical signal analyses that do not incorporate physiology.

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¹School of Physics, University of Sydney, Sydney, Australia

²Center for Integrative Brain Function, University of Sydney, Sydney, New South Wales, Australia

³Sir Peter Mansfield Imaging Center, University of Nottingham, Nottingham, UK

⁴Rotman Research Institute at Baycrest, Toronto, Canada

⁵Design Lab, School of Architecture, University of Sydney, Sydney, New South Wales, Australia

O16 Dynamic Operations of Hierarchically Interacting Canonical Microcircuits

Tim Kunze^{1,2}*, Jens Haueisen², and Thomas R. Knösche¹

Research on canonical microcircuits conceptualizes findings of the recursive occurrence of neural populations and coupling patterns in vertically and horizontally structured divisions (i.e. cortical columns) of the cerebral cortex [1]. The profound description and examination of the link between canonical architectures and the associated functionality promises a better understanding of higher level functions which emerge from the interaction of canonical microcircuits. Fundamental for this interaction is the embedding canonical microcircuits in hierarchical networks [2], mediating both bottom-up and top-down signals to specific neuronal populations. Here, computational studies can help to formulate hypotheses about constitutive mechanisms, which are experimentally identifiable in the neural substrate.

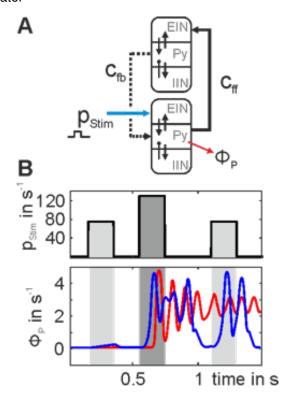


Figure 1: A. Adaptive dynamical shift of a perceptual threshold in a hierarchical configuration of two interacting neural mass models, mimicking canonical microcircuits. B. A bottom-up target input (light grey) excites the lower area only after previous application of a priming stimulus (dark grey).

We use a neural mass model [3], where a pyramidal cell population (Py) receives negative feedback from an inhibitory interneuron population (IIN) and positive feedback via a secondary excitatory population of interneurons (EIN), representing neurons in layer IV. We systematically apply transient afferent inputs, modeled by pulses of various magnitude and duration, as bottom-up signals to the EIN or as top-down signals to the Py [2] and monitor the behavior of the Py. These response behaviors are classified as: a) nonresponsive for sub-threshold transient deflections, b) transfer for supra-threshold transient deflections, and c) memory for sustained supra-threshold deflections and are mapped to the stimulation parameter range.

Single-channel stimulations, either bottom-up (to EIN) or top-down (to Py), lead to differential response behaviors, where strong and long bottom-up stimulations are preferably stored (memory behavior), in contrast to top-down signals, which predominantly show transient deflections. In a concomitant stimulation, constant top-down input modulates the modelÕs sensitivity to pulsed bottom-up stimulation in favor of the memory response behavior. We employ this modulatory influence in a hierarchical network (Fig. 1A) comprising two canonical microcircuits to show a conceivable neural mechanism for the dynamic adaptation of a perceptual threshold. In this configuration, a target stimulus is not able to excite a perceptual area, unless a priming stimulus tunes the networkÕs sensitivity.

¹Max Planck Institute for Human Cognitive and Brain Sciences, Leipzig, Germany

²Institute of Biomedical Engineering and Informatics, Ilmenau University of Technology, Ilmenau, Germany

The differential response behaviors to top-down and bottom-up stimuli indicate the functional role of separate input channels in canonical microcircuits. Exemplarily, we show one constitutive operation emerging from interacting microcircuits, but expect many more mechanisms relevant in cognitive disciplines like language or memory, such as stimulus selection or structure building computations. Further, the present results in the hierarchical setup demand a further evaluation in light of predictive coding where important findings of neural communication have been put forward.

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O17 Learning structure of 3D objects with cortical columns

Subutai Ahmad*, Yuwei Cui, Marcus Lewis, and Jeff Hawkins Numenta, Redwood City, CA, USA

The neocortex is organized in cellular layers. Connections between layers run mostly perpendicular to the surface of the neocortex, which suggests a columnar pattern of activation across layers. The cells in some layers also send their axons across long distances parallel to the surface of the neocortex, which suggests a laminar pattern of activation across multiple columns. The vertical and horizontal spread of axons is a ubiquitous feature of all neocortical regions.

In this study, we propose a network model that utilizes both intra-column and cross-column connections for robust object learning and recognition (Figure 1). The model consists of a set of cortical columns, where each cortical column processes a different subset of the sensory input space. An object consists of a set of component features at particular locations on the object. Each cortical column learns an object by forming feedforward connections from its component features to a set of active neurons in a different cellular layer. After learning, sensation of a sequence of object features leads to activations of the corresponding neural population representing the object. Since features can be shared among multiple objects, information received by a single cortical column is often ambiguous. The model uses auto-associative connections to integrate many sensations over time and can converge onto unique object representations once sufficient feature are sampled. The recognition speed and accuracy can be improved by simultaneously considering multiple cortical columns with lateral connections, where each column learns feedforward connections independently and learns cross-column lateral connections according to Hebbian rules. The lateral inputs target distal dendritic segments. Although they are not strong enough to directly activate a neuron, neurons with both lateral input and feedforward input will fire earlier and prevent other neurons from responding [1]. The cross-columnar connections bias each column to form a representation that is consistent with the partial knowledge of all the interconnected columns. We show that objects can be recognized

faster and that each cortical column can store more objects by using cross-column connections. The model is consistent with a large body of anatomical and physiological evidence and provides a number of predictions that

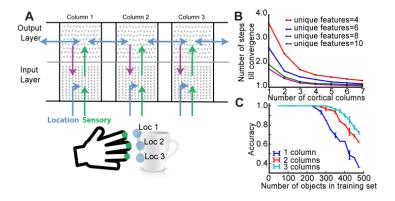


Figure 1: A. We consider the problem of object recognition with a set of cortical columns. Each column receives sensory input from a different sensor (e.g., different finger). A first layer of the network transforms the raw sensory input into sparse distributed representations that corresponds to object features. The second layer receives feedforward inputs from the first layer. It recognizes an object by converging onto a stable activation pattern through lateral connections. B. The recognition speed increases as a function of column number. C. Retrieval accuracy of object during testing vs. the number of learned objects. More objects can be learned with networks with more cortical columns.

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can be tested in future experiments.

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O18 Influence of network topology on spreading of epileptic seizure

Simona Olmi^{1*}, Spase Petkoski², Fabrice Bartolomei³, Maxime Guye⁴, and Viktor Jirsa²

In partial epilepsy, seizures originate in a local network, the so-called epileptogenic zone, before recruiting other close or distant brain regions. Correctly delineating the epileptogenic and the propagation zone is essential for successful resective surgery. In particular, the stereotaxic EEG (SEEG) is used to edge the resection zone. Nevertheless, the propagation pathways of epileptic seizures are still largely unknown. We utilize a specific dynamical model for epilepsy, the Epileptor model [1], to predict the recruitment network given the seizure origins and the structural brain connectivity. Thus, we try to understand the role played by the topology in constraining the recruitment process and we suggest a paradigm for epileptic surgery that relies on minimal invasiveness and maximum effectiveness. In particular, we schematize the brain network dynamics in terms of neural mass models able to captures the details of the autonomous slow evolution of interictal and ictal phases; these mass models are coupled among them and the coupling terms model the effective presence of nerve pathways and fibers among different brain regions [2]. In this framework it is possible to identify the minimal number of local disconnections of the epileptogenic zone that are necessary to stop seizure propagation via the application of linear stability analysis and, therefore, to define the optimal set of links to be cut in order to stop seizure propagation (see Figure 1). In order to demonstrate the potential use of this framework in practice, we apply our methods to structural connectivity matrices derived from patients affected by partial epilepsy. In all cases a partial disconnection, that counts for the resection of few pathways, is sufficient to stop seizure activity in the brain. Therefore we demonstrate that seizure spreading is thus supported and enhanced by the underlying topology and that a disconnection procedure, if well addressed, can become a fruitful procedure to improve the success rate of epilepsy surgery.

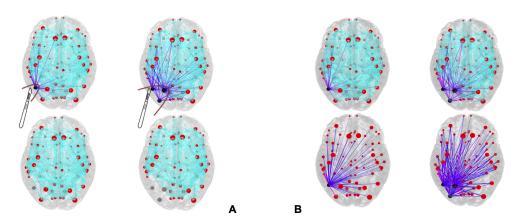


Figure 1: A: Standard resection technique, where the entire epileptogenic zone (EZ) is removed during surgical operation. Blue links represent the outgoing connections of the EZ and are completely removed during the current surgical procedures. B: Lesioning depicts the minimal number of links that are sufficient to be removed (magenta) in order to stop the seizure, versus the total number of outgoing links from the EZs (blue) that are removed during the resection of an entire EZ. Cyan links represent in both panels the full connectivity of the network.

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¹Weierstrass Institute, Berlin, Germany

²Aix Marseille Université, Inserm, Institut de Neurosciences des Systèmes, Marseille, France

³Assistance Publique Hôpitaux de Marseille, Hôpital de la Timone, Service de Neurophysiologie Clinique, CHU, Marseille, France

⁴Faculté de Médecine de la Timone, centre de Résonance Magnétique et Biologique et Médicale, Medical School of Marseille, Aix-Marseille Université, Marseille, France

O19 A model-based approach for detecting multiple change points in multivariate spike count data

Hazem Toutounji*, Daniel Durstewitz

Department of Theoretical Neuroscience, Bernstein Center for Computational Neuroscience, Central Institute of Mental Health, Medical Faculty Mannheim, Heidelberg University, Germany

Neural data often consist of multiple single unit recordings in the form of spike count time series. These time series are often highly nonstationary, where statistical moments, such as firing rates, vary to potentially encode features of the experimental paradigm, like changes in external input or different task phases. Changes in the firing rates may be sudden or gradual, and their time scale and onset may reflect information regarding neural computations, such as learning [1] or the accumulation of sensory evidence [2].

Here we develop an approach for detecting and parametrising multiple changes in multivariate spike count data within the statistical framework of State Space Models (SSM) [3]. The model assumes a nonlinear, nonstationary, autoregressive Gaussian process that captures the underlying latent neural dynamics. However, given their discrete, nonnegative nature, assumptions of normality are not guaranteed to produce consistent estimates of spike count statistical moments. Instead, the Gaussian process generates spike counts by a Poisson observation function. Both latent trajectories in phase space and latent model parameters, in addition to observation model parameters, are estimated by a 3-stage Expectation-Maximisation (EM) procedure [4]. The latter relies on Newton's method [5] to maximise, under constraints, a global Laplace approximation [6] of spike-count data's log-likelihood, given the SSM and its parameters. The dimensionality of the latent model equals the number of unknown nonstationary events, termed change points, and is selected by a cross-validation procedure. Observations, on the other hand, are generally of a much higher dimension than the latent dynamics. Due to this substantial dimensionality reduction [7], latent trajectories, thus, offer a parsimonious representation of the most relevant features in neural dynamics.

The estimation procedure is first tested on simulated data, to assure that the latent states and model parameters are correctly identified in comparison to the ground truth. As a real data example, the model is fitted to multiple single unit recordings from rat medial prefrontal cortex neurons during an operant rule switching task. The resulting reconstruction of the underlying dynamics will allow matching the neural correlates of learning to their behavioral counterpart, by relating behavioral changes to population-wide change points, as estimated by the model.

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O20 Geppetto: an open source visualisation and simulation platform for neuroscience

Matteo Cantarelli^{1*}, Adrian Quintana^{2,4}, Boris Marin⁴, Matt Earnshaw⁴, Padraig Gleeson⁴, Robert Court³, Robert A McDougal⁶, R Angus Silver⁴, Salvador Dura-Bernal⁵, Stephen Larson¹, William W Lytton⁵, and Giovanni Idili¹

Geppetto (geppetto.org) is an open-source web-based platform to explore and simulate neuroscience data and models. The platform, originally designed to support the simulation of a cell-level model of C. elegans as part of the OpenWorm project [1], has grown into a generic framework suitable for various neuroscience applications, offering out of the box solutions for data visualisation, integration and simulation. Geppetto is today used by Open Source Brain (opensourcebrain.org) (Figure 1.A), to explore and simulate computational neuroscience models described in NeuroML version 2 with a variety of simulators and by the Virtual Fly Brain (virtualflybrain.org) (Figure 1.B) to explore and visualise anatomy (including neuropil, segmented neurons and gene expression pattern data) and ontology knowledge base of *Drosophila melanogaster*. Geppetto is also being used to build a new experimental UI for the NEURON simulation environment [2][3] (Figure 1.C) based on Python and Jupyter. WormSim (wormsim.org) (Figure 1.) embeds Geppetto to let users explore dynamic mechanical and electrophysiological models of *C. elegans* produced by the OpenWorm project. Geppetto is capable of reading and visualising experimental data in the NWB format (nwb.org) to allow experimental and computational neuroscientists to share and compare data and models using a common platform. Geppetto is freely available, well documented and has an active user community. Interested potential users can try out the latest version of the platform at live.geppetto.org.

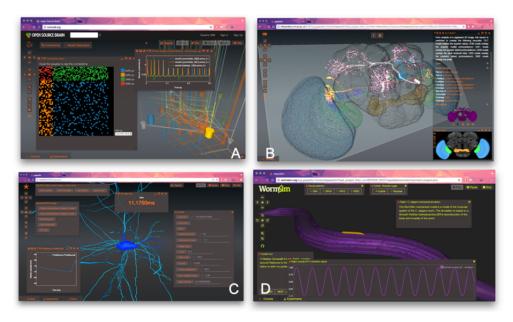


Figure 1: Gepetto in different deployments.

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¹OpenWorm Foundation, Delaware, USA

²EyeSeeTea Ltd., London, UK

³Edinburgh University, Edinburgh, UK

⁴Department of Neuroscience, Physiology and Pharmacology, University College London, London, UK

⁵State University of New York Downstate Medical Center, Brooklyn, NY, USA

⁶ Yale University, New Haven, CT, USA

O21 Position is coherently represented during flickering instabilities of place-cell cognitive maps in the hippocampus

Lorenzo Posani^{1*}, Simona Cocco¹, Karel Ježek², and Rémi Monasson¹

Place cells in hippocampus exhibit sharp spatially-related firing fields, which are formed when the animal explores new environments and are retrieved, as memories, each time the rat is placed back in those specific settings. Knowledge of the environment-specific set of place fields (map) allows for the application of Bayesian statistics to infer the position of the rodent from neuronal activity (Figure 1B, 1C). Likewise, functional-connectivity models, based only on neural correlations, i.e. with no knowledge of place fields or position, can identify the expressed map as a function of time (Figure 1A) [1]. We apply both these inference procedures to CA3 recordings from a recent "teleportation" experiment [2], in which instantaneous switches between the identity of two familiar environments trigger the instability of the recalled memory state, which flickers back and forth between the two corresponding maps (Figure 1A, 1E). Our analysis shows that the rat position is not accurately inferred during the unstable periods, under conventional approach relying on brain processing only the external input information (i.e. environment cues, Figure 1B, 1D: red curve). However, if the position is inferred using the template reflecting the decoded inner state of the network, the position error is significantly reduced, reaching values comparable to the stable conditions (Figure 1C, 1D: blue curve). Results suggest that position is robustly encoded in CA3, even during periods of conflict or ambiguity in the input information resulting in global map changes on fast dynamical time scales.

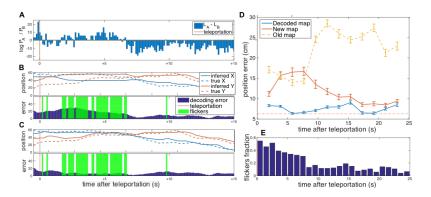


Figure 1: Position inference during flickering instabilities. A. decoded environment (log-likelihood difference) as a function of time after a teleportation in CA3. Note the flickering dynamics in the 0-5 sec interval. B. Inferred vs. real positions of the animal; place fields corresponding to light conditions were used for the inference. Freely-moving rat in a 60x60 cm box. C. same as B with position inferred using the place fields associated to the decoded map (sign of ΔL in panel A). D. positional errors averaged over 15 teleportation events; dashed line indicates the level of error for stable conditions (no light switches). E. fraction of flickering time bins (ΔL -decoded map differs from light conditions) as a function of time after the light switch.

Acknowledgements

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References

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¹Laboratories of Statistical & Theoretical Physics, Ecole Normale Supérieure, Paris, France

²Laboratory of Experimental Neurophysiology, Biomedical Center, Charles University, Prague, Czech Republic

Workshops

W1 Neuronal Oscillations: Mechanisms and Functionality

Room K.202, Wed and Thur 9:00 to 18:00

Horacio G Rotstein, New Jersey Institute of Technology

Frances Skinner, Krembil Research Institute

Vassilis Cutsuridis, University of Lincoln

Oscillations at various frequency ranges have been observed in several brain structures (hippocampus, entorhinal cortex, olfactory bulb and others). They are believed to be important for cognitive functions such as learning, memory, navigation and attention. These rhythms have been studied at the single cell level, as the result of the interaction of a neuron's intrinsic properties, at the network level, as the result of the interaction between the participating neurons and neuronal populations in a given brain region, and at higher levels of organization involving several of these regions. Recent advances in this field have benefited from the interaction between experiment and theory, and models with varying levels of detail.

The purpose of this workshop is to bring together modelers, experimentalists and theorists with the goal of sharing and discussing their current results and ideas on the underlying mechanisms that govern the generation of these rhythms at various levels of organization, and their functional implications.

An additional goal is to discuss what we mean by an explanation in the context of this workshop. To this end, speakers will be encouraged to address this issue from the perspective of Aristotle's four causes (efficient, material, formal and final causes). Please see 'What is computational neuroscience? (XVI) What is an explanation?'.

- Frances Skinner (Krembil Research Institute, Canada) "Theta oscillations in the hippocampus: generation mechanisms"
- John White (Boston University, USA) "Synaptic contributions to the theta rhythm in the isolated hippocampal formation"
- Marianne Bezaire (Boston University, U.S.A.) "Spontaneous theta oscillations arise in a detailed, large-scale model of the rodent hippocampal CA1 subfield"
- Vassilis Cutsuridis (University of Lincoln, U.K.) "Memory formation and replay in the hippocampal CA1 microcircuit"
- Jose Guzman (IST, Austria) "The synaptic microcircuit of pattern completion in the hippocampal CA3 network"
- Leonid Rubchinsky (Indiana University Purdue University Indianapolis, USA) "Temporal patterns on neural synchrony: observations, mechanisms, and functions"
- Dan Levenstein (NYU, USA) "Synchronized neocortical dynamics of the NREM slow oscillation: from mechanism to function"
- Francesco Battaglia (Donders Centre for Neuroscience, Radboud Universiteit Nijmegen, The Netherlands) "Oscillatory interaction between hippocampus and prefrontal cortex during memory consolidation"
- · Marlene Bartos (University of Freiburg, Germany) TBA
- Christoph Borgers (Tufts University, USA) "Some thoughts on the effects of synchrony"
- Carmen Canavier (LSU Health Sciences Center, USA) "Different gamma mechanisms coexist in the same excitatory/inhibitory network architecture"
- Mark Cunningham (University of Newcastle, U.K.) "Mechanisms underlying the generation of human neocortical gamma frequency oscillations"
- · Jeremie Lefebvre (University of Toronto, Canada) TBA
- Adrien Peyrache (McGill University, Canada) "Oscillatory dynamics in the limbic thalamo-cortical network reveal subcortical information flow to and from the hippocampus"
- Horacio Rotstein (New Jersey Institute of Technology, USA) "A conceptual framework for the study of resonance in neuronal systems"

- Ausra Saudargiene (Neuroscience Institute of the Lithuanian University of Health Sciences & Vytautas Magnus University, Kaunas, Lithuania) TBA
- Bijan Pesaran (New York University, USA) "Neural coherence supports functional inhibition during movement coordination"

W2 Methods of Information Theory in Computational Neuroscience

Room C.101. Wed and Thur 9:00 to 18:00

Joseph T. Lizier, University of Sydney

Viola Priesemann, Max Planck Institute for Dynamics and Self-organisation

Justin Dauwels, Nanyang Technological University

Taro Toyoizumi, RIKEN Brain Science Institute

Alexander G Dimitrov, Washington State University

Lubomir Kostal, Academy of Sciences of the Czech Republic

Michael Wibral, Goethe University, Frankfurt

Methods originally developed in Information Theory have found wide applicability in computational neuroscience. Beyond these original methods there is a need to develop novel tools and approaches that are driven by problems arising in neuroscience. A number of researchers in computational/systems neuroscience and in information/communication theory are investigating problems of information representation and processing. While the goals are often the same, these researchers bring different perspectives and points of view to a common set of neuroscience problems. Often they participate in different fora and their interaction is limited.

The goal of the workshop is to bring some of these researchers together to discuss challenges posed by neuroscience and to exchange ideas and present their latest work. The workshop is targeted towards computational and systems neuroscientists with interest in methods of information theory as well as information/communication theorists with interest in neuroscience.

For an up to date list of talks and schedule please see http://bit.ly/cns2017itw

- Selin Aviyente (Michigan State University) "Directed information: application to EEG during cognitive control"
- Lionel Barnett (University of Sussex) "Information transfer in continuous and discrete time"
- Karl Friston (University College London) "Active inference and artificial curiosity"
- Renaud Jolivet (University of Geneva) "Energy-efficient information transfer at synapses"
- Lubomir Kostal (Academy of Sciences of the Czech Republic) "Reference frame independence as a constraint on the mutual information decomposition"
- Joseph T. Lizier (The University of Sydney) "An estimator for transfer entropy between spike trains"
- Daniele Marinazzo (University of Ghent) "Synergetic and redundant information flow detected by unnormalized Granger causality: application to resting state fMRI"
- Jil Meier (Delft University of Technology) "The epidemic spreading model and the direction of information flow in brain networks"
- · Viola Priesemann (Max Planck Institute for Dynamics and Self-organization, Goettingen) TBA
- Adrià Tauste (Universitat Pompeu Fabra) "Directed information flow within the thalamo-cortical network"
- Tatjana Tchumatchenko (Max Planck Institute for Brain Research, Frankfurt) "Information coding of mean and variance modulating signals in cortical neurons"
- Taro Toyoizumi (RIKEN Brain Science Institute) "A local learning rule for independent component analysis"
- Raul Vicente (University of Tartu) TBA
- · Plus additional contributed talks ...

W3 Recent Methods and Analyses for Large-scale Neuronal Population Recordings

Room C.102. Wed and Thur 9:00 to 18:00

Michela Chiappalone, Istituto Italiano di Tecnologia, Genova Valentina Pasquale, Istituto Italiano di Tecnologia, Genova Pierre Yger, Institut de la Vision, INSERM, Paris

Understanding how assemblies of neurons encode information requires recording large populations of cells in the brain. In recent years, progress in population calcium imaging and multichannel electrophysiology opened the possibility to record from hundreds or even thousands of neurons simultaneously. While these techniques offer unprecedented chances to monitor large neural circuits, they also push for the design of new algorithms to gather and process information from such high-dimensional datasets.

This two-day workshop will gather leading experimentalists and theoreticians to discuss latest computational methods and analyses used to process such large-scale neuronal population recordings, both <u>in vivo</u> and <u>in vitro</u>. Focusing on high-density electrophysiology and calcium imaging, it will review recent advances in neuroinformatics research, including spike sorting techniques and characterization of neural assemblies' spatio-temporal activity. It will be a unique opportunity to address various questions such as:

- How to enhance the robustness of new algorithms identifying spikes, and/or design a proper validation framework ensuring the quality of the data?
- How to detect neural assemblies' activity and correlations both in space and time, and possibly link them
 to sensory perception and behavior?
- What are the links, from a signal processing point of view, between calcium imaging and high density electrophysiology recordings?

- Matthias Hennig (University of Edinburgh, Scotland) "Spike sorting for large scale multielectrode arrays: efficient methods and lessons learnt"
- Felix Franke (ETH Zurich, Switzerland) "How prewhitening can improve spike sorting performance"
- Nick Steinmetz (University College London, UK) "Recording large, distributed neuronal populations with Neuropixels electrode arrays in behaving mice"
- Pierre Yger or Olivier Marre (Institut de la Vision, France) "Towards online accurate spike sorting for thousands of channels"
- Thomas Deneux (UNIC, CNRS, France) "Spike inference from calcium signals: MLspike algorithm and general perspectives"
- Marius Pachitariu (University College London, UK) "Kilosort and Suite2p: robust and scalable frameworks for neural activity extraction in large-scale recordings"
- Stephen J. Eglen (University of Cambridge, UK) "Detecting pairwise correlations in high-density recordings: open science in action"
- Adrien Peyrache (McGill University, Canada) "Millisecond synchrony in the thalamo-cortical network of the brain's navigation system: a mechanism for efficient information transmission?"
- George Dimitriadis (Sainsbury Welcome Centre, UK) "Understanding large-scale neural recordings: Ground truth data sets and the T-sne visualizations tool"
- Gaute Einevoll (NMBU, Norway) "Biophysical modeling of benchmarking data for validation of methods for analysing electrophysiological data"
- Yannick Bornat (IMS Bordeaux, France) "Low latency hardware computing to use electrode array inputs in closed loop experiments"
- Ulisse Ferrari (Institut de la Vision, France) "Closed-loop estimation of retinal network sensitivity reveals signature of efficient coding"
- Valentina Pasquale (Istituto Italiano di Tecnologia, Italy) "Measuring similarity of endogenous and evoked activity patterns in cultured cortical networks"

- Sonja Grün (Forschungszentrum Jülich, Germany) "Analysis of massively parallel spike data for higherorder correlations"
- Arno Onken (Istituto Italiano di Tecnologia, Italy) "Matrix and tensor factorizations for analyzing neural population activity"
- Paolo Bonifazi (Ikerbasque, Bilbao, Spain) "Sparse synchronizations and neuronal network failures: astrocytes replacement recovers global synchronizations in Atm-deficient cerebellar circuits in vitro"

W4 New Advances in Theoretical Tools for the Study of Large-scale Neural Systems

Room K.101, Wed 9:00 to 18:00 and Thur 9:00 to 12:30

Simona Olmi, Institute of Complex Systems- CNR

David Angulo-Garcia, Aix-Marseille University

Benjamin Lindner, Humboldt University Berlin

New advances in mathematical and computational neurosciences require the development of mathematical tools able to explain the dynamical evolution observed at different spatiotemporal scales. Our idea is to focus on the specific issue of meso/macroscopic dynamical behaviours in neural systems, given their relevance for population coding and computation. This workshop aims at discussing some of the most recent analytical techniques and mathematical tools used to derive the evolution of neural populations and its emergent dynamics, with the hope of understanding how features of meso/macroscopic systems such as local dynamics, heterogeneity, noise and topology shape their response. With this workshop we expect to bridge the gap between models and data and to further shed light on the inter-relation between structure, dynamics and function of the brain circuitry by studying the transitions between different collective states.

- Olivier Faugueras (INRIA and LJAD, France) "Coping with correlations in the analysis of the thermodynamic limit of neuronal networks"
- Ernest Montbrió (Universitat Pompeu Fabra, Spain) "Firing rate equations require a spike synchrony mechanism to correctly describe fast oscillations in inhibitory networks"
- Alex Roxin (Campus de Bellaterra, Barcelona, Spain) "Microscopic and macroscopic states in networks of recurrently coupled spiking neurons"
- Srdjan Ostojic (Ecole Normale Supérieure, Paris, France) "From dynamics to computations in recurrent random networks with low-rank connectivity structure"
- Maurizio Mattia (Istituto Superiore di Sanità, Italy) "Low-dimensional dynamics of cortical networks explaining slow-wave activity and its transition to the asynchronous state."
- Tilo Schwalger (Ecole polytechnique fédérale de Lausanne, Switzerland) "Mesoscopic dynamics of interacting neural populations of spiking neurons"
- Boris Gutkin (Ecole Normale Supérieure, Paris, France) "Phase response curves for emergent network oscilations"
- Tatyana Sharpee (Salk Institute for Biological Studies, CA) "Dynamics of large-scale recurrent networks"
- Raffaella Burioni (University of Parma, Italy) "Chaos and correlated avalanches in excitatory neural networks with synaptic plasticity"
- Magnus Richardson (University of Warwick, United Kingdom) "Negative-feedback control of cortical activity by the neuromodulator adenosine"
- Marc de Kamps (University of Leeds, United Kingdom) TBA
- Davide Bernardi (Bernstein Center for Computational Neuroscience Berlin, Germany) "Optimal detection of single-cell stimulation in large random networks of integrate-and-fire neurons"
- David Angulo-Garcia (University of Aix-Marseille, France) "Lazarus effect: mechanism of neuron's deactivation and reactivation in sparse heterogeneous inhibitory neuronal networks"
- Simona Olmi (Weierstrass Institute Berlin, Germany) "Exact firing time statistics of neurons driven by discrete inhibitory noise"

W5 Theoretical Neuroscience in the Human Brain Project

Room K.201, Wed 9:00 to 18:00

Michele Giugliano, Universiteit Antwerpen

Alain Destexhe, Centre National de la Recherche Scientifique (CNRS)

Viktor Jirsa, Aix-Marseille University

This workshop gathers in an open and interactive forum, some of the Principal Investigators of the Theoretical Neuroscience pillar (SP4) of the Human Brain Project. SP4 provides novel mathematical descriptions for multiscale representations of the brain and its components, from cellular to network levels. We propose to focus this workshop on how microscopic processes in the brain express themselves parametrically on a higher level of organisation, ranging from populations of neurons, entire brain areas to the whole-brain networks. Individual talks will address and discuss the following research questions:

- How can mean-field models achieve a direct faithful integration between mesoscopic and macroscopic signals (e.g. LFP, EEG, fMRI)?
- How can building blocks, including different signals of the brain, be incorporated into simplified descriptions
 of dendrites, generic models, and synaptic plasticity?
- · What is the state-of-the art of large-scale models of the whole brain?
- Which do recent advances in modelling synaptic plasticity and memory tell about computational properties of the neocortex?
- · How close is our community to the modelling of the whole-brain?

- Alain Destexhe (UNIC, CNRS, Gif sur Yvette, and European Institute for Theoretical Neuroscience, Paris)
 "Welcome note and Introduction"
- Gaute Einevoll (Norvergian University of Life Sciences, Ås, Norway) TBA
- Olivier Faugeras (Inria MathNeuro & TOSCA Teams, Sophia Antipolis, France) TBA
- Michele Giugliano (Universiteit Antwerpen (Belgium) "The dynamical response properties of cortical neurons"
- · Moritz Helias (Research Centre Juelich, Germany) "Towards a field theory for neuronal networks"
- Viktor Jirsa (Institut de Neuroscience des Systemes, Marseille, France) "Translational neuroscience: from bifurcations to personalized medicine"
- Marc de Kamps (University of Leeds, UK) TBA
- Marja-Leena Linne (Tampere University of Technology, Finland) TBA
- Sacha van Albada (Jülich Research Centre, Jülich, Germany) "Spiking dynamics and inter-area interactions in a supercomputational model of cortex"
- Nicole Voges (Jülich Research Centre, Jülich, Germany) "Breaking the complexity barrier of analysis by reproducible workflows"
- Gorka Zamora-López (Center for Brain and Cognition at Universitat Pompeu Fabra, Spain) "Relation between structural and functional brain connectivities: lessons learned and future plans within HBP"

W6 Computational and Experimental Advances in Cerebellum Research

Room K.102, Wed 9:00 to 18:00

Erik De Schutter, Okinawa Institute of Science and Technology **Yunliang Zang**, Okinawa Institute of Science and Technology

With the improvement of experimental techniques, like in vivo patch-clamp recording and calcium imaging, we have gained a deeper understanding of information processing in the cerebellum. However, experiment driven computational modeling is still an indispensable tool to explore the implications of these findings. The purpose of this workshop is to foster an active dialogue between experimentalists and modelers about cerebellar physiology and function. Sufficient time will also be reserved to discuss how modeling and experiments can provide interesting data relevant to each other.

Both experimental and computational work at the level of cerebellar single cell and network studies are included in the workshop.

- Arnd Roth (UCL, London, UK) "Untangling cerebellar circuits with scanning electron microscopy and focused ion beam milling"
- Alessandro Barri (Institut Pasteur, Paris, France) "Temporal processing in the cerebellar cortex enabled by dynamical synapses"
- · Coffee break
- Philippe Isope (CNRS, Strasbourg, France) "How presynaptic short term dynamics influence Purkinje cell discharge in the cerebellum"
- · Mario Negrello (Erasmus MC, Rotterdam, Netherland) "The origin of complex spike synchrony"
- Ian Duguid (University of Edinburgh, Edinburgh, UK) "Purkinje cell dendritic responses during self-paced locomotion"
- Yunliang Zang (OIST, Okinawa, Japan) "Voltage- and branch-dependent complex spike responses in Purkinje neurons"
- Brandon Stell (CNRS, Paris, France), "In vivo imaging of Purkinje cell simple spikes"
- Paul Chadderton (Imperial College London, London, UK) "Cerebellar processing of kinematic signals during active whisking"

W7 Principles and Applications of Extracellular Potentials

Room K.203, Wed 9:00 to 18:00

Michiel Remme, Humboldt University Berlin

Torbjoern Ness, Norwegian University of life Sciences

Gaute Einevoll, Norwegian University of life Sciences - University of Oslo

Electrical signals from the cortical surface of animals were recorded as early as 1875. The high-frequency part (above $\sim 500~\text{Hz}$) of the recorded potentials provides information about the spiking activity of neurons located around the electrode, whereas the low-frequency part, the $\hat{a} \in \text{local}$ field potential $\hat{a} \in \text{local}$ (LFP), was found more difficult to interpret. Recently, the interest in LFPs has undergone a resurgence. Key reasons are the growing capacity for streaming continuous data from multiple electrodes and the development of multicontact electrodes for high-density recordings across areas and laminae. Further, the LFP captures key integrative synaptic processes that cannot be measured by observing the spiking activity of a few neurons alone. The LFP is also a promising signal for steering neuroprosthetic devices and for monitoring neural activity in human recordings because they are more easily and stably recorded in chronic settings than are spikes. The goal of the workshop is to provide a resume of the advances in understanding the generation of extracellular potentials through theoretical and experimental approaches as well as on the consequences (e.g., ephaptic interactions) and the applications of such signals (e.g., neuroprosthetic devices).

- Ad Aertsen (University of Freiburg) "Decoding motor cortex activity at multiple scales"
- Costas Anastassiou (Allen Institute for Brain Science) "Brain dynamics and associated electric fields during physiological and pathological activity"
- Florian Aspart (Technical University of Berlin) "Frequency dependent polarization of pyramidal cells models due to weak extracellular fields"
- Gaute Einevoll (Norwegian University of Life Sciences) "What can we learn from local field potentials (LFPs)?"
- Joshua Goldwyn (Ohio State University) "Generators of field potentials and (possible) ephaptic interactions in the auditory brainstem"
- Sonja Grün (Research Centre Jülich, Aachen University) "Spatial and temporal LFP-LFP and spike-LFP relationships"
- Paula Kuokkanen (Humboldt University Berlin) "Extracellular ITD potential and spike contributions in barn owl's nucleus laminaris"
- Alberto Mazzoni (Scuola Superiore Sant'Anna) "Predicting risk attitude in conflictual economic tasks from subcortical local field potentials"
- Arno Onken (Italian Institute of Technology) "Application of mixed vine copulas to model jointly neural spikes and local field potentials"
- Bartosz Teleńczuk (Centre National de la Recherche Scientifique) "Contributions of inhibitory and excitatory neurons to the focal LFP in human and monkey"

W8 Fingerprints and Applications of Brain Dynamics Estimated from Neuroimaging Data

Room K.103, Wed 9:00 to 18:00

Matthieu Gilson, University Pompeu Fabra

Tim van Hartevelt, Oxford

The functioning of the brain relies on detailed interactions between specialized neuronal subsystems, implementing joint segregation and integration of information such as sensory stimuli, memory tokens and intentions. Nowadays, neuroimaging techniques (fMRI, EEG, MEG, etc.) provide indirect measurements of the neuronal activity at the whole-brain level. Recent efforts have focused on extracting fingerprints of the measured brain dynamics to discriminate between tasks, conditions (e.g., sleep vs. awake) or individuals. For example, given a dynamic network model, whole-brain effective connectivity describes the interaction scheme between regions for each condition, which can be quantitatively compared. The goal of this workshop is to review both data-analysis methods and model-based approaches that have attacked this problem.

Abstracts available here: matthieugilson.eu/workshop_CNS2017.html

- Henrique Fernandes (U Aarhus, Denmark) "Brain fingerprints of structural connectivity in health and disease"
- Dante Mantini (KU Leuven, Belgium) "Detecting large-scale brain networks using high-density electroen-cephalography"
- Emily Finn (NIMH, Bethesda, USA) "Can we manipulate brain state to emphasize individual differences in functional connectivity?"
- Anish Mitra (U Washington, St Louis, USA) "Structured temporal sequences in spontaneous human brain activity"
- Demian Battaglia (INSERM, Marseille, France) TBA
- Joana Cabral (Oxford U, UK) "Spontaneous switching between states of functional connectivity relates to cognitive performance in healthy older adults"
- Karl Friston (UCL, London, UK) "Dynamic causal modelling and network discovery"
- Vicente Pallares (U Pompeu Fabra, Barcelona, Spain) "Whole-brain effective connectivity from fMRI restingstate data discriminates between individuals"
- Thomas Bolton (EPFL, Lausanne, Switzerland) "Shedding light on resting-state dynamic functional network interactions by sparse coupled hidden Markov models"

W9 Emerging Models in Scientific Communication and Discussion

Room K.201, Thur 9:00 to 18:00

Romain Brette, Institut de la Vision, Paris

The academic publishing system is undergoing large changes towards a more open process, including the increasing use of preprints, open access and open data repositories. This move is especially important for computational and theoretical neuroscience, which require the availability of empirical data and model code. Several recent experiments aim at opening the scientific discussion itself, where not only the article but also the reaction of the community is published. Different models are being experimented; anonymous or signed reviews; invited or spontaneous; led by authors or editors. It has also been suggested that the social web (e.g. reddit, stackexchange) might provide relevant models.

In this workshop, we will explore emerging open models in academic publishing, with speakers presenting concrete experiments. Ample room will be reserved for discussion, in particular as it relates to concrete projects for the computational neuroscience community.

- Romain Brette (Vision Institute, Paris), "Decoupling peer review and editorial selection"
- Paola Masuzzo (Ghent U, Belgium), "Do you speak open science?"
- Stephen Eglen (Cambridge U, UK, "Encouraging code and data sharing in neuroscience"
- Nicolas Rougier (Neurodegeneratives Diseases Institute, Inria, Bordeaux, France), "ReScience: Reproducible science is good. Replicated science is better."
- Brandon Stell (CNRS, Paris), "PubPeer, the online journal club"
- Thomas Ingraham (F1000 Research), "Reforming and modernizing wasteful and antiquated practices in academic publishing"
- Frances Skinner (Krembil Research Institute, Canada), "eLife from a computational neuroscience insider's perspective"
- Thierry Galli (Institut Jacques Monod, Paris), "ASAPbio: accelerating science and publication in biology"

W10 Reaction-diffusion Modeling for Neurobiology

Room K.203, Thur 9:00 to 18:00

Robert McDougal, Yale University William W Lytton, SUNY Downstate Avrama Blackwell, George Mason

Methods developed for Computational Systems Biology are finding increasing use in Computational Neuroscience to understand the details of molecular cascades that influence neuron electrical activity – many and likely most voltage-sensitive ion channels and ligand sensitive channels (synapses) are modulated by such cascades. Additionally, examination of metabolomics requires consideration of the role of ATP in the functioning of cells for applications to a variety of brain diseases. This workshop on reaction-diffusion modeling will provide an introduction to the different types research questions being addressed using reaction-diffusion modeling. It will also identify the various technical approaches and bottlenecks encountered in addressing reaction-diffusion problems – stochastic vs deterministic simulation, simulation at 1D, 2D and 3D, tetrahedra vs cubic arrays, difficulties of coupling to membrane mechanisms and challenges of parallelization.

- William W Lytton (SUNY Downstate) "Reaction-diffusion foundations of multiscale modeling for the nervous system"
- Jim Schwaber (Thomas Jefferson) "Multiscale models of schizophrenia: gene networks to information in neuronal network"
- Marja-Leena Linne (Tampere University of Technology) "Model order reduction techniques with applications to describing reaction kinetics in neuronal and glial cells"
- Ekaterina Brocke (KTH Royal Institute of Technology) "Numerical discretization schemes for co-simulation of coupled electrical chemical systems"
- Davide Lillo (Marseille) "Slow variables of epileptiform activity: metabolic candidates and computational properties"
- Joanna Jedrzejewska-Szmek (George Mason) "NeuroRD as an approach for large scale modeling of neuronal signaling pathways."
- Erik De Schutter (OIST) "Mesh based neuron modeling using STEPS: fully integrated stochastic molecular and electrophysiological simulation in 3D"
- Robert A McDougal (Yale) "Using NEURON to incorporate reaction-diffusion into cellular and network models"

W11 Recent Developments in Epilepsy Modeling

Room K.103, Thur 9:00 to 18:00

<u>Wim van Drongelen</u>, The University of Chicago **Stephan A. van Gils**, University of Twente

Epilepsy is a chronic disease of the CNS characterized by the occurrence of seizures. It affects 1% of the people worldwide, and a significant proportion of the affected population does not respond to anticonvulsant drugs. This limited success of anticonvulsant treatment is, in part, due to our limited understanding of the underlying mechanisms that are responsible for the pathological behavior.

Translational research can play an important role in the unravelling of the mechanisms that may be responsible. As more and better experimental data become available, it is important to be able to integrate these data with modeling approaches in order to test hypotheses and to improve insight and produce more individualized models. In this workshop a number of active researchers in the field come together to discuss their latest results ranging from computational models on different scales to statistical methods to evaluate the use of biomarkers.

- Wim van Drongelen (U Chicago) "A recipe for seizures and a possible "pacebreaker""
- Stephan van Gils (U Twente) "The cross-scale effects of neural interactions during human neocortical seizure activity"
- Marc Goodfellow (U Exeter) "The role of networks in seizure generation"
- Geertjan Huiskamp (U Medical Center Utrecht) and Jurgen Hebbink (U Twente) "Networks of early and delayed responses to single pulse electrical stimulation of the cortex"
- Viktor Jirsa (U Marseille) "On nature of seizure dynamics"
- Bill Lytton (Downstate Medical Center) "Multiscale modeling of epilepsy: opportunities for drug discovery"
- Stiliyan Kalitzin (SEIN) "Reconstructive computational modelling for identifying epileptic networks in neurophysiological data"
- Wytze Wadman (U Amsterdam) "Ionic homeostasis in epilepsy and the balance between excitation and inhibition"
- Fabrice Wendling (U Rennes 1) "Novel stimulation protocols for probing neural network excitability: from computational modeling to clinical application"

W12 Neuroscience Gateway: Enabling Developers and Users to Utilize Open High Performance Computing Resources for Large Scale Simulations

Room K.101, Thur 14:00 to 18:00

Amit Majumdar, University of California San Diego, La Jolla Subhashini Sivagnanam, University of California San Diego, La Jolla Ted Carnevale, Yale University

The US National Science Foundation (NSF) funded Neuroscience Gateway (NSG) catalyzes computational neuroscience research by lowering the administrative and technical barriers that make it difficult for researchers to access open supercomputer resources for large scale simulations and data processing. It provides free and open access via a web portal and programmatically to supercomputers and time on the supercomputers is acquired via the peer reviewed process of the Extreme Science and Engineering Discovery Environment in the US. It has been in operation since early 2013, it has over 450 registered users. For the 2017 calendar year alone NSG was awarded 10,000,000 core hours on various supercomputers in the US. NSG already has large number of tools and software (NEURON, PGENESIS, NEST, BRIAN, PyNN, MOOSE, Freesurfer, R, Octave, Matlab etc.), libraries (BluePyOpt, CARLsim, Tensorflow etc.) and pipelines (The Virtual Brain Pipeline etc.). NSG is open to any user from anywhere in the world. Developers (of tools, libraries, and pipelines) and users utilize NSG extensively. This workshop will bring together both the users and the developers of tools/libraries/pipelines associated with the NSG for discussion of tools, software and research where HPC resources are utilized for neuroscience.

- Amit Majumdar, Subhashini Sivagnanam (UC San Diego), Ted Carnevale (Yale University) "Introduction to the Neuroscience Gateway"
- Salvador Dura-Bernal, William W Lytton (SUNY Downstate), Samuel A Neymotin (Brown U) "Parallel simulation of NEURON-based large scale network models"
- M Migliore, CA Lupascu, LL Bologna, R Migliore (Institute of Biophysics, National Research Council, Palermo, Italy) "Interaction of the Neuroscience Gateway with the Brain Simulation Platform of the European Human Brain Project: practical examples"
- Padraig Gleeson (University College London) "Using models from the Open Source Brain repository on the NSG portal infrastructure"
- Alexandar Peyser (Jülich Supercomputing Centre, Institute for Advanced Simulation, Forschungszentrum Jülich) "Towards exascale computing in neuroscience: NEST, NestMC and TVB"
- Christina M. Weaver (Franklin & Marshall College) "Modeling the effects of aging and neurodegeneration on cortical and striatal neurons"
- Marianne Bezaire (Boston University) "Full-scale detailed modeling of a hippocampal CA1 network using the Neuroscience Gateway"

W13 Cortical Function: Towards Understanding and Developing Integrative Theories

Room K.102, Thur 14:00 to 18:00

<u>Hamish Meffin</u>, The University of Melbourne **Anthony Burkitt**, The University of Melbourne

Understanding how our brain computes and analyses sensory inputs from our external environment whilst enabling us to experience such rich and varied mental lives is one of the great scientific challenges of the 21st Century. Recent advances have uncovered much about the cerebral cortex, with its 2-4mm thick sheet of neurons having a consistent anatomical structure consisting of six well-characterised layers and network connectivity. This workshop aims to look at what progress has been made in understanding how the cortex functions and what general integrative principles underlie how it works and enable capabilities as diverse as sensory perception, control of voluntary motor activity and high-level cognitive functions.

- Marcus Diesmann (Institute of Neuroscience and Medicine, Research Centre Jýlich, Germany) "A brain-scale model of macaque visual cortex at cellular and synaptic resolution"
- Jorge Mejias (Center for Neural Science, New York University, USA) "Large-scale models of cortical dynamics: neural communication and cognitive computations"
- Subutai Ahmad (VP Research Numenta, USA) "Why the neocortex has layers and columns, a theory of learning 3D models of the world"
- Stefan Mihalas (Allen Institute for Brain Science, USA) "Cortical circuits implement optimal context integration and its gating"

W14 Postdoc Career Workshop

Room K.102, Thur 9:00 to 12:30

Joanna Jedrzejewska-Szmek, University of Warsaw

Computational neuroscience is a diverse, international and interdisciplinary community, which allows for diverse and challenging career paths. This workshop is intended to provide postdocs and students in computational neuroscience an opportunity to interact with panel of mentors with diverse careers in research and industry. Mentors will discuss their own experiences, talk about different professional paths and funding opportunities, working abroad, and give advice how to survive in both 'fat' and 'lean' funding periods. The panel of mentors will also answer questions about available career choices, their advantages and disadvantages, and relative importance of skills, work experience and teaching.

- Piotr Franaszczuk (US Army Research Laboratory Human Research and Engineering Directorate)
- Eugene Izhikevich (Brain Corporation)
- Malin Sandström (International Neuroinformatics Coordinating Facility (INCF))
- Eleni Vasilaki (University of Sheffield)
- Christina Weaver (Franklin & Marshall College)

Posters

Poster Listing

Sunday Posters Posters P1 – P104

P1 Potential functions of different temporal patterns of intermittent neural synchronization

Leonid Rubchinsky^{1,2*}, Sungwoo Ahn³

P2 NestMC: A morphologically detailed neural network simulator for modern high performance computer architectures

Wouter Klijn¹, Benjamin Cumming², Stuart Yates², Vasileios Karakasis³, and Alexander Peyser¹*

P3 Automatically generating HPC-optimized code for simulations using neural mass models

Marmaduke Woodman¹, Sandra Diaz-Pier^{2*}, and Alexander Peyser²

P4 Conjunction or co-activation? A multi-level MVPA approach to task set representations

James Deraeve¹[⋆], Eliana Vassena², and William Alexander¹

P5 Understanding Adaptation in Human Auditory Cortex with Modeling

David Beeman^{1*}, Pawel Kudela², Dana Boatman-Reich^{3,4}, and William Anderson²

P6 Silent and bursting states of Purkinje cell activity modulate VOR adaptation

Niceto Luque^{1,2,3*}, Francisco Naveros⁴, Richard Carrillo⁴, Eduardo Ros⁴, and Angelo Arleo^{1,2,3}

¹ Department of Mathematical Sciences, Indiana University Purdue University Indianapolis, Indianapolis, IN, USA

²Stark Neurosciences Research Institute, Indiana University School of Medicine, Indianapolis, IN, USA

³Department of Mathematics, East Carolina University, Greenville, NC, USA

¹ Jülich Supercomputing Centre, Forschungszentrum Jülich, Jülich, Germany

²Future Systems, Swiss National Supercomputing Centre, Zürich, Switzerland

³User Engagement & Support, Swiss National Supercomputing Centre, Lugano, Switzerland

¹ Institut de Neurosciences des Systèmes, Aix Marseille Université, Marseille, France

²Simulation Lab Neuroscience, Forschungszentrum Jülich, Jülich, Germany

¹Department of Experimental Psychology, Ghent University, Ghent, Belgium

²Donders Center for Cognitive Neuroimaging, Radboud University, Nijmegen, Netherlands

¹Department of Electrical, Computer, and Energy Engineering 1, University of Colorado, Boulder, CO, USA

²Department of Neurosurgery, Johns Hopkins School of Medicine, Baltimore, MD, USA

³Department of Neurology, Johns Hopkins School of Medicine, Baltimore, MD, USA

⁴Department of Otolaryngology, Johns Hopkins School of Medicine, Baltimore, MD, USA

¹ INSERM. U968. Paris. France

² Sorbonne Universités, UPMC University Paris 06, UMR S 968, Institut de la Vision, Paris, France

³CNRS, UMR 7210, Paris, France

⁴Department of Computer Architecture and Technology, University of Granada (CITIC), Granada, Spain

P7 The "convis" framework: Population Simulation of the Visual System with Automatic Differentiation using theano

Jacob Huth^{1*}, Timothee Masquelier², and Angelo Arleo¹

P8 Why does neural activity in ASD have low complexity: from a perspective of a small-world network model

Koki Ichinose¹*, Jihoon Park¹, Yuji Kawai¹, Junichi Suzuki¹, Hiroki Mori², and Minoru Asada¹

P9 Phase-locked mode prediction with generalized phase response curve

Sorinel A Oprisan*, Dave I Austin

Department of Physics and Astronomy, College of Charleston, Charleston, SC, USA

P10 Neural Field Theory of Corticothalamic Prediction and Attention

Tahereh Babaie^{1,2}*, Peter Robinson^{1,2}

P11 Top-down dynamics of cortical pitch processing explain the emergence of consonance and dissonance in dyads

Alejandro Tabas^{1,2*}, Martin Andermann³, André Rupp³, and Emili Balaguer-Ballester^{2,4}

P12 Modeling sensory cortical population responses in the presence of background noise

Henrik Lindén^{1*}, Rasmus K. Christensen¹, Mari Nakamura², and Tania R. Barkat²

P13 Cortical circuits from scratch: A metaplastic rule for inducing lognormal firing rates in a cortical model

Zachary Tosi¹*, John Beggs²

P14 Investigating the effects of horizontal interactions on RGCs responses in the mice retina with high resolution pan-retinal recordings

Davide Lonardoni¹*, Fabio Boi¹, Stefano Di Marco², Alessandro Maccione¹, and Luca Berdondini¹

¹Sorbonne Universités, UPMC Univ Paris 06, INSERM, CNRS, Institut de la Vision, Paris, France

²CERCO UMR5549, CNRS, University Toulouse, France

¹Department of Adaptive Machine Systems, Osaka University, Osaka, Japan

²Department of Computer Science, University of Cergy-Pontoise, Cergy-Pontoise, France

¹School of Physics, Faculty of Science, University of Sydney, Sydney, Australia

²Center of Excellence for Integrative Brain Function, Australian Research Council, Australia

¹Max Planck Institute for Human Cognitive and Brain Sciences, Saxony, Leipzig, Germany

²Dept. of Computing and Informatics. Faculty of Science and Technology, Bournemouth University, UK

³Biomagnetism Section, Heidelberg University, Baden-Württemberg, Germany

⁴Bernstein Centre for Computational Neuroscience Heidelberg-Mannheim, Heidelberg University, Germany

¹Center for Neuroscience, University of Copenhagen, Copenhagen, Denmark

²Brain and Sound Lab, Department of Biomedicine, Basel University, Basel, Switzerland

¹Cognitive Science, Indiana University, Bloomington, IN, USA

²Physics, Indiana University, Bloomington, IN, USA

¹Neuroscience and Brain Technology Department, Fondazione Istituto Italiano di Tecnologia, Genova, Italy

²Scienze cliniche applicate e biotecnologiche, Università dell'Aquila, L'Aquila, Italy

P15 Calcium base plasticity rule can predict plasticity direction for a variety of stimulation paradigms

Joanna Jedrzejewska-Szmek^{1*}, Daniel Dorman^{1,2}, and Kim Avrama Blackwell^{1,2}

P16 Unstructured network topology begets privileged neurons and rank-order representation

Christoph Bauermeister^{1,2*}, Hanna Keren^{3,4}, and Jochen Braun^{1,2}

P17 X047 Finer parcellation reveals intricate correlational structure of resting-state fMRI signals

Joao Vicente Dornas*, Jochen Braun

Institute of Biology, Otto von Guericke University, Magdeburg, Saxony-Anhalt, Germany

P18 Modelling human choices: MADeM and decision-making

Eirini Mavritsaki^{1,2*}, Silvio Aldrovandi¹, and Emma Bridger¹

P19 The interplay between synaptic plasticity and firing rate adaptation sharpens response dynamics with visual learning

Sukbin Lim¹*, Nicolas Brunel^{2,3}

P20 Adaptation and inhibition control the pathologic synchronization in the model of a focal epileptic seizure.

Anatoly Buchin^{1,2*}, Clifford Charles Kerr³, Anton Chizhov^{4,5}, Gilles Huberfeld^{6,7}, Richard Miles⁸, and Boris Gutkin^{9,10}

P21 Efficient and Effective Neural Activity Shaping for a Retinal Implant

Martin Spencer¹*, Hamish Meffin^{1,2}, Tatiana Kameneva¹, David B Grayden¹, and Anthony N Burkitt¹

¹Krasnow Institute, George Mason University, Fairfax, VA, USA

²Bioengineering Department, George Mason University, Fairfax, VA, USA

¹ Institute of Biology, Otto-von-Guericke University, Magdeburg, Germany

²Center for Behavioral Brain Sciences, Magdeburg, Germany

³Network Biology Research Laboratory, Technion - Israel Institute of Technology, Haifa, Israel

⁴Department of Physiology, Technion - Israel Institute of Technology, Haifa, Israel

¹Department of Psychology, Birmingham City University, Birmingham, UK

²School of Psychology, University of Birmingham, Birmingham, UK

¹Neural and Cognitive Sciences, NYU Shanghai, Shanghai, China

²Department of Neurobiology, University of Chicago, Chicago, Illinois, USA

³Department of Statistics, University of Chicago, Chicago, Illinois, USA

¹Department of Physiology and Biophysics, University of Washington, Seattle, WA, USA

²Allen Institute for Brain Science, Seattle, WA, USA

³SUNY Downstate Medical Center, New York City, NY, USA

⁴Computational Physics Laboratory, Ioffe Institute, St Petersburg, Russian Federation

⁵Sechenov Institute of Evolutionary Physiology and Biochemistry, St Petersburg, Russian Federation

⁶ Pitié-Salpêtrière Hospital, University Pierre and Marie Curie, Paris, France

⁷Inserm U1129 Infantile Epilepsies and Brain Plasticity, Paris Descartes University, Paris, France

⁸Cortex and Epilepsy Group, Brain and Spine Institute, Paris, France

⁹Department of Cognitive Neuroscience, Group for Neural Theory, École Normale Supérieure, Paris, France

¹⁰Center for Cognition and Decision Making, NRU Higher School of Economics, Moscow, Russian Federation

¹Department of Biomedical Engineering, University of Melbourne, Melbourne, Australia

²NVRI, Department of Optometry & Vision Sciences, University of Melbourne, Melbourne, Australia

P22 Application of control theory to neural learning in the brain

Catherine Davey^{1*}, David B Grayden^{1,2}, and Anthony N Burkitt¹

P23 Modeling dynamic oscillations: A method of inferring neural behavior through mean field network models

Tao Liangyu^{1*}, Vineet Tiruvadi^{1,2}, Rehman Ali⁴, Helen Mayberg³, and Rob Butera¹

P24 Synaptic strengths dominate phasing of motor neurons by a central pattern generator

Cengiz Gunay^{1,2*}, Anca Doloc-Mihu¹, Damon G Lamb^{1,3}, and Ronald L Calabrese¹

P25 PumpHCO-db: A database of half-center oscillator computational models for analyzing the influence of Na+/K+ pump on the bursting activity

Anca Doloc-Mihu*, Ronald Calabrese

Department of Biology, Emory University, Atlanta, GA, USA

P26 Encoding of memories: effective connectivity on the hippocampus and the role of inhibition in the information flow.

Victor J. Lopez-Madrona^{1*}, Fernanda S. Matias², Ernesto Pereda³, Claudio Mirasso⁴, and Santiago Canals¹

P27 Extended generalized leaky integrate and fire neuron for cerebellum modeling

Alice Geminiani^{1*}, Alessandra Pedrocchi¹, Egidio Dangelo², and Claudia Casellato¹

P28 Saccade Velocity Driven Oscillatory Networkmodel of Grid cells

Ankur Chauhan*, Karthik Soman, and Srinivasa Chakravarthy

Department of Biotechnology, Indian Institute of Technology Madras, Chennai, Tamilnadu, India

¹Department of Biomedical Engineering, University of Melbourne, Victoria, Australia

²Centre for Neural Engineering, University of Melbourne, Victoria, Australia

¹Department of Biomedical Engineering, Georgia Institute of Technology, Atlanta, GA, USA

²Department of Biomedical Engineering, Emory University, Atlanta, GA, USA

³Department of Psychiatry and Behavioral Sciences, Emory University, Atlanta, GA, USA

⁴Department of Electrical Engineering, Stanford University, Stanford, CA, USA

¹Department of Biology, Emory University, Atlanta, GA, USA

²School of Science and Technology, Gerogia Gwinnett College, Lawrenceville, GA, USA

³Department of Neurology, Univ. Florida, Gainesville, FL, USA

¹ Instituto de Neurociencias, Consejo Superior de Investigaciones Científicas, Universidad Miguel Hernández, Sant Joan d'Alacant, Spain

²Instituto de Física, Universidade Federal de Alagoas, Maceió, Alagoas, Brazil

³Departamento de Ingeniería Industrial, Escuela Superior de Ingeniería y Tecnología, Universidad de La Laguna Avda. Astrofísico Fco. Sanchez, s/n, La Laguna, Tenerife, Spain

⁴Instituto de Física Interdisciplinar y Sistemas Complejos, CSIC-UIB, Campus Universitat de les Illes Balears, Palma de Mallorca, Spain

¹NEARLab, Dept. of Electronics, Information and Bioengineering, Politecnico di Milano, Milan, Italy

²Department of Brain and Behavioral Sciences, University of Pavia, Pavia, Italy

P29 Programmed cell death in substantia nigra due to subthalamic nucleus-mediated excitotoxicity: a computational model of Parkinsonian neurodegeneration

Vignayanandam Muddapu*, Srinivasa Chakravarthy

Bhupat and Jyoti Mehta School of Biosciences, Department of Biotechnology, IIT-Madras, Chennai, TN, India

P30 A novel approach for determining how many distinct types of neurons are in the Drosophila brain by sequencing neural structure

Chaochun Chuang*, Nan-Yow Chen

National center for high-performance computing, Taiwan

P31 Generating sequences in recurrent neural networks for storing and retrieving episodic memories

Mehdi Bayati^{1,2*}, Jan Melchior¹, Laurenz Wiskott¹, and Sen Cheng^{1,2}

P32 Modeling replay and theta sequences in a 2-d recurrent neural network with plastic synapses

Amir Hossein Azizi1*, Kamran Diba2, and Sen Cheng1

P33 Biophysically detailed model of cortical activity in response to moving gratings

Elena Smirnova^{1,2*}, Elena Yakimova³, and Anton Chizhov^{1,2}

P34 NeuriteSLIM – Shrink the Neuro Fibers for Visualization the Connectome

Nan-Yow Chen^{1*}, Chi-Tin Shih², and Chaochun Chuang¹

P35 Identification of models of sensory neural circuits consisting of a nonlinear filter in series with a leaky integrate-and-fire neuron

Dorian Florescu, Daniel Coca*

Department of Automatic Control and Systems Engineering, University of Sheffield, Sheffield, UK

P36 Modelling fluctuations in resting-state functional connectivity in epilepsy

Julie Courtiol*, Spase Petkoski, and Viktor K Jirsa

Aix Marseille Univ, Inserm, INS, Institut de Neurosciences des Systèmes, Marseille, France

P37 Exact solutions to a Wilson-Cowan network of excitatory and inhibitory neurons whose dynamics is triggered by one single spike

Roberto Covolan*

Department of Neurology, State University of Campinas, Campinas, SP, Brazil

¹ Institut für Neuroinformatik, Ruhr-Universität Bochum, Bochum, Germany

²Mercator Research Group 'Structure of Memory', Ruhr-University Bochum, Bochum, Germany

¹ Institut für Neuroinformatik, Ruhr University Bochum (RUB), Bochum, Germany

²Psychology faculty, university of Wisconsin-Milwaukee, WI, USA

¹ Ioffe Institute, St. Petersburg, Russia

²Sechenov Institute of Evolutionary Physiology and Biochemistry of RAS, St. Petersburg, Russia

³Pavlov Institute of Physiology, St. Petersburg, Russia

¹ High Performance Computing Division, National Center for High-Performance Computing, Taiwan

²Department of Applied Physics, Tunghai University, Taiwan

P38 Encoding variable cortical states with short-term spike patterns

Bartosz Telenczuk¹*, Richard Kempter², Gabriel Curio³, and Alain Destexhe¹

P39 Cat Paw-shaking as a Transient Response to Sensory Input to Locomotion CPG

Jessica Parker^{1*}, Alexander Klishko², Boris Prilutsky², and Gennady Cymbalyuk¹

P40 Population Coding with Two-Dimensional Feature Maps in the Retina

Felix Franke^{1*}, Andreas Hierlemann¹, and Rava Azeredo Da Silvera^{2,3}

P41 A detailed computational reconstruction of the cerebellum granular layer network predicts large scale spatiotemporal dynamics of neuronal activity

Stefano Casali¹, Stefano Masoli¹*, Martina Francesca Rizza^{1,3}, and Egidio Dangelo^{1,2}

P42 A Biophysically Detailed Cerebellar Stellate Neuron Model Predicts Local Synaptic Interactions

Martina Francesca Rizza^{1,2*}, Stefano Masoli¹, and Egidio Dangelo^{1,3}

P43 Neuromodulation of Subgenual Cingulate Activity Localizable from EEG

Yinming Sun^{1,2⋆}, Willy Wong^{1,3}, Faranak Farzan², Daniel Blumberger^{2,4}, and Zafiris Daskalakis^{2,4}

P44 Phase dynamics in a GO/NOGO finger tapping task

Svitlana Popovych^{1,2*}, Shivakumar Viswanathan^{2,3}, Nils Rosjat^{1,2}, Christian Grefkes^{2,3}, Gereon Fink^{2,3}, and Silvia Daun^{1,2}

¹ Unité de Neurosciences, Information et Complexité, CNRS, 91198 Gif-sur-Yvette, France; European Institute for Theoretical Neuroscience, CNRS, Paris, France

²Institute for Theoretical Biology, Humboldt-Universität zu Berlin, Berlin, Germany

³Dept. of Neurology, Universitätsmedizin Charité, Berlin, Germany

¹Neuroscience Institute, Georgia State University, Atlanta, GA, USA

²School of Biological Sciences, Georgia Institute of Technology, Atlanta, GA, USA

¹Department of Biosystems Science and Engineering, ETH Zürich, Basel, Switzerland

²Ecole Normale Supérieure, Paris, France

³Centre National de la Recherche Scientifique, Paris, France

¹Department of Brain and Behavioral Sciences, University of Pavia, Pavia, Italy

²Brain Connectivity Center, C. Mondino National Neurological Institute, Pavia, Italy

³Dipartimento di Informatica, Sistemistica e Comunicazione, Università degli Studi di Milano-Bicocca, Viale Sarca, Italy

¹Department of Brain and Behavioral Sciences, University of Pavia, Pavia, Italy

²Dipartimento di Informatica, Sistemistica e Comunicazione, Università degli Studi di Milano-Bicocca, Milan, Italy

³Brain Connectivity Center, Istituto Neurologico IRCCS C. Mondino, Pavia, Italy

¹Institute of Biomaterials and Biomedical Engineering, University of Toronto, Toronto, ON, M5S3G9, Canada

²Centre for Addiction and Mental Health, Toronto, ON, M5T1R8, Canada

³Dept. of Electrical and Computer Engineering, University of Toronto, Toronto, ON, M5S3G4, Canada

⁴Dept. of Psychiatry, University of Toronto, Toronto, ON, M5S3G4, Canada

¹Heisenberg Research Group of Computational Neuroscience - Modeling Neural Network Function, Department of Animal Physiology, Institute of Zoology, University of Cologne, Cologne, Germany

²Cognitive Neuroscience, Institute of Neuroscience and Medicine (INM-3), Research Center Juelich, Juelich, Germany

³Department of Neurology, University Clinic Cologne, Cologne, Germany

P45 Mechanisms of focal seizure generation in a realistic small-network model with ionic dynamics

Damiano Gentiletti¹*, Piotr Suffczynski¹, Vadym Gnatkovski², and Marco de Curtis²

P46 Pre-allocation of working memory modulates memory performance

Hyeonsu Lee^{1*}, Woochul Choi^{1,2}, and Se-Bum Paik^{1,2}

P47 Temporal dynamics of bistable perception reveals individual time window for perceptual decision making

Woochul Choi*, Se-Bum Paik

P48 Regularly structured retinal mosaics can induce structural correlation between orientation and spatial frequency maps in V1

Jaeson Jang¹*, Se-Bum Paik^{1,2}

P49 Distinct role of synaptic and nonsynaptic plasticity in memory ensemble formation, allocation, and linkage

Youngiin Park1*, Se-Bum Paik1,2

P50 Frequency- and Location-Dependence of Auditory Influence on Human Visual Perception

Jun Ho Song*, Se-Bum Paik

P51 Developmental model for ocular dominance column seeded from retinal

Min Song^{1,2}*, Se-Bum Paik^{1,2}

P52 Reliability of effective connectivity from fMRI resting-state data: discrimination between individuals

Vicente Pallares^{1*}, Matthieu Gilson¹, Simone Kuhn², Andrea Insabato¹, and Gustavo Deco¹

¹Department of Experimental Physics, University of Warsaw, Warsaw, Poland

²Istituto Neurologico Carlo Besta, Milan, Italy

¹Department of Bio and Brain Engineering

²Program of Brain and Cognitive Engineering, Korea Advanced Institute of Science and Technology, Daejeon, Republic of Korea

¹ Department of Bio and Brain Engineering, Korea Advanced Institute of Science and Technology, Daejeon, Republic of Korea

²Program of Brain and Cognitive Engineering, Korea Advanced Institute of Science and Technology, Daejeon, Republic of Korea

¹ Department of Bio and Brain Engineering, Korea Advanced Institute of Science and Technology, Daejeon, Republic of Korea

²Program of Brain and Cognitive Engineering, Korea Advanced Institute of Science and Technology, Daejeon, Republic of Korea

¹Department of Bio and Brain Engineering

²Program of Brain and Cognitive Engineering, KAIST, Daejeon, Republic of Korea

¹Department of Information and Communication Technologies, Universitat Pompeu Fabra, Barcelona, Spain

²Max Plank Institute for Human Development, Berlin, Germany

P53 Temporal dynamics of resting state networks on a whole-brain level

Katharina Glomb^{1*}, Adrian Ponce-Alvarez¹, Matthieu Gilson¹, Petra Ritter², and Gustavo Deco^{1,3}

P54 Non-parametric estimation of network connectivity using MVAR processes in multiunit activity

Matthieu Gilson¹*, Adria Tauste Campo^{1,2}, Alexander Thiele³, and Gustavo Deco^{1,4}

P55 Dependence of Absence Seizure Dynamics on Physiological Parameters

Farah Deeba^{1,2*}, Paula Sanz-Leon^{1,2}, and Pa Robinson^{1,2}

P56 NEST-SpiNNaker comparison of large-scale network simulations

Sacha J van Albada^{1*}, Andrew Rowley², Johanna Senk¹, Michael Hopkins², Maximilian Schmidt^{1,3}, Alan Stokes², David Lester², Steve Furber², and Markus Diesmann^{1,4,5}

P57 Temporal processing in the cerebellar cortex enabled by dynamical synapses

Alessandro Barri^{1*}, Martin Wiechert², and David Digregorio¹

P58 Emergence of perceptual invariances in biological sensory processing

Alexander G Dimitrov*

Department of Mathematics and Statistics, Washington State University Vancouver, Vancouver, WA, USA

P59 A non-linear stochastic strategy to estimate synaptic conductances under the presence of subthreshold ionic currents.

Catalina Vich Llompart^{1*}, Rune W. Berg², Antoni Guillamon³, and Susanne Ditlevsen⁴

¹Center for Brain and Cognition, Dept. of Technology and Information, Universitat Pompeu Fabra, Barcelona, Spain

²Dept. of Neurology, Charité - University Medicine, Charitéplatz, Berlin, Germany

³Institució Catalana de la Recerca i Estudis Avançats, Universitat Barcelona, Barcelona, Spain

¹Computational Neuroscience Group, Dept. de Tecnologies de la Informació i les Comunicacions, Universitat Pompeu Fabra, Barcelona, Spain

²Epilepsy Monitoring Unit, Department of Neurology, Hospital del Mar Medical Research Institute, Barcelona, Spain

³Institute of Neuroscience, Newcastle University, Newcastle upon Tyne, UK

⁴Institució Catalana de Recerca i Estudis Avançats, Barcelona, Spain

¹School of Physics, University of Sydney, Sydney, Australia

²Center for Integrative Brain Function, University of Sydney, Sydney, Australia

¹Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6), Jülich Research Centre and JARA BRAIN Institute I, Jülich, Germany

²School of Computer Science, University of Manchester, Manchester, UK

³Laboratory for Neural Circuit Theory, RIKEN Brain Science Institute, Wako, Japan

⁴Department of Psychiatry, Psychotherapy and Psychosomatics, Medical Faculty, RWTH Aachen University, Aachen, Germany

⁵Department of Physics, Faculty 1, RWTH Aachen University, Aachen, Germany

¹Unite d'Imagerie Dynamique du Neurone, Institut Pasteur, Paris, France

²Department of Physiology, Universität Bern, Switzerland

¹Dept. of Mathematics and Computer Science, Universitat de les Illes Balears, Palma, Spain

²Dept. of Neuroscience and Pharmacology, University of Copenhagen, Copenhagen, Denmark

³Dept. of Applied Mathematics I, EPSEB, Universitat Politècnica de Catalunya, Barcelona, Spain

⁴Dept. of Mathematical Science, University of Copenhagen, Copenhagen, Denmark

P60 Involvement of randomness in reinforcement learning

Romain Cazé*, Benoît Girard, and Stéphane Doncieux

ISIR. Université Pierre et Marie Curie, Paris, France

P61 Modelling the impact of dendritic spine geometry on electrical and calcic signalling with the Finite Element Method

Nicolas Doyon*, Frank Boahen

Department of Mathematics and Statistics, Laval University, Quebec, Canada

P62 Resilience in dynamical neural networks with synaptic adaptation

Patrick Desrosiers^{1,2*}, Edward Laurence², Nicolas Doyon^{1,3}, and Louis J. Dubé²

P63 Cell assemblies: a computational challenge

Eleonora Russo*, Daniel Durstewitz

Department of Theoretical Neuroscience, ZI - Central Institute for Mental Health, Mannheim, Germany

P64 Reconstructing neural dynamics from experimental data using radial basis function recurrent neural networks

Dominik Schmidt*, Daniel Durstewitz

Department of Theoretical Neuroscience, Bernstein Center for Computational Neuroscience, Central Institute of Mental Health, Medical Faculty Mannheim, Heidelberg University, Germany

P65 Layer V pyramidal cells as mediators of delta oscillations: Insights from biophysically detailed modeling and connections with schizophrenia genetics.

Tuomo Mäki-Marttunen¹*, Florian Krull¹, Francesco Bettella¹, Christoph Metzner², Anna Devor^{3,4}, Srdjan Djurovic⁵, Anders M. Dale^{3,4}, Ole A Andreassen¹, and Gaute T. Einevoll^{6,7}

P66 Biophysical modeling of single-neuron contributions to ECoG and EEG signals

Solveig Næss^{1,2}, Torbjørn V Næss³, Geir Halnes³, Eric Halgren⁴, Anders M. Dale⁴, and Gaute T. Einevoll^{3,5}*

¹Centre de recherche de l'Institut universitaire en santé mentale de Québec, Québec, Québec, Canada

²Département de physique, de génie physique et d'optique, Université Laval, Québec, Québec, Canada

³ Département de mathématiques et de statistique, Université Laval, Québec, Québec, Canada

¹NORMENT, Institute of Clinical Medicine, University of Oslo, Norway

²Centre for Computer Science and Informatics Research, University of Hertfordshire, Hatfield, UK

³Department of Neurosciences, University of California San Diego, La Jolla, CA, USA

⁴Department of Radiology, University of California San Diego, La Jolla, CA, USA

⁵Department of Medical Genetics, Oslo University Hospital, Oslo, Norway

⁶Faculty of Science and Technology, Norwegian University of Life Sciences, Ås, Norway

⁷Department of Physics, University of Oslo, Norway

¹Department of Informatics, University of Oslo, Oslo, Norway

²Simula-UiO-UCSD Research and PhD (SUURPh) training program

³Faculty of Science and Technology, Norwegian University of Life Sciences, Ås, Norway

⁴Department of Neuroscience and Radiology, School of Medicine, UC San Diego, CA, USA

⁵Department of Physics, University of Oslo, Oslo, Norway

P67 Extracellular diffusion can introduce errors in current source density estimates

Geir Halnes 1* , Tuomo Maki-Marttunen 2 , Klas H Pettersen 3,4 , Ole A Andreassen 2 , and Gaute T. Einevoll 1,5

P68 Estimation of metabolic oxygen consumption from optical measurements in cortex

Marte Sætra^{1*}, Anders M Dale², Anna Devor², and Gaute T. Einevoll^{1,3}

P69 Computing Brain Signals: Concurrent simulation of network activity, extracellular electric potentials and magnetic fields.

Espen Hagen¹*, Solveig Næss², Torbjørn V Næss³, and Gaute T. Einevoll^{1,3}

P70 Integration of orientation and spatial frequency in a model of visual cortex

Alina Schiffer^{1*}, Axel Grzymisch¹, Malte Persike², and Udo A Ernst¹

P71 Performance-optimization guided distribution of attentional resources

Daniel Harnack, Udo A Ernst*

Computational Neuroscience Lab, Institute for Theoretical Physics, University Bremen, Germany

P72 Feature integration with critical dynamics in cortical subnetworks

Nergis Tomen*, Udo A Ernst

Computational Neuroscience Lab, Institute for Theoretical Physics, University of Bremen, Germany

P73 Interneuronal contribution to state transition in the mouse neocortex

Stefano Zucca^{1,2}, Valentina Pasquale^{3*}, Giuseppe Pica^{2,4}, Manuel Molano-Mazon^{2,4}, Michela Chiappalone³, Stefano Panzeri^{2,4}, and Tommaso Fellin^{1,2}

¹Faculty of Science and Technology, Norwegian University of Life Sciences, As, Norway

²NORMENT, Institute of Clinical Medicine, University of Oslo, Oslo, Norway

³Letten Centre and Glialab, Dept. of Molecular Medicine, Inst. of Basic Medical Sciences, University of Oslo, Oslo, Norway

⁴Centre for Molecular Medicine Norway, University of Oslo, Oslo, Norway

⁵Department of Physics, University of Oslo, Oslo, Norway

¹Department of Physics, University of Oslo, Oslo, Norway

²Department of Neurosciences, UC San Diego, La Jolla, CA, USA

³Faculty of Science and Technology, Norwegian University of Life Sciences, Ås, Norway

¹Department of Physics, University of Oslo, Oslo, Norway

²Department of Informatics, University of Oslo, Oslo, Norway

³Faculty of Science and Technology, Norwegian University of Life Sciences, Aas, Norway

¹Computational Neuroscience Lab, Institute for Theoretical Physics, Univ. of Bremen, Germany

²Department of Psychology, Methods Section, Johannes Gutenberg University Mainz, Germany

¹Optical Approaches to Brain Function Laboratory, Department of Neuroscience and Brain Technologies, Istituto Italiano di Tecnologia, Genova, Italy

²Neural Coding Laboratory, Istituto Italiano di Tecnologia, Genova and Rovereto, Italy

³Department of Neuroscience and Brain Technologies, Istituto Italiano di Tecnologia, Genova, Italy

⁴Neural Computation Laboratory, Center for Neuroscience and Cognitive Systems @UniTn, Istituto Italiano di Tecnologia, Rovereto, Italy

P74 Embodiment, connectivity, and critical states in neural systems

Kelvin Oie^{1*}, David Boothe¹, Joshua Crone¹, Alfred Yu¹, and Melvin Felton²

¹U.S. Army Research Laboratory, Aberdeen Proving Ground, MD, USA

P75 A computational model of temporal processing in the human auditory cortex

Isma Zulfigar^{1*}, Michelle Moerel^{1,2}, Peter de Weerd^{1,2}, and Elia Formisano^{1,2}

P76 The dependence of simulated local field potential (LFP) frequency content on local and long range connectivity

David Boothe¹, Alfred Yu¹, Joshua Crone¹, Melvin Felton¹, Kelvin Oie¹, and Piotr J Franaszczuk^{1,2*}

P77 Pre-whitening as a means to improve dimensionality reduction and simplify clustering in spikesorters for multi-electrode recordings

Roland Diggelmann^{1,2*}, Michele Fiscella^{1,2}, Andreas Hierlemann¹, and Felix Franke¹

P78 An integrative model explaining many functions of corticothalamic feedback

Domenico Guarino^{1*}, Jan Antolik^{1,2}, Andrew P Davison¹, and Yves Fregnac¹

P79 State-Dependent Control of Oscillatory Brain Dynamics

Benjamin Xavier Etienne¹, Flavio Frohlich², and Jérémie Lefebvre^{1,3}*

P80 Switch of preference as a signature of heterogeneous excitability of neurons in the primate prefrontal cortex

Encarni Marcos^{1*}, Maurizio Mattia², and Aldo Genovesio¹

P81 Modeling of the perceptual dynamics of the perception of body motion

Leonid Fedorov^{1,2*}, Tjeerd Dijkstra¹, Louisa Sting^{1,3}, Howard Hock⁴, and Martin Giese^{1,2}

²U.S. Army Research Laboratory, Adelphi, MD, USA

¹ Maastricht Centre for Systems Biology, Maastricht University, Maastricht, The Netherlands

²Department of Cognitive Neuroscience, Maastricht University, Maastricht, The Netherlands

¹US Army Research Laboratory, Aberdeen, Maryland, USA

²Department of Neurology, The Johns Hopkins University School of Medicine, Baltimore, Maryland, USA

¹Department of Biosystems Science and Engineering, ETH Zurich, Basel, Switzerland

²Neural Circuit Laboratories, Friedrich Miescher Institute for Biomedical Research, Basel, Switzerland

¹UNIC, CNRS-FRE3693, Gif-sur-Yvette, France

²Institut de la Vision, Paris, France

¹Krembil Research Institute, University Health Network, Toronto, Ontario, Canada

²Department of Psychiatry and Cell Biology, University of North Carolina at Chapel Hill, Chapel Hill, North Carolina, USA

³Department of Mathematics, University of Toronto, Toronto, Ontario, Canada

¹Department of Physiology and Pharmacology, Sapienza University of Rome, Rome, Italy

² Istituto Superiore di Sanità, Rome, Italy

¹ Section for Comput. Sensomotorics, Dept. Cognitive Neurology, CIN&HIH, Univ. of Tübingen, Germany

²GTC, International Max Planck Research School, University of Tübingen, Tübingen, Germany

³Department of Cognitive Science, University of Tübingen, Tübingen, Germany

⁴Center for Complex Systems and the Brain Sciences, Department of Psychology, Florida Atlantic University

P82 Modelling the effects of propofol on neuronal synchronization in network of interneurons

Laure Buhry^{1,2}, Clément Langlet², and Francesco Giovannini^{1,2}*

P83 The dynamical response properties of in silico neurons from the Blue Brain Project digitally reconstructed neocortical microcircuitry

Christophe Verbist¹*, Stefano Salvade^{1,2}, and Michele Giugliano¹

¹TNB, Department of Biomedical Sciences, University of Antwerp, Belgium

P84 Self-Organized Balanced Spiking Neural Networks To Encode Natural Stimuli

James Henderson^{1,2}[⋆], Pulin Gong^{1,2}

P85 Transiently attracting states in recurrent neural networks

Hendrik Wernecke*, Bulcsu Sandor, and Claudius Gros

Institute for Theoretical Physics, Goethe University, Frankfurt am Main, Germany

P86 Characterization of resting state dynamics in monkey motor cortex

Nicole Voges¹*, Paulina Dabrovska¹, Johanna Senk¹, Espen Hagen², Alexa Riehle^{1,3}, Thomas Brochier³, and Sonja Grün^{1,4}

P87 Sharp wave ripples as propagating patterns emerging from spatially extended neural circuits

Yifan Gu, Pulin Gong*

School of Physics and Australian Research Council Centre of Excellence for Integrative Brain Function, University of Sydney, NSW, Australia

P88 Macroscopic Phase-Resetting Curve of Spiking Neural Networks: Theory and Application

Gregory Dumont*, Boris Gutkin

Group for Neural Theory, Ecole Normale Supérieure, Paris, France

P89 Semi-numerical method for computationally effective analysis of working memory models

Nikita Novikov¹*, Boris Gutkin^{1,2}

¹Neurosys Team, LORIA, CNRS, INRIA CR Nancy Grand Est, Villers-lès-Nancy, France

²Université de Lorraine, Vandoeuvre-lès-Nancy, France

²DIBRIS, University of Genova, Genova, Italy

¹School of Physics, The University of Sydney, Sydney, NSW, Australia

²Centre for Integrative Brain Function, The University of Sydney, Sydney, NSW, Australia

¹ Institute of Neuroscience and Medicine (INM-6) & Institute for Advanced Simulations (IAS-6) & JARA Brain Institute I, Juelich Research Centre, Juelich, Germany

²Department of Physics, University of Oslo, Oslo, Norway

³Institut de Neurosciences de la Timone (INT), CNRS - Aix Marseille Universitée, Marseille, France

⁴Theoretical Systems Neurobiology, RWTH Aachen University, Aachen, Germany

¹Centre for Cognition and Decision Making, National Research University Higher School of Economics, Moscow, Russia

²Department of Cognitive Studies, Ecole Normale Superieure PSL* Research University, Paris, France

P90 Workflow for model building, parameter estimation and uncertainty analysis applied to calciumand G-protein dependent subcellular signaling underlying synaptic plasticity

Parul Tewatia¹*, Olivia Eriksson^{1,2}, Andrei Kramer², Joao Santos³, Alexandra Jauhiainen⁴, Kim Avrama Blackwell⁵, and Jeanette Hällgren Kotaleski^{1,3}

P91 The role of striatal feedforward inhibition in propagation of cortical oscillations

Jovana Belic^{1,2,3}[⋆], Arvind Kumar², and Jeanette Hällgren Kotaleski^{1,2,4}

P92 Electrical propagation on Cortical Connectome and Communicability

Masanori Shimono^{1,3*}, Naomichi Hatano²

P93 A cortical model for learning complex temporal structure in sensory streams

Subutai Ahmad*, Yuwei Cui, and Jeff Hawkins

Numenta, Redwood City, CA, USA

P94 Conditions for traveling waves in spiking neural networks obtained from a rigorous mapping to a neural-field model

Johanna Senk¹*, Karolína Korvasová¹, Jannis Schuecker¹, Espen Hagen^{1,2}, Tom Tetzlaff¹, Markus Diesmann^{1,3,4}, and Moritz Helias^{1,4}

P95 Temporal structure of synchrony and Unitary Events in periodically-driven balanced networks

Tobias Kuehn¹*, Michael Denker¹, Piergianluca Mana¹, Sonja Grün^{1,2}, and Moritz Helias^{1,3}

¹KTH Royal Institute of Technology, School of Computer Science and Communication, Sweden

²Stockholm University, Department of Numerical Analysis and Computer Science, Sweden

³Karolinska Institute, Department of Neuroscience, Sweden

⁴Early Clinical Biometrics, AstraZeneca AB R&D, Gothenburg, Sweden

⁵Computational and Experimental Neuroplasticity Laboratory, Krasnow Institute for Advanced Study, George Mason University, USA

¹ Science for Life Laboratory, Royal Institute of Technology, Solna, Sweden

²Department of Computational Science and Technology, Royal Institute of Technology, Stockholm, Sweden

³Bernstein Center Freiburg, University of Freiburg, Freiburg, Germany

⁴Department of Neuroscience, Karolinska Institute, Solna, Sweden

¹Osaka University, Toyonaka, Osaka, Japan

²University of Tokyo, Bunkyo, Tokyo, Japan

³Riken Brain Science Institute, Saitama, Japan

¹ Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6) and JARA BRAIN Institute I, Forschungszentrum Jülich, Jülich, Germany

²Department of Physics, University of Oslo, Oslo, Norway

³Department of Psychiatry, Psychotherapy and Psychosomatics, Medical Faculty, RWTH Aachen University, Aachen. Germany

⁴Department of Physics, Faculty 1, RWTH Aachen University, Aachen, Germany

¹ Institute of Neuroscience and Medicine (INM-6) and Institute of Advanced Simulation (IAS-6) and JARA Brain Institute I, Jülich Research Centre, Jülich, Germany

²Theoretical Systems Neurobiology, Faculty I, RWTH Aachen University, Aachen, Germany

³Department of Physics, Faculty I, RWTH Aachen University, Aachen, Germany

P96 Distributed correlations in motor cortex suggest virtually unstable linearized dynamics

David Dahmen^{1*}, Markus Diesmann^{1,2,3}, and Moritz Helias^{1,3}

P97 Transition to chaos and short-term memory in driven random neural networks

Jannis Schuecker^{1*}, Sven Goedeke¹, and Moritz Helias^{1,2}

P98 Dynamics of cell assemblies in binary neuronal networks

Christian Keup^{1,2*}, Tobias Kuehn^{1,2}, and Moritz Helias^{1,2}

P99 A dynamic mean-field approach for the largest Lyapunov exponent of random neural networks

Sven Goedeke^{1*}, Jannis Schuecker¹, and Moritz Helias^{1,2}

P100 Microdraw: Online platform for the collaborative editing of cytoarchitectonic brain atlases

Katja Heuer¹, Rembrandt Bakker^{2,3*}, Paul Tiesinga³, and Roberto Toro⁴

P101 Single-compartment models of retinal ganglion cells with different morphologies

Wei Qin¹, Alex Hadjinicolaou¹, Hamish Meffin^{2,3}, David B Grayden¹, Anthony N Burkitt¹, Michael Ibbotson^{2,3}, and Tatiana Kameneva¹*

¹ Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6) and JARA BRAIN Institute I, Jülich Research Centre, Jülich, Germany

²Department of Psychiatry, Psychotherapy and Psychosomatics, Medical Faculty, RWTH Aachen University, Aachen, Germany

³Department of Physics, Faculty 1, RWTH Aachen University, Aachen, Germany

¹Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6) and JARA BRAIN Institute I, Jülich Research Centre, Jülich, Germany

²Department of Physics, Faculty 1, RWTH Aachen University, Aachen, Germany

¹ Institute of Neuroscience and Medicine (INM-6) and Institute of Advanced Simulation (IAS-6) and JARA, Jülich Research Centre, Jülich, Germany

²Department of Physics, Faculty I, RWTH Aachen University, Aachen, Germany

¹Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6) and JARA BRAIN Institute I, Jülich Research Centre, Jülich, Germany

²Department of Physics, Faculty 1, RWTH Aachen University, Aachen, Germany

¹Dept. of Neuropsychology, Max Planck Institute for Human Cognitive and Brain Sciences, Leipzig, Germany

²Neuroinformatics dept., Donders Institute for Brain, Cognition and Behaviour, Radboud University Nijmegen, Niimegen, The Netherlands

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

⁴Applied and Theoretical neuroanatomy group, Institut Pasteur, Paris, France

¹Department of Biomedical Engineering, University of Melbourne, Victoria, Australia

²National Vision Research Institute, Australian College of Optometry, VIC, Australia

³Department of Optometry and Vision Sciences, University of Melbourne, Victoria, Australia

P102 Credibility, Replicability, Reproducibility in Simulation for research and clinical application

William W Lytton¹*, Lealem Mulugeta², Andrew Drach³, Jerry Myers⁴, Marc Horner⁵, Rajanikanth Vadigepalli⁶, Tina Morrison⁷, Marlei Walton⁸, Martin Steele⁹, and Anthony Hunt¹⁰

P103 Computational social interaction in reciprocity and empathic behavior as behavioral economics and risk tasking behavior

Nicoladie Tam*

Department of Biological Sciences, University of North Texas, Denton, TX, USA

P104 On the need for standardized real-time software technology in closed-loop neuroscience

Rodrigo Amaducci, Carlos Muñiz, Manuel Reyes-Sánchez, Francisco B Rodríguez, and Pablo Varona*

Grupo de Neurocomputación Biológica, Dpto. de Ingeniería Informática, Escuela Politécnica Superior, Universidad Autónoma de Madrid, Spain

¹Dept. Of Physiology, SUNY Downstate Medical Center, Brooklyn, NY, USA

²InSilico Labs LLC, Houston, TX, USA

³Institute for Computational Engineering and Sciences, University of Texas at Austin, Austin, TX, USA

⁴John H Glenn Research Center, NASA, Houston, TX, USA

⁵ANSYS, Inc. Canonsburg, PA, USA

⁶Thomas Jefferson University, Philadelphia, PA, USA

⁷U.S. Food and Drug Administration, Washington DC, USA

⁸KBRWyle, El Segundo, CA, USA

⁹Kennedy Space Center, NASA, Houston, TX, USA

¹⁰Bioengineering and Therapeutic Sciences, University of California, San Francisco, CA, USA

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P105 Discovering Connectivity Changes with rescaled Energy-Based Models

Joseph Cronin*, Matthias Hennig

Institute for Adaptive and Neural Computation, School of Informatics, The University of Edinburgh, United Kingdom

P106 Fitting and analysis pipeline to build data-driven models of tonic and burst firing in thalamic neurons

Elisabetta lavarone^{1*}, Christian O Reilly¹, Jane Yi², Ying Shi², Bas-Jan Zandt¹, Werner van Geit³, Christian Rössert³, Henry Markram^{2,3}, and Sean Hill¹

P107 A data-driven pipeline for digital reconstruction of somatosensory thalamic microcircuitry

Christian O Reilly^{1*}, Elisabetta lavarone¹, Jane Yi², Ying Shi², Rodrigo Perin², Huanxiang Lu¹, Bas-Jan Zandt¹, Henry Markram², and Sean Hill¹

P108 Sensitivity analysis of somatic and dendritic features, parameters, and fitting error of biophysically detailed neuron models

Bas-Jan Zandt¹*, Elisabetta lavarone¹, Alexander Bryson², Werner van Geit¹, Christian O Reilly¹, Christian Rössert¹, and Sean Hill¹

P109 Music gone crazy: neural oscillators to the rescue?

Michal Hadrava^{1,2,3*}, Jaroslav Hlinka^{2,3}

P110 Bistability generates highly irregular spike trains with weakly fluctuated inputs

Ryosuke Hosaka*

Department of Applied Mathematics, Fukuoka University, Fukuoka, Japan

P111 Big bang bifurcations structure the parameter space of a two-cell inhibitory network with synaptic depression.

Mark Olenik¹*, Conor Houghton²

¹Laboratory for the Neural Basis of Brain States, Blue Brain Project, EPFL, Geneva, Switzerland

²Laboratory of Neural Microcircuity, Brain Mind Institute, EPFL, Lausanne, Switzerland

³Blue Brain Project, EPFL, Geneva, Switzerland

¹Laboratory for the Neural Basis of Brain States, Blue Brain Project, EPFL, Geneva, Switzerland

²Laboratory of Neural Microcircuity, Brain Mind Institute, EPFL, Lausanne, Switzerland

¹Blue Brain Project, École Polytechnique Fédérale de Lausanne, Geneva, Switzerland

²Florey Institute of Neuroscience & Mental Health, University of Melbourne, Melbourne, Vic, Australia

¹Department of Cybernetics, Faculty of Electrical Engineering, Czech Technical University in Prague, Prague, Czech Republic

²Department of Nonlinear Dynamics and Complex Systems, Institute of Computer Science, The Czech Academy of Sciences, Prague, Czech Republic

³National Institute of Mental Health, Klecany, Czech Republic

¹School of Biological Sciences, University of Bristol, Bristol, UK

²Department of Computer Science, University of Bristol, Bristol, UK

P112 Morphology and balance influences Dendritic mosaic formation.

Nicolangelo lannella1*, Thomas Launey2

P113 Comparison between extracellular and intracellular stimulation

Tatiana Kameneva^{1*}, Rebecca Kotsakidis², Hamish Meffin^{2,3}, and Michael Ibbotson^{2,3}

P114 X106 Bifurcations in a temperature-dependent neural mass model reveal heterogeneous effect of focal cooling on epileptic discharges

Jaymar Soriano^{1,2*}, Takatomi Kubo¹, Takao Inoue³, Hiroyuki Kida³, Toshitaka Yamakwa⁴, Michiyasu Suzuki³, and Kazushi Ikeda¹

P115 Robust Transmission of Rate Coding in the Inhibitory Purkinje Cell to Cerebellar Nuclei Pathway in Awake Mice

Samira Abbasi², Amber Hudson¹, Detlef Heck³, and Dieter Jaeger¹★

P116 A method to generate realistic artificial spike trains as inputs to biophysical neuron models: cerebellar mossy fibers as a case study

Joel Lee^{1*}, Samira Abbasi², Amber Hudson¹, Detlef H Heck³, and Dieter Jaeger¹

P117 Serotonergic fiber densities may emerge from random walks

Skirmantas Janusonis*

Department of Psychological and Brain Sciences, University of California, Santa Barbara, CA, USA

P118 A taxonomy of seizures based on dynamics

Maria Luisa Saggio^{1*}, Andreas Spiegler¹, William Stacey², Christophe Bernard¹, and Viktor Jirsa¹

P119 Computational model for diffusion-induced bursting of biophysically realistic HH-type neuron: mathematical characterization

Davide Lillo*, Christophe Bernard, and Viktor Jirsa

Institut de Neurosciences des Systèmes, Aix Marseille Univ, Marseille, France

¹ School of Mathematical Sciences, University of Nottingham, Nottingham, UK

²Lab for Synaptic Molecules of Memory Persistence, RIKEN, Brain Science Institute, Saitama, Japan

¹Department of Biomedical Engineering, University of Melbourne, Victoria, Australia

²National Vision Research Institute, Australian College of Optometry, VIC, Australia

³Department of Optometry and Vision Sciences, University of Melbourne, Victoria, Australia

¹Nara Institute of Science and Technology, Nara, Japan

²University of the Philippines - Diliman, Quezon City, Philippines

³Yamaguchi University, Ube, Japan

⁴Kumamoto University, Kumamoto, Japan

¹Department Biology, Emory University, Atlanta, GA, USA

²Department of Biomedical Engineering, Hamedan University of Technology, Hamedan, Iran

³Department of Anatomy and Neurobiology, UT Health Science Center, Memphis, TN, USA

¹Department of Biology, Emory University, Atlanta, GA, USA

²Department of Biomedical Engineering, Hamedan University of Technology, Hamedan, Iran

³Department of Anatomy and Neurobiology, University of Tennessee Health Science Center, Memphis, TN, USA

¹INSERM UMR 1106 Institut de Neurosciences des Systèmes - Aix-Marseille Université, Marseille, France

²Department of Neurology, Department of Biomedical Engineering, University of Michigan, Ann Arbor, MI, USA

P120 Phase-lags in large scale brain synchronization

Spase Petkoski*, Viktor Jirsa

INSERM UMR 1106 Institut de Neurosciences des Systemes - Aix-Marseille Universite, Marseille, France

P121 Network dynamics after focal stimulation in a connectome-based network model of the mouse brain

Andreas Spiegler*, Viktor Jirsa

Institut de la Santé et de la Recherche Médical, Institut de Neurosciences des Systèmes UMR_S 1106, Aix Marseille Université, 13005 Marseille, France

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P123 Analysing the impact of sodium channels in Alzheimer's disease using a computational model

Seyed Ali Sadegh Zadeh*, Chandra Kambhampati

Department of Computer Science, University of Hull, Hull, UK

P124 Models of brain design: is physics more important than evolutionary optimization?

Jan Karbowski*

University of Warsaw, Warsaw, Poland

P125 Perceptual Attractors and Neural Confusions in Phoneme Manifolds

Zeynep Gokcen Kaya¹*, Yair Lakretz², and Alessandro Treves^{1,3}

P126 Network analysis of task-oriented neuroimaging data via multivariate information-theoretic measures

Lily W Li¹, Joseph Lizier², Paula Sanz-Leon^{1,3}, and Cliff C Kerr^{1,3}*

P127 Optimal localist and distributed coding of spatiotemporal spike patterns through STDP and coincidence-detection

Timothee Masquelier¹[⋆], Saeed Reza Kheradpisheh²

P128 Muscle force potentiation induced by active dendrites of spinal motoneuron during locomotorlike movement

Hojeong Kim*

Convergence Research Institute, DGIST, Daegu, Korea

¹Cardiff University Brain Research Imaging Centre, Cardiff University, Cardiff, UK

²Neuroscience and Mental Health Research Institute, Cardiff University, Cardiff, UK

¹Cognitive Neuroscience Sector, SISSA, Trieste, 34136, ITALY

²Sagol School of Neuroscience, Tel Aviv University, Tel Aviv, ISRAEL

³Kavli Institute, Norwegian University of Science and Technology, Trondheim, NORWAY

¹School of Physics, University of Sydney, NSW, Australia

²School of Civil Engineering, University of Sydney, NSW, Australia

³Centre for Integrative Brain Function, University of Sydney, NSW, Australia

¹CERCO UMR 5549, CNRS – Université de Toulouse, France

²Dept. of Computer Sc., School of Mathematics, Statistics, and Computer Science, Univ. of Tehran, Tehran, Iran

P129 X180 Perceptual Mechanics of the Brian and the Free Energy Principle

Chang Sub Kim*

Department of Physics, Chonnam National University, Gwangju, Republic of Korea

P130 Influence of handedness on the response inhibition in Stroop task: ERP study

Julia Marakshina^{1,2}, Alexander Vartanov¹, Anastasia Neklyudova^{1*}, Stanislav Kozlovskiy¹, and Andrey Kiselnikov¹

P131 Contribution of short-term plasticity of the bipolar-ganglion synapse to the activity both in the normal and the degenerating rd1 retina

Kanako Taniguchi¹, Katsunori Kitano²*

P132 Differential connectomics of the rat thalamus

Oliver Schmitt*, Felix Lessmann, Sebastian Schwanke, Peter Eipert, Jennifer Meinhardt, Julia Beier, Kanar Kadir, Adrian Karnitzki, Linda Sellner, Ann-Christin Klünker, Lena Kuch, Frauke Ruß, Jörg Jenssen, and Andreas Wree

Department of Anatomy, University of Rostock, Rostock, Germany

P133 X351 Nonuniform neural field modeling of nonlinear dynamics spreading

Sanz-Leon Paula^{1,2*}, Stuart Knock^{1,2}

P134 Excitatory-to-inhibitory plasticity for sequence learning

Shih-Cheng Chien*, Burkhard Maess, and Thomas R. Knösche

Max Planck Institute for Human Cognitive and Brain Sciences, Leipzig, Germany

P135 Experimental and modelling evidence for coaxial conduction in myelinated axons

Charles Cohen^{1,2*}, Marko A Popovic¹, Jan Klooster¹, and Maarten Hp Kole^{1,2}

P136 GIMBL-Vis: A GUI-Based Interactive Multidimensional Visualization Toolbox for Matlab

Erik Roberts^{1*}, Nancy Kopell²

P137 Neural relativity principle

Daniel Kepple¹, Hamza Giaffar¹, Dmitry Rinberg², and Alexei Koulakov¹*

¹Lomonosov Moscow State University, Moscow, Russia

²Psychological Institute of Russian Academy of Education, Moscow, Russia

¹ Graduate School of Information Science and Engineering, Ritsumeikan University, Kusatsu, Japan

²Department of Human and Computer Intelligence, Ritsumeikan University, Kusatsu, Japan

¹School of Physics, University of Sydney, New South Wales, Australia

²Center for Integrative Brain Function, University of Sydney, New South Wales, Australia

¹ Axonal Signalling Group, Netherlands Institute for Neuroscience, Royal Netherlands Academy for Arts and Sciences (KNAW), Amsterdam, The Netherlands

²Cell Biology, Faculty of Science, Utrecht University, Utrecht, The Netherlands

¹Department of Biomedical Engineering, Boston University, Boston, MA, USA

²Department of Mathematics and Statistics, Boston University, Boston, MA, USA

¹Cold Spring Harbor Laboratory, Cold Spring Harbor, NY, USA

²New York University, New York, NY, USA

P138 Disrupted cerebrum-cerebellum network in Schizophrenia revealed by network-based statist and graph theory

Caroline Garcia Forlim¹*, Leonie Klock^{1,2}, Johanna Bächle², Laura Stoll², Patrick Giemsa², Marie Fuchs², Nikola Schoofs², Christiane Montag², Jurgen Gallinat¹, and Simone Kuhn¹

P139 Computational behavioral analysis of acute psychosocial trauma reveals gradually increasing stress reactions in adult mice

Ray Lee^{1,2*}, Greg Stephens^{2,3}, and Bernd Kuhn¹

P140 Relationship between population sparse coding and short-term synaptic plasticity

Luiz Tauffer^{1,2*}, Philippe Isope³, and Arvind Kumar²

P141 The contribution of topology for inclusion of feedforward network and biased synaptic strength to the long-term memory effect in a cortical microcircuit

Katsuma Inoue*, Yoshiyuki Ohmura, Shogo Yonekura, and Yasuo Kuniyoshi

Department of Mechano-Informatics, University of Tokyo, Bunkyo, Tokyo, JPN

P142 Propagation of spatio-temporally complex spike pattern in feedforward network of the barrel cortex: in vitro multi-electrode array and in silico neural network model study

Hyun Jae Jang*, Jeehyun Kwag

Dept. of Brain and Cognitive Engineering, Korea University, Seoul, Korea

P143 Computational Geometry for the Simulation of Neural Circuits with Population Density Techniques

de Kamps Marc*, Yi Ming Lai

School of Computing, University of Leeds, Leeds, UK

P144 Neuron classification in the Stomatogastric ganglion using voltage-sensitive dye imaging and signal processing tools

Filipa Dos Santos*, Kp Lam, and Peter Andras

School of Computing and Mathematics, Keele University, Newcastle-under-Lyme, ST5 5BG, UK

P145 Reconstructed attractors from optogenetics experiments

Sorinel A Oprisan¹*, Julia Imperatore¹, Jessica Helms¹, Tamas Tompa², and Antonieta Lavin²

¹Clinic and Policlinic for Psychiatry and Psychotherapy, University Medical Center Hamburg-Eppendorf, Hamburg, Germany

²Charité University Medicine and St. Hedwig-Krankenhaus, Department of Psychiatry and Psychotherapy, Berlin, Germany

¹Optical Neuroimaging Unit, Okinawa Institute of Science and Technology Graduate University (OIST), Okinawa, Japan

²Biological Physics Theory Unit, Okinawa Institute of Science and Technolgy Graduate University (OIST), Okinawa, Japan

³Department of Physics and Astronomy, Vrije Universiteit Amsterdam, Amsterdam, The Netherlands

¹Bernstein Center Freiburg, University of Freiburg, Germany

²School of Computer Science and Communication, KTH Royal Institute of Technology, Stockholm, Sweden

³CNRS, University of Strasbourg, Strasbourg, France

¹Department of Physics and Astronomy, College of Charleston, Charleston, SC, USA

²Department of Neuroscience, Medical University of South Carolina, Charleston, SC, USA

P146 A Simple Parameter Landscape for Optimisation of Conductance-Based Models of Neurons

Felicity H Inkpen^{1*}, Michael Ashby¹, and Nathan Lepora²

P147 Neural Synchronization Through Electric Field Effects

Aaron Shifman^{1,2,3*}, John Lewis^{1,2,3}

P148 How Spines Cross-talk: Compartmental Model of Heterosynaptic Plasticity

Zhong Zhang¹, Yeqian Feng¹, Christian Tetzlaff³, Tomas Kulvicius⁴, and Yinyun Li²*

P149 Determination of the spike-train power spectrum statistics in modular networks with mixtures of different excitatory and inhibitory populations

Rodrigo Pena¹, Davide Bernardi^{2,3}, Antonio C Roque^{1*}, and Benjamin Lindner^{2,3}

P150 Optimal detection of single-cell stimulation in large random networks of integrate-and-fire neurons

Davide Bernardi^{1,2*}, Benjamin Lindner^{1,2}

P151 Self-consistent power spectra from an iterative scheme for recurrent heterogeneous networks

Sebastian Vellmer^{1,2*}, Benjamin Lindner^{1,2}

P152 Astrocyte-modulated synaptic plasticity in sensory cortex: A computational study

Ausra Saudargiene^{1,2}, Tiina Manninen³, Riikka Havela³, and Marja-Leena Linne³*

P153 Up and down states statistics for Gamma oscillations

Arthur Powanwe*, Andre Longtin

Department of Physics and Centre for Neural Dynamics, University of Ottawa, Ottawa, Canada

¹School of Physiology, Pharmacology and Neuroscience, University of Bristol, Bristol, UK

²Department of Engineering Mathematics, University of Bristol, Bristol, UK

¹Department of Biology, University of Ottawa, Ottawa, Ontario, Canada

²Center for Neural Dynamics, University of Ottawa, Ottawa, Ontario, Canada

³Ottawa Brain and Mind Research Institute, University of Ottawa, Ottawa, Ontario, Canada

¹Department of Management Science, Beijing Normal University, Beijing, China

²School of Systems Science, Beijing Normal University, Beijing, China

³Institute of Physics-Biophysics, Georg-August-University, Goettingen, Germany

⁴Maersk Mc-Kinney Moller Institute, University of Southern Denmark, Odense, Denmark

¹Department of Physics, University of Sao Paulo, Ribeirao Preto, Sao Paulo, Brazil

²Theory of Complex Systems and Neurophysics, Bernstein Center for Computational Neuroscience, Berlin, Germany

³Department of Physics, Humboldt University of Berlin, Berlin, Germany

¹Bernstein Center for Computational Neuroscience - Berlin, Berlin, Germany

²Department of Physics, Humboldt University, Berlin, Germany

¹Theory of Complex Systems and Neurophysics, Bernstein Center for Computational Neuroscience, Berlin, Germany

²Department of Physics, Humboldt Universität zu Berlin, Berlin, Germany

¹Neuroscience Institute, Lithuanian University of Health Sciences, Kaunas, Lithuania

²Department of Informatics, Vytautas Magnus University, Kaunas, Lithuania

³Computational Neuroscience Group, BioMediTech Institute and Faculty of Biomedical Sciences and Engineering, Tampere University of Technology, Tampere, Finland

P154 EDLUT: a real-time spiking neural network simulator for embodiment experiments

Francisco Naveros¹*, Jesús Garrido¹, Richard Carrillo¹, Eduardo Ros¹, and Niceto Luque^{2,3,4}

P155 Embedded Ensemble Encoding — a hypothesis for reconciling cortical coding strategies

Joe Graham¹, Salvador Dura-Bernal^{1*}, Sergio Angulo¹, Samuel Neymotin^{1,2}, Srdjan Antic³, and William W Lytton^{1,4}

P156 Multiscale modeling of ischemic stroke with the NEURON reaction-diffusion module

Adam Newton^{1,2*}, Alexandra Seidenstein^{2,3}, Robert A McDougal¹, and William W Lytton^{2,4}

P157 Accelerating NEURON reaction-diffusion simulations

Robert A McDougal^{1*}, William W Lytton^{2,3}

P158 Computation of invariant objects in the analysis of periodically forced neural oscillators

Alberto Perez-Cervera*, Gemma Huguet, and Tere M-Seara

Departament de Matemàtica Aplicada, Universitat Politècnica de Catalunya, Barcelona, Spain

P159 Computational model of spatio-temporal coding in CA3 with speed-dependent theta oscillation

Caroline Haimerl^{1,2}, David Angulo-Garcia^{1,3*}, Alessandro Torcini^{1,3,4}, Rosa Cossart¹, and Arnaud Malvache¹

P160 The effect of progressive degradation of connectivity between brain areas on the brain network structure

Kaoutar Skiker*, Mounir Maouene

Department of mathematics and computer science, ENSAT, Abdelmalek Essaadi's University, Tangier, Morocco

¹Department of Computer Architecture and Technology, University of Granada (CITIC), Granada, Spain

²INSERM, U968, Paris, France

³Sorbonne Universités, UPMC University Paris 06, UMR_S 968, Institut de la Vision, Paris, France

⁴CNRS, UMR 7210, Paris, France

¹Department of Physiology and Pharmacology, SUNY Downstate Medical Center, Brooklyn, NY, USA

²Department of Neuroscience, Brown University, Providence, RI, USA

³Institute for Systems Genomics, University of Connecticut Health Center, Farmington, CT, USA

⁴Department of Neurology, Kings County Hospital, Brooklyn NY, USA

¹Department of Neuroscience, Yale University, New Haven, CT, USA

²Department Physiology & Pharmacology, SUNY Downstate, Brooklyn, NY, USA

³NYU School of Engineering, 6 MetroTech Center, Brooklyn, NY, USA

⁴Kings County Hospital Center, Brooklyn, NY, USA

¹Neuroscience, Yale University, New Haven, CT, USA

²Physiology & Pharmacology, SUNY Downstate Medical Center, Brooklyn, NY, USA

³Kings County Hospital, Brooklyn, NY, USA

¹Institut de Neurobiologie de la Méditerrannée (INMED), INSERM, UMR901, Aix-Marseille Univ, Marseille, France

²Center of Neural Science, New York University, NYC, NY, USA

³Aix-Marseille Univ, INSERM, INS, Inst Neurosci Syst, Marseille, France

⁴Laboratoire de Physique Théorique et Modélisation, CNRS UMR 8089, Université de Cergy-Pontoise, Cergy-Pontoise Cedex, France

P161 A network architecture for comparing the behavior of a neurocomputational model of rewardbased learning with human

Gianmarco Ragognetti^{1*}, Letizia Lorusso², Andrea Viggiano², and Angelo Marcelli¹

P162 Distributed plasticity in the cerebellum: how do cerebellar cortex and nuclei plasticity cooperate for learning?

Rosa Senatore, Antonio Parziale, and Angelo Marcelli*

Laboratory of Natural Computation, Department of Information and Electrical Engineering and Applied Mathematics, University of Salerno, Italy

P163 Ising Model with conserved magnetization on the Human Connectome: implications on the relation structure-function in wakefulness and anesthesia

Sebastiano Stramaglia¹, Mario Pellicoro¹, Leonardo Angelini¹, Enrico Amico^{2,3}, Hannelore Aerts², Jesus Cortes⁴, Steven Laureys³, and Daniele Marinazzo²*

P164 Multiscale Granger causality analysis by à trous wavelet transform

Sebastiano Stramaglia^{1*}, lege Bassez², Luca Faes³, and Daniele Marinazzo²

P165 New (spectral) dynamic causal modeling scheme improves effective connectivity estimation within resting state networks in longitudinal data

Hannes Almgren^{1*}, Frederik van de Steen¹, Adeel Razi^{2,3}, and Daniele Marinazzo¹

P166 Effective connectivity modulations of win-and loss feedback: A dynamic causal modeling study of the human connectome gambling task.

Frederik van de Steen¹*, Ruth Krebs², and Daniele Marinazzo¹

P167 Modeling global brain dynamics in brain tumor patients using the Virtual Brain

Hannelore Aerts*, Daniele Marinazzo

Department of Data Analysis, Ghent University, Belgium

¹Laboratory of Natural Computation, Department of Information and Electrical Engineering and Applied Mathematics, University of Salerno, Italy

²Department of Medicine, University of Salerno, 84083 Lancusi (SA), Italy

¹ Dipartimento di Fisica, Università degli Studi Aldo Moro, Bari, and INFN, Sezione di Bari, Italy

²Data Analysis Department, Ghent University, Ghent, Belgium

³Coma Science Group, University of Liège, Belgium

⁴Cruces Hospital and Ikerbasque Research Center, Bilbao, Spain

¹Dipartimento di Fisica, Università degli Studi Aldo Moro, Bari, and INFN, Sezione di Bari, Italy

²Data Analysis Department, Ghent University, Ghent, Belgium

³BIOtech, Dept. of Industrial Engineering, University of Trento, and IRCS-PAT FBK, Trento, Italy

¹Department of Data Analysis, Ghent University, Ghent, Belgium

²The Wellcome Trust Centre for Neuroimaging, University College London, London, UK

³ Department of Electronic Engineering, NED University of Engineering and Technology, Karachi, Pakistan

¹Department of data analysis, Ghent University, Ghent, Belgium

²Department of experimental psychology, Ghent University, Ghent, Belgium

P168 Representation of Neuronal Morphologies

Lida Kanari^{1*}, Pawel Dlotko², Martina Scolamiero³, Ran Levi⁴, Julian Shillcock¹, Christiaan de Kock⁵, Kathryn Hess³, and Henry Markram¹

P169 X034 Firing Rate Heterogeneity and Consequences for Stimulus Estimation in the Electrosensory System

Cheng Ly1*, Gary Marsat2

P170 KnowledgeSpace: a community encyclopedia linking brain research concepts to data, models and literature

Tom Gillespie², Willy Wong², Malin Sandstroem¹*, Mathew Abrams¹, Jeffrey S. Grethe², and Maryann Martone³

P171 Evaluating the computational capacity of a cerebellum model

Robin de Gernier^{1*}, Sergio Solinas², Christian Rössert³, Marc Haelterman¹, and Serge Massar¹

P172 Complexity of cortical connectivity promotes self-organized criticality

Valentina Pasquale^{1*}, Vito Paolo Pastore², Sergio Martinoia², and Paolo Massobrio²

P173 Attractor dynamics of cortical assemblies underlying brain awakening from deep anesthesia.

Cristiano Capone^{1,2⋆}, Núria Tort-Colet³, Maria V. Sanchez-Vives^{3,4}, and Maurizio Mattia¹

¹Blue Brain Project, École polytechnique fédérale de Lausanne, Switzerland

²Departement of Mathematics, Swansea University, UK

³Laboratory for Topology and Neuroscience at the Brain Mind Institute, École polytechnique fédérale de Lausanne, Switzerland

⁴Institute of Mathematics, University of Aberdeen, UK

⁵Department of Integrative Neurophysiology, Center for Neurogenomics and Cognitive Research, VU Universiteit Amsterdam, The Netherlands

¹Department of Statistical Sciences and Operations Research, Virginia Commonwealth University, Richmond, VA, USA

²Biology Department, West Virginia University, Morgantown, WV, USA

¹INCF Secretariat, Karolinska Institute, Stockholm, Sweden

²Center for Research in Biological Systems, UCSD, La Jolla, CA, USA

³Neurosciences, UCSD, La Jolla, CA, USA

¹École polytechnique de Bruxelles, Université libre de Bruxelles, Brussels, Belgium

²Department of Biomedical Science, University of Sassari, Sassari, Italia

³Blue Brain Project, École polytechnique fédérale de Lausanne, Geneva, Switzerland

¹Neuroscience and Brain Technologies Department, Istituto Italiano di Tecnologia (IIT), Genova, ITALY

²Department of Informatics, Bioengineering, Robotics, System Engineering (DIBRIS), University of Genova, Genova, ITALY

¹ Istituto Superiore di Sanità (ISS), Rome, Italy

²PhD Program in Physics, Sapienza University, Rome, Italy

³Institut d'Investigacions Biomèdiques August Pi i Sunyer (IDIBAPS), Barcelona, Spain

⁴Institució Catalana de Recerca i Estudis Avançats (ICREA), Barcelona, Spain

P174 Are receptive fields in visual cortex quantitatively consistent with efficient coding?

Ali Almasi^{1,2}, Shaun Cloherty⁴, David B Grayden², Yan Wong^{3,4}, Michael Ibbotson^{1,5}, and Hamish Meffin^{1,5}*

P175 Cholinergic Modulation of DG-CA3 microcircuit dynamics and function

Luke Prince^{1*}, Krasimira Tsaneva-Atanasova^{2,3}, and Jack Mellor¹

P176 Subthalamic nucleus low frequency fluctuations carry information about future economic decisions in parkinsonian gamblers

Alberto Mazzoni^{1*}, Manuela Rosa², Jacopo Carpaneto¹, Luigi Romito³, Alberto Priori^{2,4}, and Silvestro Micera^{1,5}

P177 Data-driven computational modeling of CA1 hippocampal principal cells and interneurons

Rosanna Migliore^{1*}, Carmen Alina Lupascu¹, Francesco Franchina¹, Luca Leonardo Bologna¹, Armando Romani², Christian Rossert², Sara Saray³, Jean-Denis Courcol², Werner van Geit², Szabolcz Kali³, Alex Thomson⁴, Audrey Mercer⁴, Sigrun Lange^{4,5}, Joanne Falck⁴, Eilif Muller², Felix Schurmann², and Michele Migliore¹

P178 The interplay between basal ganglia and cerebellum in motor adaptation

Dmitrii Todorov, Robert Capps, William Barnett, and Yaroslav Molkov*

Department of Mathematics and Statistics, Georgia State University, Atlanta, GA, USA

P179 Microscopic and macroscopic dynamics of neural populations with delays

Federico Devalle^{1,2}*, Diego Pazó³, and Ernest Montbrió¹

¹National Vision Research Institute, Australian College of Optometry, Australia

²NeuroEngineering Laboratory, Dept. Biomedical Eng., University of Melbourne, Australia

³Dept. of Physiology, Monash University, Australia

⁴Dept. of Electrical & Computer Systems Eng., Monash University, Australia

⁵ARC Centre of Excellence for Integrative Brain Function, University of Melbourne, Australia

¹Centre for Synaptic Plasticity, School of Physiology, Pharmacology, and Neuroscience, University of Bristol, Bristol, UK

²Department of Mathematic, College of Engineering, Mathematics and Physical Sciences, University of Exeter, Exeter, UK

³EPRSC Centre for Predictive Modelling in Healthcare, University of Exeter, Exeter, UK

¹ Translational Neural Engineering, The Biorobotics Institute, Scuola Superiore Sant'Anna, Pontedera, Italy

²Clinical Center for Neurostimulation, Neurotechnology and Movement Disorders Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Milan, Italy

³Movement Disorders Department, Neurological Institute Carlo Besta, Milan, Italy

⁴Department of Health Sciences, University of Milan & ASST Santi Paolo e Carlo, Milan, Italy

⁵Bertarelli Foundation Chair in Translational NeuroEngineering, Institute of Bioengineering and Center for Neuroprosthetics, Ecole Polytechnique Federale De Lausanne, Lausanne, Switzerland

¹Institute of Biophysics, National Research Council, Palermo, Italy

²Blue Brain Project, École Polytechnique Fédérale de Lausanne Biotech Campus, Geneva, Switzerland

³Institute of Experimental Medicine, Hungarian Academy of Sciences, Budapest, Hungary

⁴University College London, United Kingdom

⁵University of Westminster, London, United Kingdom

¹Center for Brain and Cognition, Universitat Pompeu Fabra, Barcelona, Spain

²Department of Physics, Lancaster University, Lancaster, UK

³Instituto de Fisica de Cantabria (IFCA). CSIC-Universidad de Cantabria, Santander, Spain

P180 Motivation signal in anterior cingulate cortex during economic decisions

Gabriela Mochol¹*, Habiba Azab², Benjamin Hayden², and Rubén Moreno - Bote¹

P181 A simple computational model of altered neuromodulation in cortico-basal ganglia dynamics underlying bipolar disorder

Pragathi Priyadharsini Balasubramani¹, Srinivasa Chakravarthy², and Vignayanandam Muddapu²*

P182 Theta/alpha coordination of pre-motor and parietal networks during free behavior in rats

Medorian Gheorghiu¹*, Bartul Mimica², Jonathan Withlock², and Raul C Muresan¹

P183 Information theoretic approach towards identifying changes in cellular-level functional connectivity and synchrony across animal models of schizophrenia

Jennifer Zick^{1,2*}, Kelsey Schultz⁴, Rachael Blackman^{1,2,3}, Matthew Chafee^{1,3}, and Theoden Netoff^{1,4}

P184 Neural Suppression with Deep Brain Stimulation using a Linear Quadratic Regulator

Nicholas Roberts^{1*}, Vivek Nagaraj², Andrew Lamperski³, and Theoden Netoff¹

P185 Reinforcement learning for phasic disruption of pathological oscillations in a computational model of Parkinson's disease

Logan Grado^{1*}, Matthew Johnson^{1,2}, and Theoden Netoff¹

P186 Metrics for detection of delayed and directed coupling

David Darrow¹*, Theoden Netoff²

¹Center for Brain and Cognition and Department of Information and Communications Technologies, University Pompeu Fabra, Barcelona, Spain

²Department of Brain and Cognitive Sciences and Center for Visual Sciences, University of Rochester, Rochester, NY, USA

¹Brain and Cognitive Sciences, University of Rochester, Rochester, New York, USA

²Bhupat and Jyoti Mehta School of Biosciences, Department of Biotechnology, IIT- Madras, Chennai, TN, India

¹Romanian Institute of Science and Technology, Cluj-Napoca, Cluj, Romania

²Centre for Neural Computation, Kavli Institute for Systems Neuroscience, Trondheim, Norway

¹Graduate Program in Neuroscience, University of Minnesota, Minneapolis, MN, USA

²Medical Scientist Training Program (MD/PhD), University of Minnesota, Minneapolis, MN, USA

³Brain Sciences Center, VA Medical Center, Minneapolis, MN, USA

⁴Department of Biomedical Engineering, University of Minnesota, Minneapolis, MN, USA

¹Department of Biomedical Engineering, University of Minnesota, Minneapolis, MN, USA

²Graduate Program in Neuroscience, University of Minnesota, Minneapolis, MN, USA

³Department of Electrical and Computer Engineering, University of Minnesota, Minneapolis, MN, USA

¹Department of Biomedical Engineering, University of Minnesota, Minneapolis, MN, United States

²Institute for Translational Neuroscience, University of Minnesota, Minneapolis, MN, United States

¹Department of Neurosurgery, University of Minnesota, Minneapolis, MN, USA

²Department of Biomedical Engineering, University of Minnesota, Minneapolis, MN, USA

P187 Insurgence of network bursting events in formed neuronal culture networks: a computational approach

Davide Lonardoni^{1*}, Hayder Amin¹, Stefano Di Marco², Alessandro Maccione¹, Luca Berdondini¹, and Thierry Nieus^{1,3}

P188 Brian2GeNN: Free GPU Acceleration for Brian 2 Users

Marcel Stimberg¹, Dan F M Goodman², and Thomas Nowotny³*

P189 Spike counts in the visual cortex consistently encode both stimuli and behavioral choices in a change-detection task.

Veronika Koren^{1,2}*, Valentin Dragoi³, and Klaus Obermayer^{1,2}

P190 Local topology of connectome stabilizes critical points in mean field model

Samy Castro^{1,2}*, Mariano Fernandez³, Wael El-Deredy⁴, and Patricio Orio^{1,5}

P191 How chaos in neural oscillators determine network behavior

Kesheng Xu¹, Jean Paul Maidana¹, and Patricio Orio^{1,2}*

P192 STEPS 3: integrating stochastic molecular and electrophysiological neuron models in parallel simulation

Weiliang Chen^{1*}, Iain Hepburn¹, Francesco Casalegno², Adrien Devresse², Aleksandr Ovcharenko², Fernando Pereira², Fabien Delalondre², and Erik De Schutter¹

P193 A conductance-based model of cerebellar molecular layer interneurons

Peter Bratby, Erik De Schutter*

Okinawa Institute of Science and Technology Graduate University, Okinawa, Japan

¹Neuroscience and Brain Technology Department, Fondazione Istituto Italiano di Tecnologia, Genova, Italy

²Scienze cliniche applicate e biotecnologiche, Università dell'Aquila, L'Aquila, Italy

³Dept. of Biomedical and Clinical Sciences 'Luigi Sacco', University of Milan, Italy

¹ Sorbonne Universités, UPMC Univ Paris 06, INSERM, CNRS, Institut de la Vision, Paris, France

²Department of Electrical and Electronic Engineering, Imperial College, London, UK

³School of Engineering and Informatics, University of Sussex, Brighton, UK

¹Neural Information Processing Group, Institute of Software Engineering and Theoretical Computer Science, Technische Universität Berlin, Berlin, Germany

²Bernstein Center for Computational Neuroscience Berlin, Germany

³Department of Neurobiology and Anatomy, University of Texas Medical School, Houston, Texas, US

¹Centro Interdisciplinario de Neurociencia de Valparaíso, Universidad de Valparaíso, Valparaíso, 2360102, Chile ²Programa de Doctorado en Ciencias, mención en Neurociencia, Facultad de Ciencias, Universidad de Valparaíso, Valparaíso, 2360102, Chile

³Laboratorio de Electrónica Industrial, Control e Instrumentación, Universidad Nacional de La Plata, La Plata, Argentina

⁴Escuela de Ingeniería Biomédica, Universidad de Valparaíso, 2362905, Valparaíso, Chile

⁵ Instituto de Neurociencia, Universidad de Valparaíso, Facultad de Ciencias, Universidad de Valparaíso, Valparaíso, 2360102. Chile

¹ Centro Interdisciplinario de Neurociencia de Valparaíso, Universidad de Valparaíso, Valparaíso, Chile

² Facultad de Ciencias, Instituto de Neurociencia, Universidad de Valparaíso, Valparaíso, Chile

¹ Computational Neuroscience Unit, Okinawa Institute of Science and Technology Graduate University, Japan

²Blue Brain Project, École Polytechnique Fédérale de Lausanne, Switzerland

P194 An Ultrasensitive ON/OFF Switch Mechanism Controls the Early Phase of Cerebellar Plasticity

Andrew Gallimore*, Erik De Schutter

Computational Neuroscience Unit, Okinawa Institute of Science and Technology Graduate University, Okinawa, Japan

P195 The use of hardware accelerators in the STochastic Engine for Pathway Simulation (STEPS)

Guido Klingbeil*, Erik De Schutter

Computational Neuroscience Unit, Okinawa Institute of Science and Technology, Okinawa, Japan

P196 A model of CaMKII sensitivity to the frequency of Ca2+ oscillations in Cerebellar Long Term Depression

Criseida Zamora*, Erik De Schutter

Computational Neuroscience Unit, Okinawa Institute of Science and Technology Graduate University, Okinawa, Japan

P197 Exploring the response to climbing fiber input in Purkinje neurons by a new experimental data based model

Yunliang Zang*, Erik De Schutter

Computational Neuroscience Unit, Okinawa Institute of Science and Technology Graduate University, Okinawa, Japan

P198 Effects of network topology perturbations on memory capacity in a hippocampal place cell model

Patrick Crotty*, Eric Palmerduca

Department of Physics and Astronomy, Colgate University, Hamilton, NY, USA

P199 A NEST-simulated cerebellar spiking neural network driving motor learning

Alberto Antonietti^{1*}, Claudia Casellato¹, Csaba Erö², Egidio Dangelo³, Marc-Oliver Gewaltig², and Alessandra Pedrocchi¹

P200 Spike-based probabilistic inference with correlated noise

Ilja Bytschok¹*, Dominik Dold¹, Johannes Schemmel¹, Karlheinz Meier¹, and Mihai A. Petrovici^{1,2}

P201 Optimal refractoriness from a rate-distortion perspective

Hui-An Shen*, Simone Surace, and Jean-Pascal Pfister

Institute of Neuroinformatics, UZH and ETHZ, Zurich, Switzerland

P202 Towards online accurate spike sorting for hundreds of channels

Baptiste Lefebvre, Marcel Stimberg, Olivier Marre, and Pierre Yger*

Institut de la Vision, INSERM UMRS 968, CNRS, Paris, France

¹Department of Electronics, Information and Bioengineering, Politecnico di Milano, Milano, Italy

²Blue Brain Project, Ecole Polytechnique Fédérale de Lausanne (EPFL), Biotech Campus, Geneva, Switzerland

³Department of Brain and Behavioral Sciences, University of Pavia, Pavia, Italy

¹Kirchhoff-Institute for Physics, Heidelberg University, Heidelberg, Germany

²Department of Physiology, University of Bern, Bern, Switzerland

P203 Modeling orientation preference in the apical and basal trees of L2/3 V1 neurons

Athanasia Papoutsi^{1*}, Jiyoung Park², Ryan Ash², Stelios Smirnakis², and Panayiota Poirazi¹

P204 Dual recordings in the mouse auditory brainstem and midbrain reveal differences in the processing of vocalizations

Richard Felix¹, Alexander G Dimitrov^{1,2*}, and Christine Portfors¹

¹Department of Integrative Biology and Neuroscience, Washington State University Vancouver, Vancouver WA, USA

P205 Modelling of leg decoupling in the stick insect and its possible significance for understanding the workings of the locomotor system

Silvia Daun^{1,2*}, Tibor Toth¹

P206 Spatio-temporal dynamics of key signaling molecules in growth cones

Joanna Jedrzejewska-Szmek¹*, Nadine Kabbani^{1,2}, and Kim Avrama Blackwell^{1,3}

P207 A simulation of EMG signal generation following TMS

Bahar Moezzi^{1,2*}, Natalie Schaworonkow³, Lukas Plogmacher³, Mitchell R. Goldsworthy^{2,4}, Brenton Hordacre², Mark D McDonnell¹, Nicolangelo lannella^{1,5}, Michael C. Ridding², and Jochen Triesch³

P208 The effect of LTP, LTD and non-specific LTD on the Recognition of Sparse Noisy Patterns in Simplified and Detailed Purkinje Cell Models

Reinoud Maex¹, Karen Safaryan², and Volker Steuber³*

P209 Modeling causality of the smoking brain

Rongxiang Tang¹, Yi-Yuan Tang²*

¹ IMBB, FORTH, Heraklion, Crete, Greece

²Neurology, Baylor College of Medicine, Houston, Texas, USA

²Department of Mathematics and Statistics, Washington State University Vancouver, Vancouver WA, USA

¹Department of Animal Physiology, Institute of Zoology, University of Cologne, Cologne, Germany

²Cognitive Neuroscience, Institute of Neuroscience and Medicine (INM-3), Research Center Juelich, Juelich, Germany

¹Krasnow Institute, George Mason University, Fairfax, VA, USA

²School of Systems Biology, George Mason University, Fairfax, VA, USA

³Bioengineering Department, George Mason University, Fairfax, VA, USA

¹Computational and Theoretical Neuroscience Laboratory, School of Information Technology and Mathematical Sciences, University of South Australia, Australia

²Robinson Research Institute, School of Medicine, University of Adelaide, Australia

³Frankfurt Institute for Advanced Studies, Germany

⁴Discipline of Psychiatry, School of Medicine, University of Adelaide, Australia

⁵School of Mathematical Sciences, University of Nottingham, UK

¹Department of Cognitive Sciences, Ecole Normale Superieure, rue d'Ulm 25, Paris, France

²Department of Physics and Astronomy, Knudsen Hall, University of California, Los Angeles, CA, USA

³Centre for Computer Science and Informatics Research, University of Hertfordshire, College Lane, Hatfield, UK

¹Department of Psychology, Washington University in St. Louis, St. Louis, MO, USA

²Department of Psychological Sciences, Texas Tech University, TX, USA

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P209 Modeling causality of the smoking brain

Rongxiang Tang¹, Yi-Yuan Tang²*

P210 Modelling of calcium waves in astrocytic networks induced by neural activity

Darya V. Verveyko¹*, Alexey Brazhe², Andrey Verisokin¹, and Dmitry Postnov³

P211 Simulated voltage clamp: Offline biophysical reconstruction of fast ionic currents in large cells with uncompensated series resistance

Cengiz Gunay^{1,2*}, Gabriella Panuccio³, Michele Giugliano³, and Astrid A Prinz¹

P212 Representing and implementing cognitive sequential interactions

Pablo Varona¹*, Mikhail I. Rabinovich²

P213 An integrated neuro-mechanical model of C. elegans locomotion

Jack Denham*, Thomas Ranner, and Netta Cohen

School of Computing, University of Leeds, Leeds, UK

P214 A computational approach to understanding functional synaptic diversity: the role of nanoscale topography of Ca2+ channels and synaptic vesicles

Maria Reva^{1*}, Nelson Rebola¹, Tekla Kirizs², Zoltan Nusser², and David Digregorio¹

P215 Is object saliency perceived different cross-culturally: A computational modelling study

Eirini Mavritsaki^{1,2}*, Panos Rentzelas¹

¹Department of Psychology, Washington University in St. Louis, St. Louis, MO, USA

²Department of Psychological Sciences, Texas Tech University, TX, USA

¹Department of Theoretical Physics, Kursk State University, Kursk, Russian Federation

²Department of Biophysics, Lomonosov Moscow State University, Moscow, Russian Federation

³Department of Physics, Saratov State National Research University, Saratov, Russian Federation

¹Dept. Biology, Emory University, Atlanta, GA, USA

²School of Science and Technology, Georgia Gwinnett College, Lawrenceville, GA, USA

³Theoretical Neurobiology & Neuroengineering Lab, Dept. Biomedical Sciences, University of Antwerp, Belgium

¹ Grupo de Neurocomputación Biológica, Dpto. de Ingeniería Informática, Escuela Politécnica Superior, Universidad Autónoma de Madrid, Spain

²BioCircuits Institute, University of California, San Diego, USA

¹Laboratory of Dynamic Neuronal Imaging, Neuroscience Department, Institute Pasteur, Paris, France

²Institute of Experimental Medicine, Hungarian Academy of Sciences, Budapest, Hungary

¹Department of Psychology, Birmingham City University, Birmingham, UK

²School of Psychology, University of Birmingham, Birmingham, UK

P216 NeuroNLP: a natural language portal for aggregated fruit fly brain data

Nikul Ukani¹, Adam Tomkins²*, Chung-Heng Yeh¹, Wesley Bruning³, Allison Fenichel⁴, Yiyin Zhou¹, Yu-Chi Huang⁵, Dorian Florescu², Carlos Luna Ortiz², Paul Richmond⁶, Chung-Chuan Lo⁵, Daniel Coca², Ann-Shyn Chiang⁵, and Aurel A Lazar¹

P217 Towards prediction of plasticity response to paired cTBS from resting state network connectivity

Bahar Moezzi^{1*}, Brenton Hordacre¹, Mitchell R. Goldsworthy^{1,2}, and Michael Ridding¹

P218 Mathematical Analysis of Transient "domino effect" like Brain Dynamics

Jennifer Creaser^{1*}, Congping Lin¹, Peter Ashwin¹, Jonathan Brown², and Thomas Ridler²

P219 Synchronized neocortical dynamics during NREM sleep

Daniel Levenstein^{1,2*}, Brendon Watson^{2,3}, Gyorgy Buzsaki^{1,2}, and John Rinzel^{1,4}

P220 Accumulation process and multi-layer mechanisms of perceptual alternation in auditory streaming

Rodica Curtu^{1*}, Anh Nguyen¹, and John Rinzel²

P221 The Necessity of Sleep and Wake: Synaptic Homeostasis via System-Level Plasticity and the Ascending Arousal System

Sahand Assadzadeh^{1,2*}, Peter Robinson^{1,2}

P222 Low- and high-mode waking states in the corticothalamic system

Sanz-Leon Paula^{1,2*}, Peter Robinson^{1,2}

¹ Department of Electrical Engineering, Columbia University, New York, NY, USA

²Department of Automatic Control & Systems Engineering, The University of Sheffield, Sheffield, UK

³Department of Computer Science, Columbia University, New York, NY, USA

⁴Data Science Institute, Columbia University, New York, NY, USA

⁵Brain Research Center, National Tsing Hua University, Hsinchu, Taiwan

⁶Department of Computer Science, The University of Sheffield, Sheffield, UK

¹Robinson Research Institute, School of Medicine, University of Adelaide, Australia

²Discipline of Psychiatry, School of Medicine, University of Adelaide, Australia

¹Department of Mathematics, University of Exeter, Exeter, UK

²Institute of Biomedical and Clinical Sciences, University of Exeter Medical School, Exeter, UK

¹Center for Neural Science, New York University, New York, NY, USA

²NYU Neuroscience Institute, New York University, New York, NY, USA

³Dept. of Psychiatry, Weill Cornell Medical Center, New York, NY, USA

⁴Courant Institute for Mathematical Sciences, New York University, New York, NY, USA

¹Department of Mathematics, The University of Iowa, Iowa City, IA, USA

²Courant Institute of Mathematical Sciences, New York University, New York, NY, USA

¹School of Physics, The University of Sydney, NSW, Australia

²Center for Integrative Brain Function, The University of Sydney, NSW, Australia

¹School of Physics, University of Sydney, New South Wales, Australia

²Center for Integrative Brain Function, University of Sydney, New South Wales, Australia

P223 Closed-loop temporally structured light stimulation in weakly electric fish

Caroline Garcia Forlim^{1,2*}, Lírio O. B. de Almeida³, Ángel Lareo⁴, Reynaldo D Pinto³, Pablo Varona⁴, and Francisco B Rodríguez⁴

P224 Information-theoretic analysis of temporal code-driven stimulation applied to electroreception

Angel Lareo^{1*}, Caroline Garcia Forlim², Reynaldo D Pinto³, Pablo Varona¹, and Francisco B Rodríguez¹

P225 Gain control mechanism based on lateral inhibition of antennal lobe improves pattern recognition performance under wide concentration variability

Aaron Montero¹*, Thiago Mosqueiro², Ramon Huerta^{1,2}, and Francisco B Rodríguez¹

P226 Maximum Relative Area as a Feature for Adaptability in ERP-based BCI Systems

Vinicio Changoluisa^{1,2}*, Pablo Varona¹, and Francisco B Rodríguez¹

P227 Intrinsically stochastic neuron models for use in network simulations

Vinícius Cordeiro, César Ceballos, Nilton Kamiji, and Antonio C Roque*

Departamento de Física-FFCLRP, Universidade de São Paulo, Brazil

P228 Modeling action potential and network effects after site-directed RNA editing of sodium channels

William W Lytton^{1,2}*, Andrew Knox³, and Joshua Rosenthal⁴

¹Clinic and Policlinic for Psychiatry and Psychotherapy, University Medical Center Hamburg-Eppendorf, Hamburg, Germany

²Departamento de Física Geral, Universidade de Sao Paulo, Sao Paulo, Brazil

³Instituto de Física de Sao Carlos, Universidade de Sao Paulo, Sao Carlos, Brazil

⁴Escuela Politécnica Superior, Universidad Autonoma de Madrid, Madrid, Spain

¹ Grupo de Neurocomputación Biológica, Departamento de Ingeniería Informática, Escuela Politécnica Superior, Universidad Autónoma de Madrid, Spain

²Clinic and Policlinic for Psychiatry and Psychotherapy, University Medical Center, Hamburg-Eppendorf, Hamburg, Germany

³Lab. Neurodynamics/Neurobiophysics - Dept. Physics and Interdisciplinary Sciences - Institute of Physics of São Carlos, Universidade de São Paulo - Brazil

¹ Grupo de Neurocomputación Biológica, Dpto. de Ingeniería Informática, Escuela Politécnica Superior, Universidad Autónoma de Madrid, Madrid, Spain

²BioCircuits Institute, University of California, San Diego, La Jolla, CA, USA.

¹ Grupo de Neurocomputación Biológica, Dpto. de Ingeniería Informática. Escuela Politécnica Superior. Universidad Autónoma de Madrid, Spain

²Universidad Politécnica Salesiana, Quito, Ecuador

¹Depts. of Physiology&Pharmacology and Neurology, SUNY Downstate, Brooklyn, NY, USA

²Dept. of Neurology, Kings County Hospital, Brooklyn, NY, USA

³Dept. of Neurology, University of Wisconsin, Madison, WI, USA

⁴Dept. of Neurobiology, Marine Biological Laboratory, Woods Hole, MA, USA

P229 Movement-related delta-theta synchronization in young and elderly healthy subjects

Silvia Daun^{1,2*}, Svitlana Popovych^{1,2}, Liqing Liu^{1,2}, Bin Wang¹, Tibor Toth², Christian Grefkes^{1,3}, Gereon Fink^{1,3}, and Nils Rosjat^{1,2}

P230 ePyNN: a low cost embedded system for simulating Spiking Neural Networks

Abraham Perez-Trujillo¹, Andres Espinal², Marco A. Sotelo-Figueroa², Ivan Cruz-Aceves³, and Horacio Rostro-Gonzalez¹*

P231 Temporal structure of bilateral coherence in essential and physiological hand tremor

Martin Zapotocky^{1,2⋆}, Soma Chakraborty^{1,2}, Martina Hoskovcova², Jana Kopecka², Olga Ulmanova², and Evzen Ruzicka²

P232 Detecting joint pausiness in parallel spike trains

Matthias Gaertner¹*, Sevil Duvarci², Jochen Roeper², and Gaby Schneider¹

P233 A stochastic model relates responses to bistable stimuli to underlying neuronal processes

Stefan Albert^{1*}, Katharina Schmack², and Gaby Schneider¹

P234 Function and energy consumption constrain biophysical properties of neurons - an example from the auditory brainstem

Michiel Remme^{1*}, John Rinzel², and Susanne Schreiber¹

P235 The Brain Simulation Platform of the Human Brain Project: collaborative web applications and tools for data-driven brain models

Michele Migliore¹, Carmen Alina Lupascu¹*, Luca Leonardo Bologna¹, Rosanna Migliore¹, Stefano M. Antonel², Jean-Denis Courcol², and Felix Schürmann²

¹Cognitive Neuroscience, Institute of Neuroscience and Medicine (INM-3), Research Center Juelich, Juelich, Germany

²Heisenberg Research Group of Computational Neuroscience – Modeling Neural Network Function, Department of Animal Physiology, Institute of Zoology, University of Cologne, Cologne, Germany

³Department of Neurology, University Hospital Cologne, Cologne, Germany

¹Department of Electronics, University of Guanajuato, Guanajuato, Mexico

²Department of Organizational Studies, University of Guanajuato, Guanajuato, Mexico

³CONACYT, Mathematics Research Center (CIMAT), Guanajuato, Mexico

¹Institute of Physiology, Czech Academy of Sciences, Prague, Czech Republic

²Department of Neurology, First Faculty of Medicine, Charles University in Prague, Czech Republic

¹Institute of Mathematics, Goethe-University, Frankfurt, Germany

²Neuroscience Center, Institute of Neurophysiology, Goethe-University, Frankfurt, Germany

¹ Institute of Mathematics, Goethe-University, Frankfurt, Germany

²Department of Psychiatry and Psychotherapy, Charité Universitätsmedizin, Berlin, Germany

¹Institute for Theoretical Biology, Humboldt University, and Bernstein Center for Computational Neuroscience, Berlin, Germany

²Center for Neural Science and Courant Institute of Mathematical Sciences, New York University, New York, NY, USA

¹Institute of Biophysics, National Research Council (CNR), Palermo, Italy

²Blue Brain Project, École Polytechnique Fédérale de Lausanne (EPFL), Geneva, Switzerland

P236 A Single Pyramidal-Cell and Network Computational Model of the Hippocampal CA3 Region

Sami Çelikok*, Eva M. Navarro-López, and Neslihan Serap Sengör

P237 Functional connectivity between prefrontal cortex and striatum showed by computational model

Rahmi Elibol*, Neslihan Serap Sengör

Electronics and Communication Engineering, Istanbul Technical University, Istanbul, Turkey

P238 A spiking neural network model of basal ganglia-thalamocortical circuit with Brian2

Mustafa Özdemir*, Neslihan Serap Sengör

Electronic-Communication Department, Istanbul Technical University, Istanbul, Türkiye

P239 Coordinate-transformation spiking neural network for spatial navigation

Tianyi Li*, Angelo Arleo, and Denis Sheynikhovich

Sorbonne Universités, UPMC Univ Paris 06, INSERM, CNRS, Institut de la Vision, Paris, France

P240 Micro-connectomics with cognitive task selectivity

Akihiro Nakamura¹, Masanori Shimono^{1,2}*

P241 Does reinforcement leaning explain zone-allocation behavior between two competing mice?

Youngio Song¹*, Sol Park^{1,2}, Ilhwan Choi², Jaeseung Jeong^{1,3}, and Hee-Sup Shin²

P242 Optimal synaptic scaling emerges from Hebbian learning rules in balanced networks

Sadra Sadeh, Padraig Gleeson*, and R Angus Silver

Department of Neuroscience, Physiology and Pharmacology, University College London, London, UK

P243 Deciphering the contributions of oriens-lacunosum/moleculare (OLM) cells during local field potential (LFP) theta rhythms in CA1 hippocampus

Alexandra Chatzikalymniou^{1,2}★, Frances Skinner^{1,3,2}

P244 Nonlinear optimal control of brain networks

Lazaro Sanchez-Rodriguez*, Roberto Sotero

Hotchkiss Brain Institute and Department of Radiology, University of Calgary, Calgary, Alberta, Canada

P245 An inhibitory microcircuit that amplifies the redistribution of somatic and dendritic inhibition

Loreen Hertaeg*, Owen Mackwood, and Henning Sprekeler

Modelling of Cognitive Processes, Berlin Institute of Technology and Bernstein Center for Computational Neuroscience, Berlin, Germany

¹Osaka University, Toyonaka, Osaka, Japan

²Riken Brain Science Institute, Saitama, Japan

¹Department of Bio and Brain Engineering, KAIST, Daejeon, Rep. of Korea

²Center for Cognition and Sociality, IBS, Daejeon, Rep. of Korea

³Program of Brain and Cognitive Engineering, KAIST, Daejeon, Rep. of Korea

¹Krembil Research Institute, University Health Network, Toronto, ON, Canada

²Department of Physiology, University of Toronto, Toronto, ON, Canada

³Department of Medicine (Neurology), University of Toronto, Toronto, ON, Canada

P246 Learning grid cells in recurrent neural networks

Steffen Puhlmann¹*, Simon N. Weber^{1,2}, and Henning Sprekeler^{1,2}

P247 A model of perceptual learning, biases, and roving

David Higgins^{1,2*}, Henning Sprekeler^{1,2}

P248 Presynaptic inhibition provides a rapid stabilization of recurrent excitation in the face of plasticity

Laura Naumann^{1,2*}, Henning Sprekeler^{1,2}

P249 A grid score for individual spikes of grid cells

Simon N. Weber^{1,2*}, Henning Sprekeler^{1,2}

P250 Cortical circuits implement optimal integration of context

Ramakrsinan Iyer, Stefan Mihalas*

Allen Institute for Brain Science, Seattle, WA, USA

P251 Neural cross-frequency coupling functions in the resting state with eyes open and eyes closed

Valentina Ticcinelli¹*, Tomislav Stankovski^{1,2}, Peter V.e. McClintock¹, and Aneta Stefanovska¹

P252 Dissecting the total astrocytic potassium current in a computational model

Predrag Janjic¹*, Dimitar Solev³, Gerald Seifert², Ljup_o Kocarev¹, and Christian Steinhäuser²

P253 Information rate of multiple synaptic release sites with separately released vesicles during short-term depression

Mehrdad Salmasi^{1,2,3*}, Stefan Glasauer^{1,2,3,4}, and Martin Stemmler^{2,5}

¹MKP, Modelling of cognitive processes, Berlin Institute of Technology, Berlin, Germany

²Bernstein Center for Computational Neuroscience, Berlin, Germany

¹ Modelling of Cognitive Processes, TU Berlin, Germany

²Bernstein Center for Computational Neuroscience, Berlin, Germany

¹ Modelling of Cognitive Processes, Berlin Institute of Technology, Berlin, Germany

²Bernstein Center for Computational Neuroscience, Berlin, Germany

¹Berlin Institute of Technology, Berlin, Germany

²Bernstein Center for Computational Neuroscience, Berlin, Germany

¹Department of Physics, Lancaster University, Lancaster, United Kingdom

²Faculty of Medicine, Ss Cyril and Methodius University, Skopje, Macedonia

¹Laboratory for Complex Systems and Networks, Macedonian Academy of Sciences and Arts, Skopje, Macedonia

²Institute of Cellular Neurosciences, University of Bonn Medical School, Bonn, Germany

³Unaffiliated, dimitar.solev@gmail.com

¹ Graduate School of Systemic Neurosciences, Ludwig-Maximilian University, Munich, Germany

²Bernstein Center for Computational Neuroscience, Munich, Germany

³German Center for Vertigo and Balance Disorders, Ludwig-Maximilian University, Munich, Germany

⁴Department of Neurology, Ludwig-Maximilian University, Munich, Germany

⁵Department of Biology II, Ludwig-Maximilian University, Munich, Germany

P254 Properties of recurrent networks at maximum capacity for storing sequences of network states

Danke Zhang, Chi Zhang, and Armen Stepanyants*

Department of Physics and Center for Interdisciplinary Research on Complex Systems, Northeastern University, Boston, MA, USA

P255 Modulation of epileptic activity in thalamo-cortical networks by input from the cerebellar nuclei

Julia Goncharenko¹, Lieke Kros², Neil Davey¹, Christoph Metzner¹, Chris de Zeeuw², Freek Hoebeek², and Volker Steuber¹*

P256 The effect of homeostatic structural plasticity on associative memory in a network with spiketime dependent inhibitory synaptic plasticity

Ankur Sinha*, Christoph Metzner, Roderick Adams, Michael Schmuker, Neil Davey, and Volker Steuber *Biocomputation Group, University of Hertfordshire, Hatfield, UK*

P257 The dependence of arithmetic operations on input location in cerebellar nucleus and cortical pyramidal neurons

Maria Psarrou[⋆], Maria Schilstra, Neil Davey, Benjamin Torben-Nielsen, Michael Schmuker, and Volker Steuber

Centre for Computer Science and Informatics Research, University of Hertfordshire, UK

P258 A Framework for Automated Validation and Comparison of Models of Neurophysiological and Neurocognitive Biomarkers of Psychiatric Disorders

Christoph Metzner^{1*}, Achim Schweikard², Tuomo Maki-Marttunen³, Bartosz Zurowski⁴, and Volker Steuber¹

P259 Synergetic and redundant information flow in dynamical systems: an operative definition based on prediction

Daniele Marinazzo¹*, Luca Faes², and Sebastiano Stramaglia³

P260 Forming and Using Hierarchical Cognitive Maps: a Neural Network Model

Henry Jordan*, Simon Stringer

OFTNAI, Dept. Experimental Psychology, University of Oxford, UK

P261 Harmonic SSEP Spectra are Determined by Modulation of Population Firing Rate – a Modeling Study

Elzbieta Gajewska-Dendek*, Piotr Suffczynski

Department of Biomedical Physics, Institute of Experimental Physics, University of Warsaw, Warsaw, Poland

¹Centre for Computer Science and Informatics Research, University of Hertfordshire, Hatfield, UK

²Department of Neuroscience, Erasmus MC, Wytemaweg 80, Rotterdam, The Netherlands

¹Centre for Computer Science and Informatics Research, University of Hertfordshire, United Kingdom

²Institute for Robotics and Cognitive Systems, University of Luebeck, Luebeck, Germany

³NORMENT, Institute of Clinical Medicine, University of Oslo, Norway

⁴Department of Psychiatry, University of Luebeck, Schleswig-Holstein, Luebeck, Germany

¹Department of Data Analysis, Ghent University, Ghent, Belgium

²BIOtech, Dept. of Industrial Engineering, University of Trento, and IRCS-PAT FBK, Trento, Italy

³Dipartimento di Fisica, Università degli Studi Aldo Moro, Bari, and INFN, Sezione di Bari, Bari, Italy

P262 Computational measure to account for erroneous neural deactivation when oxygen supply cannot meet metabolic demand in neuroimaging studies

Nicoladie Tam¹*, George Zouridakis², and Luca Pollonini²

P263 Mapping large-scale brain networks using multivariate pattern analysis

Yi-Yuan Tang1*, Rongxiang Tang2, and J Lewis-Peacock3

P264 Interplay between propagation delay and frequency of oscillation determines emergent structures of neuronal networks driven by triplet-based STDP

Mojtaba Madadi Asl¹*, Alireza Valizadeh^{1,2}, and Peter A. Tass³

P265 X078 Plasticity and network implications of a synaptic LPA-signalling pathway

Andreas Nold^{1*}, Wei Fan², Sara Konrad¹, Heiko Endle², Johannes Vogt², and Tatjana Tchumatchenko¹

P266 Postsynaptic Activity-Dependent Synaptic Scaling Enables the Functional Organization of Memories

Juliane Herpich^{1,2*}, Christian Tetzlaff^{1,2}

P267 Input-dependent Synaptic Consolidation of Memory Representations

Jannik Luboeinski¹*, Christian Tetzlaff²

P268 Why working memory is not a reservoir: the role of transient dynamics and attractors when processing unreliably timed inputs

Timo Nachstedt^{1,2*}, Christian Tetzlaff^{1,2}

¹Department of Biological Sciences, University of North Texas, Denton, TX, USA

²Departments of Engineering Technology, Computer Science, and Electrical and Computer Engineering, University of Houston, Houston, TX, USA

¹Department of Psychological Sciences, Texas Tech University, TX, USA

²Department of Psychological & Brain Sciences, Washington University in St. Louis, St. Louis, MO, USA

³Department of Psychology, The University of Texas at Austin, Austin, TX, USA

¹Department of Physics, Institute for Advanced Studies in Basic Sciences (IASBS), Zanjan, Iran

²School of Cognitive Sciences, Institute for Research in Fundamental Sciences (IPM), Tehran, Iran

³Department of Neurosurgery, School of Medicine, Stanford University, Stanford, CA, USA

¹Theory of Neural Dynamics, Max Planck Institute for Brain Research, Frankfurt, Germany

²Institute for Microscopic Anatomy and Neurobiology, University Medical Center, Johannes Gutenberg University, Mainz, Germany

¹Third Institute of Physics – Biophysics, Department of Computational Neuroscience, University of Goettingen, Goettingen, Germany

²Bernstein Center for Computational Neuroscience, University of Goettingen, Goettingen, Germany

¹ Third Institute of Physics – Biophysics, Department of Computational Neuroscience, University of Göttingen, Göttingen, Germany

²Bernstein Center for Computational Neuroscience, University of Göttingen, Göttingen, Germany

¹Third Institute of Physics, Universität Göttingen, Göttingen, Germany

²Bernstein Center for Computational Neuroscience, Göttingen, Germany

P269 Application of spike train synchrony measure Spike-contrast to quantify the effect of bicuculline on cortical networks grown on microelectrode arrays

Manuel Ciba¹*, Andreas Bahmer², and Christiane Thielemann¹

P270 Cortical states affect the optimal linear readout of network dynamics

Eric Kuebler^{1*}, Joseph S. Tauskela², and Jean-Philippe Thivierge¹

¹School of Psychology, Faculty of Social Science, University of Ottawa, Ottawa, Ontario, Canada

P271 Workflow, data format and tools to register neuron morphologies to a reference brain atlas

Rembrandt Bakker^{1,2*}, María García-Amado³, Marian Evangelio³, Francisco Clascá³, and Paul Tiesinga¹

¹Neuroinformatics department, Donders Institute for Brain, Cognition and Behaviour, Radboud University Nijmegen, Nijmegen, The Netherlands

²Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6) and JARA BRAIN Institute I, Jülich Research Centre, Jülich, Germany

³Departamento de Anatomía, Histología y Neurociencia, Facultad de Medicina, Universidad Autónoma de Madrid, Madrid, Spain

P272 Simple models of closed-loop cortical-environment interactions

Christopher L Buckley^{1*}, Taro Toyoizumi²

¹Informatics, University of Sussex, Brighton, UK

P273 Encoding multiple spaces in grid-cells networks

Alexis Dubreuil¹*, Rémi Monasson¹, and Alessandro Treves²

¹Laboratoire de Physique Théorique, Ecole Normale Supérieure, Paris, France

P274 Storage capacity of threshold-linear networks for gridlike continuous attractors

Davide Spalla*, Sophie Rosay, and Alessandro Treves

Cognitive Neuroscience Sector, SISSA, Trieste, Italy

P275 A network model of the turnover dynamics of excitatory and inhibitory synapses in cortex

Florence Kleberg*, Jochen Triesch

Neuroscience Lab, Frankfurt Institute for Advanced Studies, Frankfurt am Main, Germany

P276 Retinal prostheses from stimulation to seeing

Willy Wong^{1,2}*, Bruno de Oliveira Floriano¹, Toshihiko Matsuo³, and Tetsuya Uchida⁴

¹Biomems lab, Faculty of Engineering, UAS Aschaffenburg, Aschaffenburg, Germany

²Comprehensive Hearing Center, University ENT-Clinic Würzburg, Würzburg, Germany

²Human Health Therapeutics, National Research Council of Canada, Ottawa, Ontario, Canada

²Brain Science Institute, RIKEN, Wako, Saitama, Japan

²Cognitive Neuroscience, SISSA, Trieste, Italy

¹Dept. of Electrical and Computer Engineering, University of Toronto, Toronto, ON, Canada

²Institute of Biomaterials and Biomedical Engineering, University of Toronto, Toronto, ON, Canada

³Ophthalmology, Okayama University Medical School, Okayama, Japan

⁴Polymer Materials Science, Faculty of Engineering, Okayama University, Okayama, Japan

P277 Metabolic cost of neuronal oscillations

Domenica Dibenedetto¹*, Kamil Uludag^{1,2}

P278 Multi-cluster structure and dynamics in networks of coupled phase oscillators through different classes of STDP profiles

Abdorreza Goodarzi Nick1*, Mojtaba Madadi Asl1, and Alireza Valizadeh1,2

P279 Multi-scale network structure of macaque visual cortex: connectivity map, cortical architecture, and layer-specific pathways

Maximilian Schmidt^{1,2}, Rembrandt Bakker^{1,3}, Claus Hilgetag^{4,5}, Markus Diesmann^{1,6,7}, and Sacha J van Albada¹*

P280 The interaction of synaptic and structural plasticity in recurrent networks

Michael Fauth*, Mark van Rossum

School of Informatics, University of Edinburgh, Edinburgh, United Kingdom

P281 Automatic calibration for hybrid circuits of living and artificial neurons

Manuel Reyes-Sánchez*, Irene Elices Ocon, Rodrigo Amaducci, Carlos Muñiz, Francisco B Rodríguez, and Pablo Varona

1 Grupo de Neurocomputación Biológica, Dpto. de Ingeniería Informática, Escuela Politécnica Superior, Universidad Autónoma de Madrid, Spain

P282 Role of asymmetry in shaping spiking-bursting activity of Central Pattern Generators

Irene Elices Ocon¹*, David Arroyo¹, Rafael Levi^{1,2}, Francisco B Rodríguez¹, and Pablo Varona¹

P283 Rivalry with irregular spiking: resolving mutual inhibition and the balanced state

Ben Cohen*, Carson Chow, and Shashaank Vattikuti

Lab of Biological Modeling, NIDDK/NIH, Bethesda, MD, USA

¹Maastricht Centre for Systems Biology (MaCSBio), Maastricht University, The Netherlands

²Maastricht Brain Imaging Centre (MBIC), Faculty of Psychology & Neuroscience, Maastricht University

¹ Department of Physics, Institute for Advanced Studies in Basic Sciences (IASBS), Zanjan, Iran

²School of Cognitive Sciences, Institute for Research in Fundamental Sciences (IPM), Tehran, Iran

¹ Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6), Jülich Research Centre and JARA, Jülich, Germany

²Laboratory for Neural Circuit Theory, RIKEN Brain Science Institute, Wako, Japan

³Donders Institute for Brain, Cognition and Behavior, Radboud University Nijmegen, Nijmegen, Netherlands

⁴Department of Computational Neuroscience, University Medical Center Eppendorf, Hamburg, Germany

⁵Department of Health Sciences, Boston University, Boston, MA, USA

⁶Department of Psychiatry, Psychotherapy and Psychosomatics, Medical Faculty, RWTH Aachen University, Aachen, Germany

⁷Department of Physics, Faculty 1, RWTH Aachen University, Aachen, Germany

¹Grupo de Neurocomputación Biológica, Dpto. de Ingeniería Informática, Escuela Politécnica Superior, Universidad Autónoma de Madrid, Spain

²Department of Biological Sciences, University of Southern California, CA, USA

P284 Interplay between inhibition and connectivity structure in driving synchronization and functional properties of neural networks

Elena Bertolotti^{1,2*}, Raffaella Burioni^{1,2}, Matteo Di Volo^{3,4,5}, and Alessandro Vezzani^{1,6}

P285 A mechanistic model of dopamine modulated learning in the olfactory system of Drosophila

Bayar Menzat*, Tim Vogels

Centre for Neural Circuits and Behaviour, University of Oxford, Oxford, UK

P286 Saliency-based Gaze Prediction based on the Neural Population for Integrating the Direction of Figure

Nobuhiko Wagatsuma*

School of Science and Engineering, Tokyo Denki University, Hiki, Saitama 350-0394, Japan

P287 Investigating the differences in ion channel properties between ON and OFF ganglion cells: a combination of modelling and optimization approach

Susmita Saha^{1*}, Reena Kapoor¹, Robert Kerr², and John Wagner¹

P288 Response reversal during top-down modulation in cortical circuits with multiple interneuron types

Luis Garcia Del Molino¹*, Guangyu Robert Yang¹, Jorge F Mejias¹, and Xiao-Jing Wang^{1,2}

P289 Cellular and Network Modeling of the Striatal Microcircuit in a Mouse Model of Huntington's Disease

Hanbing Song^{1*}, Joseph Goodliffe², Jennifer Luebke², and Christina M Weaver¹

P290 Convolutional Neural Network-based Interictal Epileptiform Discharge Detection

John Thomas¹, Nishant Sinha^{2,3}, Nikhita Shaju^{1,4}, Tomasz Maszczyk¹, Jing Jin¹, Sydney Cash⁵, Justin Dauwels¹*, and Brandon Westover⁵

¹Department of Mathematical, Physical and Computer Sciences, University of Parma, Parma, Italy

²INFN, Gruppo Collegato di Parma, Parma, Italy

³Group for Neural Theory, Departément des Etudes Cognitives, Ecole Normale Supérieure, Paris, France

⁴Centro Interdipartimentale per lo Studio delle Dinamiche Complesse, Sesto Fiorentino, Italy

⁵Indiana University-Purdue University, Indianapolis, Indiana, USA

⁶IMEM-CNR, Parma, Italy

¹IBM Research Australia, Australia

²NeuroEngineering Laboratory, Electrical and Electronic Engineering, The University of Melbourne, Australia

¹Center for Neural Science, New York University, New York, NY, USA

²NYU-ECNU Institute of Brain and Cognitive Science, NYU Shanghai, Shanghai, China

¹Department of Mathematics, Franklin and Marshall College, Lancaster, PA, USA

²Department of Anatomy and Neurobiology, Boston University School of Medicine, Boston, MA, USA

¹School of Electrical and Electronic Engineering, Nanyang Technological University, Singapore

²Institute of Neuroscience, Faculty of Medical Sciences, Newcastle University, Newcastle upon Tyne, UK

³School of Computing Science, Newcastle University, Newcastle upon Tyne, UK

⁴Department of Electrical and Electronics Engineering, Vellore Institute of Technology, India

⁵Neurology Department, Massachusetts General Hospital and Harvard Medical School, MA, USA

P291 Effects of Hebbian learning on networks of Kuramoto phase oscillators with time delay

Maryam Karimian^{1*}, Domenica Dibenedetto¹, Michelle Moerel^{1,2}, Peter de Weerd^{1,2}, Thomas Burwick⁴, and Ronald L Westra^{1,3}

P292 Biophysical modelling of resting brain states and inhibitory synaptic plasticity

Romesh Abeysuriya^{1*}, Jonathan Hadida^{1,2}, Stamatios Sotiropoulos², Saad Jbabdi², and Mark Woolrich^{1,2}

P293 Temporal pattern recognition and control of animat foraging by evolving small networks of adaptive exponential integrate-and-fire neurons

Chama Bensmail^{1,2*}, Volker Steuber¹, and Borys Wrobel^{2,3}

P294 Learning Continuous Attractor Neural Networks from Continuously Morphed Patterns

Xiaolong Zou^{1,2*}, Zilong Ji^{1,2}, Xiao Liu^{1,2}, Yan Xia^{1,2}, and Si Wu^{1,2}

P295 Learning Peri-saccadic Receptive Field Remapping in Lateral Intraparietal Area from Visual Experience

Xiao Wang*, Mingsha Zhang, and Si Wu

State Key Laboratory of Cognitive Neuroscience & Learning, IDG/McGovern Institute for Brain Research, Beijing Normal University, Beijing, CHINA

P296 Axonal tree morphology dictates information coding

Netanel Ofer^{1,2*}, Orit Shefi^{1,2}, and Gur Yaari¹

P297 The Neuroscience Gateway Portal-High Performance Computing for Neuroscientists

Nicholas Carnevale^{1*}, Amit Majumdar², Subhashini Sivagnanam², and Kenneth Yoshimoto²

P298 Changes of electrophysiological neuronal properties in lithium-pilocarpine model of epilepsy

Elena Smirnova^{1,2*}, Dmitry Amakhin², Sergey Malkin², Anton Chizhov^{1,2}, and Aleksey Zaitsev²

¹Maastricht Centre for Systems Biology, Maastricht University, The Netherlands

²Department of Cognitive Neuroscience, Maastricht University, The Netherlands

³ Department of Data Science and Knowledge Engineering, Maastricht University, The Netherlands

⁴ Frankfurt Institute for Advanced Studies, Goethe University Frankfurt, Frankfurt am Main, Germany

¹Oxford Centre for Human Brain Activity, Oxford, United Kingdom

²Oxford Centre for Functional MRI of the Brain, Oxford, United Kingdom

¹Centre for Computer Science and Informatics Research, University of Hertfordshire, UK

²Biology, Adam Mickiewicz University, Poznan, Poland

³Systems Modelling IOPAN, Sopot, Poland

¹ State Key Laboratory of Cognitive Neuroscience & Learning, Beijing Normal University, Beijing, CHINA

²IDG/McGovern Institute for Brain Research, Beijing Normal University, Beijing, CHINA

¹ Faculty of Engineering, Bar Ilan University, Ramat Gan, Israel

²Institute for Nanotechnology and Advanced Materials, Bar Ilan University, Ramat Gan, Israel

¹Neuroscience Department, Yale University, New Haven, CT, USA

²San Diego Supercomputer Center, University of California San Diego, San Diego, CA, USA

¹ Ioffe Institute, St. Petersburg, Russia

²Sechenov Institute of Evolutionary Physiology and Biochemistry of RAS, St. Petersburg, Russia

P299 Depolarizing GABA leads to interneuron-based interictal discharges: experimental and mathematical models

Anton Chizhov^{1,2*}, Dmitry Amakhin², and Aleksey Zaitsev²

P300 Neural activity in distinct navigation modes of flying pigeons

Margarita Zaleshina*, Alexander Zaleshin

Moscow Institute of Physics and Technology, Moscow, Russia

P301 The Role of the Receptive Field Structure in Neuronal Compressive Sensing Signal Processing

Victor Barranca*, George Zhu

Department of Mathematics and Statistics, Swarthmore College, Swarthmore, PA, USA

P302 FuNS with E/I balance: critical dynamics maximize stability of neural networks

Quinton Skilling¹*, Daniel Maruyama², Nicolette Ognjanovski³, Sara Aton³, and Michal Zochowski^{1,2}

P303 Neural oscillations modulate the network dynamics around E-I balance in memory consolidation

Jiaxing Wu^{1*}, Nicolette Ognjanovski², Sara Aton², and Michal Zochowski³

P304 Cellular and network properties of interneuron networks dictate variable clustering patterns in both strictly inhibitory and E-I neural networks

Scott Rich1*, Victoria Booth2, and Michal Zochowski3

P305 Synaptic failure in functional network activity

Maral Budak^{1*}, Michal Zochowski^{1,2}

P306 Data-driven multiscale model of mouse M1 microcircuits

Salvador Dura-Bernal^{1*}, Samuel Neymotin^{1,3}, Benjamin A Suter⁴, Gordon M Shepherd², and William W Lytton^{1,5}

¹ Ioffe Institute, St. Petersburg, Russia

²Sechenov Institute of Evolutionary Physiology and Biochemistry of RAS, St. Petersburg, Russia

¹Biophysics Program, University of Michigan, Ann Arbor, MI, USAXS

²Department of Physics, University of Michigan, Ann Arbor, MI, USA

³Department of Cellular, Molecular, and Developmental Biology, Ann Arbor, MI, USA

¹Applied Physics Program, University of Michgan, Ann Arbor, MI, USA

²Department of Molecular, Cellular, and Developmental Biology, University of Michgan, Ann Arbor, MI, USA

³Biophysics Program and Department of Physics, University of Michgan, Ann Arbor, MI, USA

¹Applied and Interdisciplinary Mathematics Program, University of Michigan, Ann Arbor, MI, USA

²Departments of Anesthesiology and Mathematics, University of Michigan, Ann Arbor, MI, USA

³Departments of Physics and Biophysics, University of Michigan, Ann Arbor, MI, USa

¹Department of Biophysics, University of Michigan, Ann Arbor, Michigan, USA

²Department of Physics, University of Michigan, Ann Arbor, Michigan, USA

¹Department Physiology & Pharmacology, SUNY Downstate, Brooklyn, NY, USA

²Department Physiology, Northwestern University, Chicago, IL, USA

³Department Neuroscience, Brown University, Providence, RI, USA

⁴Institute of Science and Technology (IST) Austria, Klosterneburg, Austria

⁵Kings County Hospital Center, Brooklyn, NY, USA

P307 Membrane resonance and oscillation preferences of a multi-compartment model pyramidal neu-

Melvin Felton¹, Alfred Yu², David Boothe², Kelvin Oie², and Piotr J Franaszczuk^{2,3}*

P308 Network cloning using DNA barcodes

Sergey Shuvaev, Batuhan Baserdem, Anthony Zador, and Alexei Koulakov*

Cold Spring Harbor Laboratory, Cold Spring Harbor, NY, USA

P309 Triads of synchronized theta cycles boost Cross-Frequency Coupling during novelty exploration

Victor J. Lopez-Madrona^{1*}, Ernesto Pereda², Claudio Mirasso³, and Santiago Canals¹

P310 Simulations of synaptic integration in a detailed Purkinje cell model

Stefano Masoli^{1*}, Egidio Dangelo^{1,2}

P311 A model for tactile stimuli processing in cuneate nucleus

Udaya Bhaskar Rongala¹, Alberto Mazzoni¹*, Anton Spanne², Henrik Jorntell², and Calogero Maria Oddo¹

P312 A new method of system visualization of cognitive functioning for fMRI

Alexander Vartanov¹, Anastasia Neklyudova¹*, Stanislav Kozlovskiy¹, Andrey Kiselnikov¹, and Julia Marakshina^{1,2}

P313 Synaptic distribution predicts unitary LFP fields in the hippocampus and in the neocortex

Maria Telenczuk, Bartosz Telenczuk*, and Alain Destexhe

European Institute for Theoretical Neuroscience, Paris, France; Unité de Neurosciences, Information et Complexité, Gif-sur-Yvette, France

¹US Army Research Laboratory, Adelphi, MD, USA

²US Army Research Laboratory, Aberdeen Proving Ground, MD, USA

³Department of Neurology, Johns Hopkins University School of Medicine, Baltimore, MD, USA

¹ Instituto de Neurociencias, Consejo Superior de Investigaciones Científicas, Universidad Miguel Hernández, Sant Joan d'Alacant, Spain

²Departamento de Ingeniería Industrial, Escuela Superior de Ingeniería y Tecnología, Universidad de La Laguna Avda. Astrofísico Fco. Sanchez, s/n, La Laguna, Tenerife, Spain

³Instituto de Física Interdisciplinar y Sistemas Complejos, CSIC-UIB, Campus Universitat de les Illes Balears, Palma de Mallorca, Spain

¹Department of Brain and Behavioural Science, Neurophysiology and Neurocomputation Unit, University of Pavia, Pavia, Italy

²Brain Connectivity Center, Istituto Neurologico IRCCS C. Mondino, Pavia, Italy

¹The BioRobotics Institute, Scuola Superiore Sant'Anna, Pontedera, Pisa, Italy

²Neural Basis of Sensorimotor Control, Department of Experimental Medical Science, Lund University, Sweden

¹Department of Psychology, Lomonosov Moscow State University, Moscow, Russia

²Psychological Institute of Russian Academy of Education, Moscow, Russia

P314 Differential tuning of the low and high-frequency components of the neurophonic spectrum reveals the spike contribution of barn owl's nucleus laminaris neurons

Paula Kuokkanen^{1,2,3*}, Anna Kraemer³, Thomas McColgan¹, Catherine Carr³, and Richard Kempter^{1,2}

¹ Inst. For Theoretical Biology, Humboldt-Universitaet Zu Berlin, Berlin, Germany

²Bernstein Center for Computational Neuroscience Berlin, Berlin, Germany

³Dept. of Biol., Univ. of Maryland, College Park, MD, USA

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