

# Predicting neural activity by modelling the nuts and bolts of language trees

Martin Perez-Guevara<sup>1</sup>, Marc De Kamps<sup>2</sup> and Christophe Pallier<sup>1</sup>

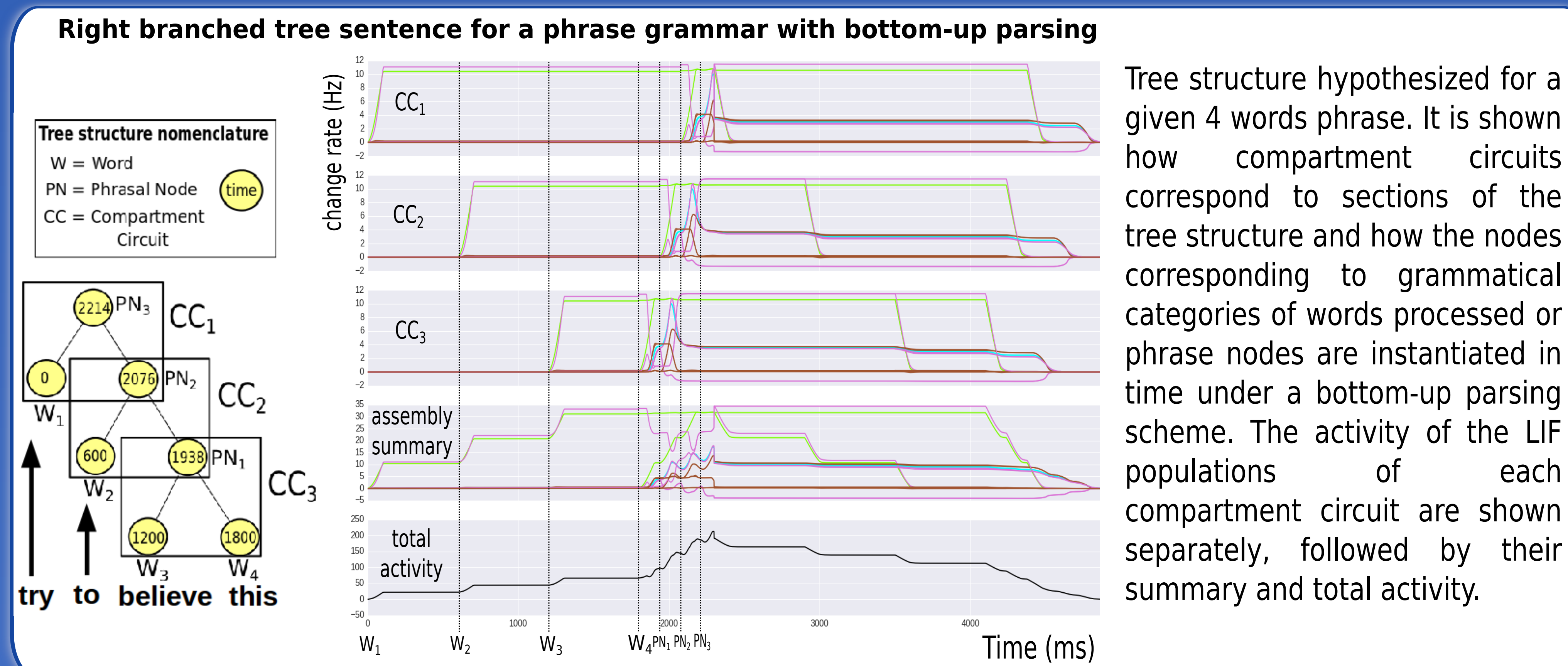
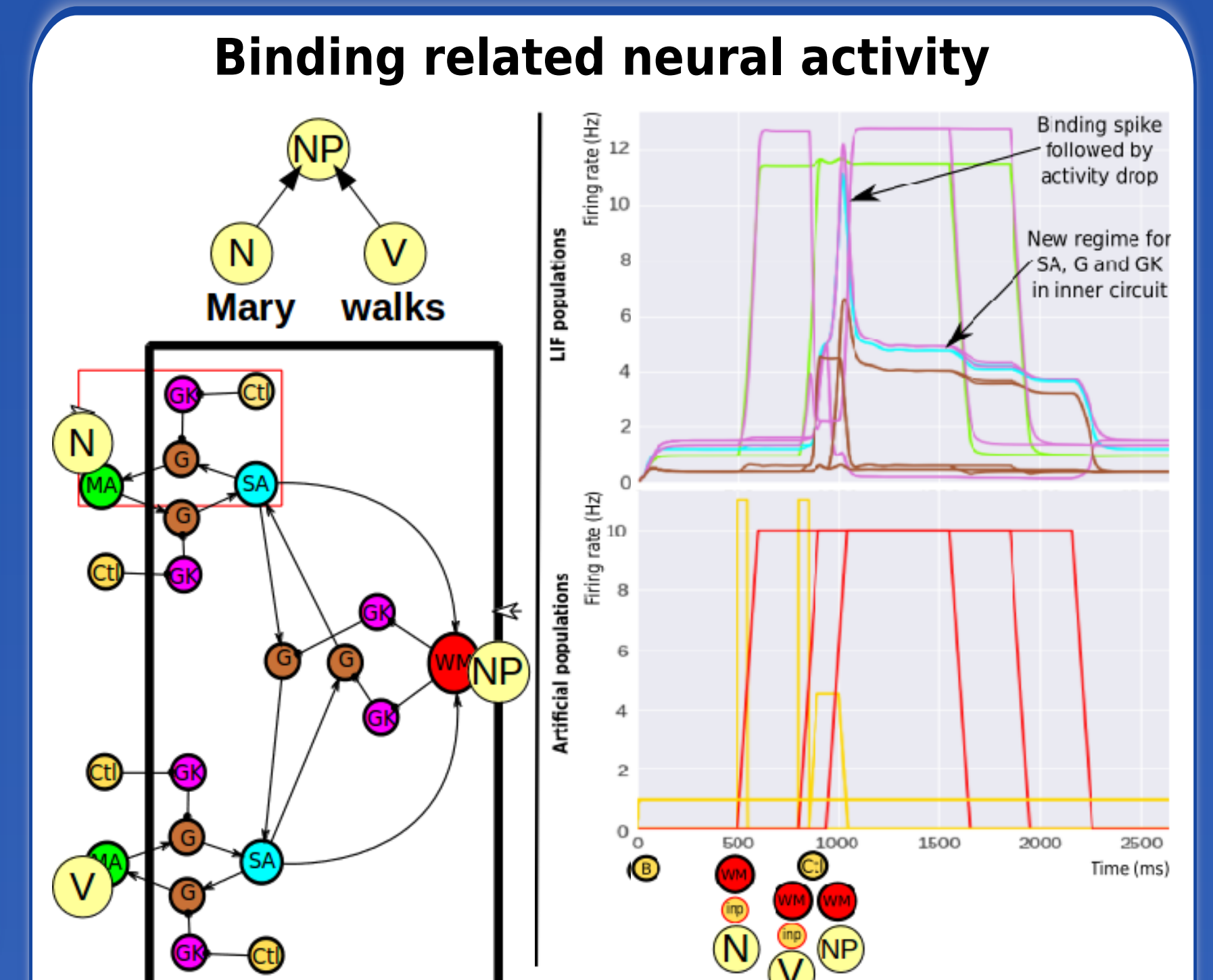
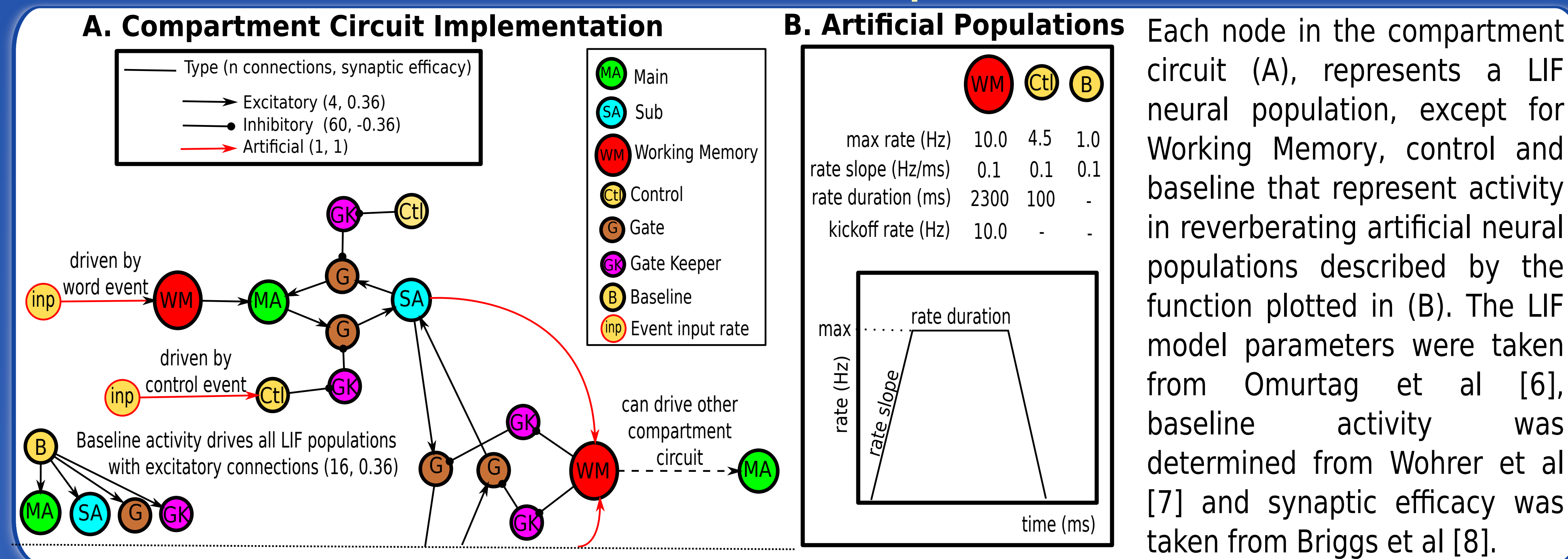
1. UNICOG INSERM/CEA/SAC/DSV/DRM/Neurospin center. Bât 145, Point Courier 156. F-91191 Gif-sur-Yvette Cedex FRANCE

2. Institute for Artificial Intelligence and Biological Systems. School of Computing. University of Leeds. LS2 9JT Leeds. UK

## Abstract

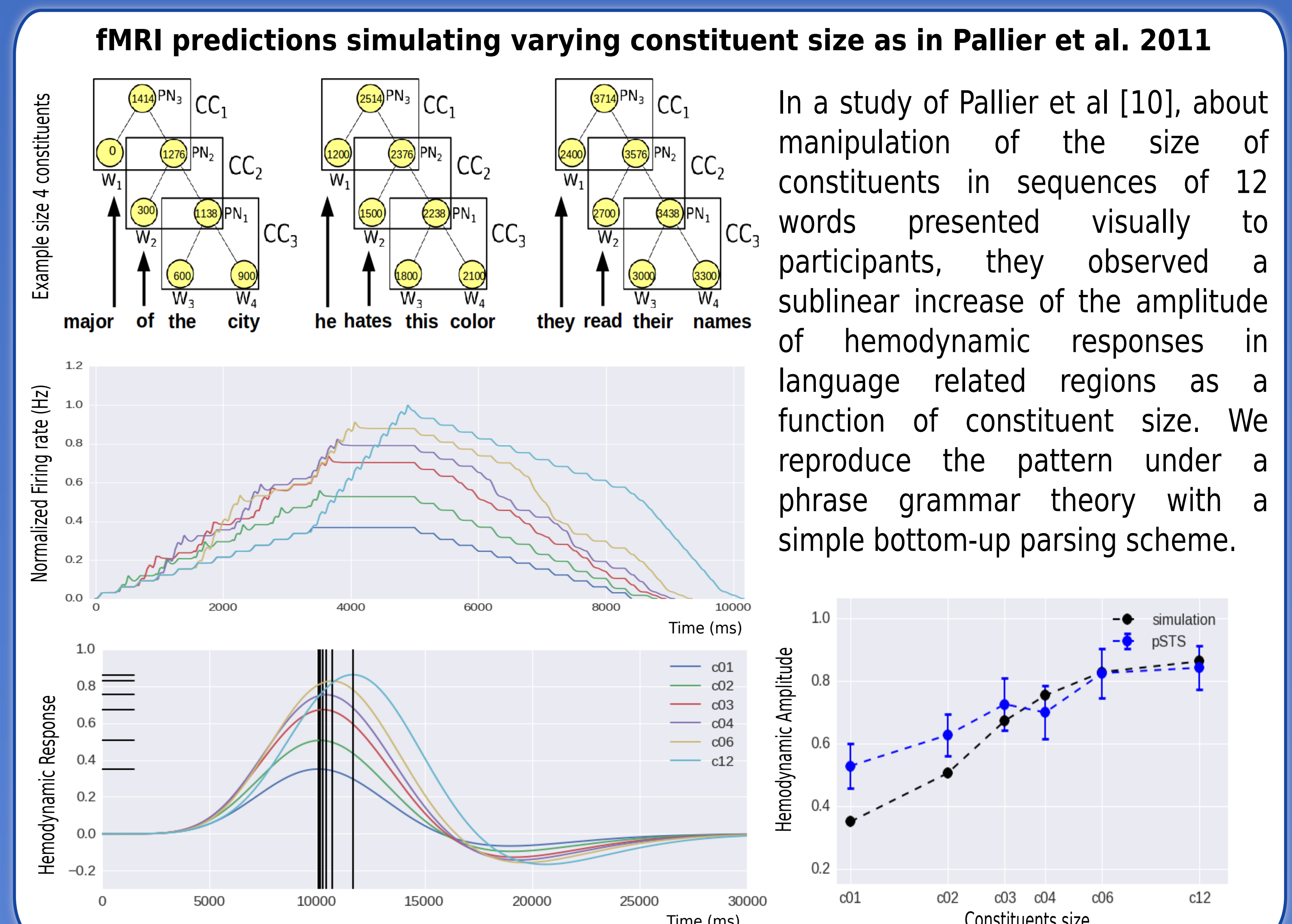
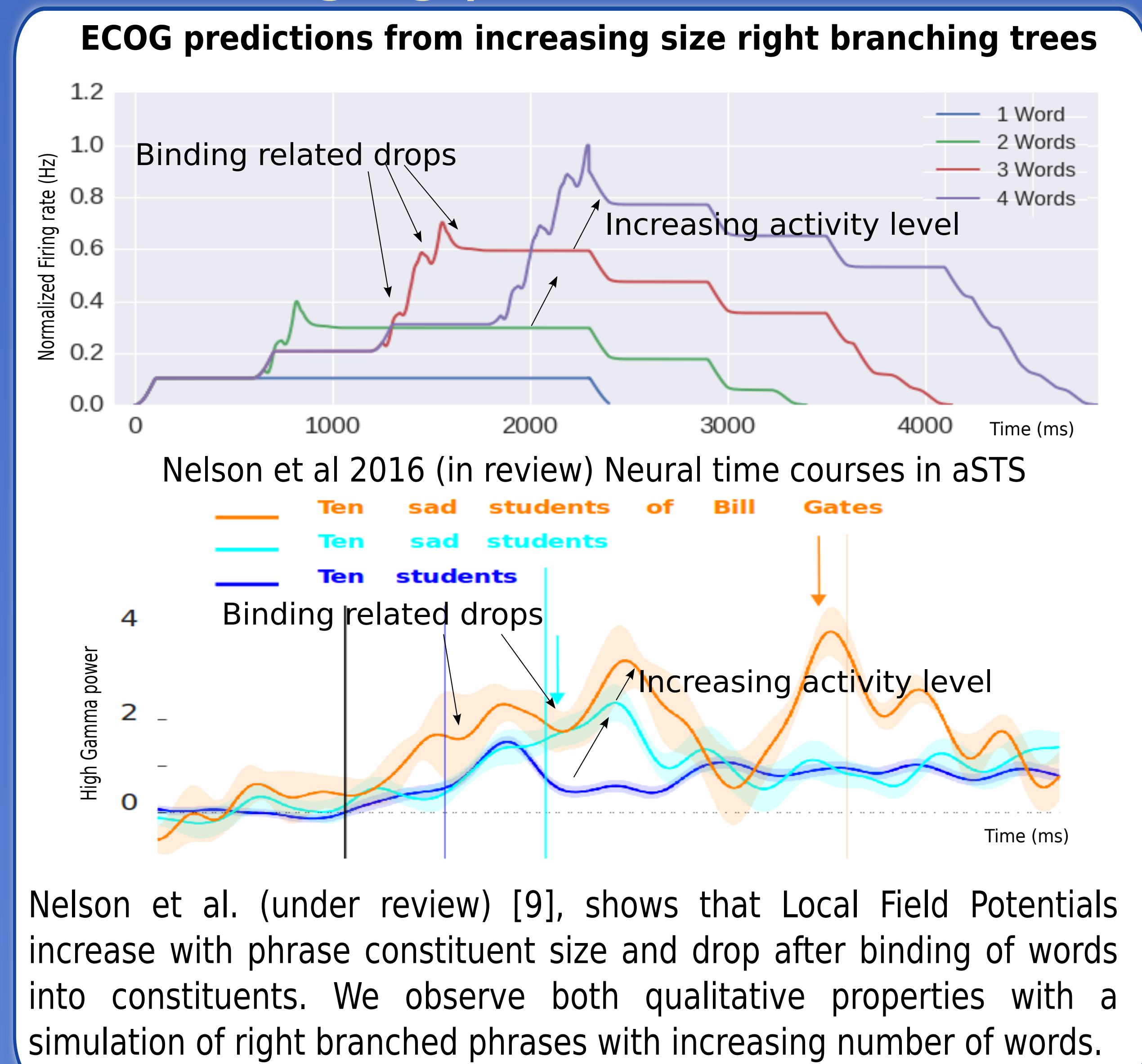
Few attempts have been made to model language in biological neural networks. The Neural Blackboard Architecture (NBA), proposed by Van der Velde and De Kamps [1] is one of them. It was designed to answer many challenges in the neural modeling of sentence processing, including the ones detailed by Jackendoff [2]. Here we expand on previous simulations of the Blackboard Architecture [3] on leaky-integrate-and-fire (LIF) populations with population density techniques [4] implemented in MIIND [5], to compare simulated time courses of neural activity associated to sentence representation and parsing with functional magnetic resonance data (fMRI) and intracranial recordings (electro-corticography; ECOG).

## Neural Blackboard Architecture Implementation



In case both MAs are active and Ctl activates to allow flow from MAs to their SAs, then binding will take place by activation of WM. Moreover other interesting non trivial dynamics take place. During the process of binding the reverberating activity of WM that inhibits the GKs connecting SAs creates a sudden burst of activity leading to a pronounced spike, this is due to the fact that SAs create a self excitatory loop, also elevating the activity of Gs and GKs between them. This burst of activity quickly drops back to a steady state that leaves the inner circuit in a level of activity far greater than in its original resting state, facilitating possible future communication between MAs.

## Neuroimaging predictions



**What next?**  
 We plan to extend the simulation to at least three parsing schemes: bottom-up, top-down and left corner. Then we will fit the nodes activity in the circuit and the timing of working memory nodes and controls to ECOG data in specific regions. The models fitted to specific regions will allow quantitative predictions on diverse fMRI dataset

## References

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