## A whole-brain spiking neural network **P73** model linking basal ganglia, cerebellum, cortex and thalamus

Carlos Gutierrez<sup>1</sup>, Zhe Sun<sup>2</sup>, Hiroshi Yamaura<sup>3</sup>, Heidarinejad Morteza<sup>2</sup>, Jun\_Igarashi<sup>2</sup>, Tadashi Yamazaki<sup>3</sup>, Markus Diesmann<sup>4</sup>, Jean Lienard<sup>1</sup>, Benoit Girad<sup>5</sup>, Gordon Arbuthnott<sup>1</sup>, Hans Ekkerhard Plesser<sup>6</sup>, Kenji Doya<sup>1</sup>

<sup>1</sup>Neural Computation Unit, Okinawa Institute of Science and Technology Graduate University, Okiniwa, Japan <sup>2</sup> Computational Engineering Applications Unit, Head Office for Information Systems and Cybersecurity RIKEN, Saitama, Japan <sup>3</sup> Department of Computer and Network Engineering, The University of Electro-Communications, Tokyo, Japan <sup>4</sup> Institute of Neuroscience and Medicine (INM-6) and Institute for Advanced Simulation (IAS-6), Jülich Research Centre, Jülich, <sup>5</sup> Institut des Systèms Intelligents et de Robotique, Université Pierre et Marie CURIE, Paris, France <sup>6</sup>Faculty of Science and Technology, Norwegian University of Life Sciences, Ås, Norway



## Introduction

In order to investigate the dynamic nature of the whole-brain network, we built biologically constrained spiking neural network models of the basal ganglia [1,2,3], cerebellum, thalamus, and the cortex [4,5] and ran an integrated simulation using K supercomputer [8] using NEST 2.16.0 [6,7,9].

S1: Sensory Motor cortex, M1: Motor cortex, BG: Basal ganglia, **TH**: Thalamus, and **CB**: Cerebellum.



## **Resting State Simulations on K computer**

We replicated resting state activities of 2.5 biological seconds of time in models with increasing scales, from 1x1mm<sup>2</sup> to 7x7mm<sup>2</sup> of cortical surface, and observed plausible values of excitatory and inhibitory populations firing rates. Largest simulated model includes 51 million neurons and 54 billion synapses, more than an entire hemisphere of a mouse brain [Herculano-Houzel, 2009] To our knowledge, this is the first time a



**Future work**: further investigate closed loops dynamics. Implement reinforcement learning using D1 and D2 MSN and integration of a virtual/robotic arm. Test model scaling using plastic synapses. Model preparation for Fugaku supercomputer.

[1] Liénard, Jean, and Benoît Girard. "A biologically constrained model of the whole basal ganglia addressing the paradoxes of connections and selection." Journal of computational neuroscience 36.3 (2014): 445-468. [2] Liénard et al., (2018). SBDM/SfN. [3] Gutierrez et al., (2018). AINI. [4] Igarashi J., Moren J., Yoshimoto J., Doya K. Selective activation of columnar neural population by lateral inhibition in a realistic model of primary motor cortex, Abstract of 44th Annual Meeting of the Society for Neuroscience, (2014)

[5] Zhe Sun and Igarashi, Abstract of JNNS2018, (2018). [6] Gewaltig, Marc-Oliver, and Markus Diesmann. "Nest (neural simulation tool)." Scholarpedia 2.4 (2007): 1430. [7] Linssen, Charl et al. (2018). NEST 2.16.0. Zenodo. 10.5281/zenodo.1400175. [8] Miyazaki, Hiroyuki, et al. "Overview of the K computer system." Fujitsu Sci. Tech. J 48.3 (2012): 302-309. [9] Jordan J, et al. (2018) Extremely Scalable Spiking Neuronal Network Simulation Code: From Laptops to Exascale Computers. Front. Neuroinform. 12:2. doi: 10.3389/ fninf.2018.00002. [10] Georgopoulos, Apostolos P., Andrew B. Schwartz, and Ronald E. Kettner. "Neuronal population coding of movement direction." Science 233.4771 (1986): 1416-1419