

# Sequence learning with the BCPNN rule

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## Sequence Learning

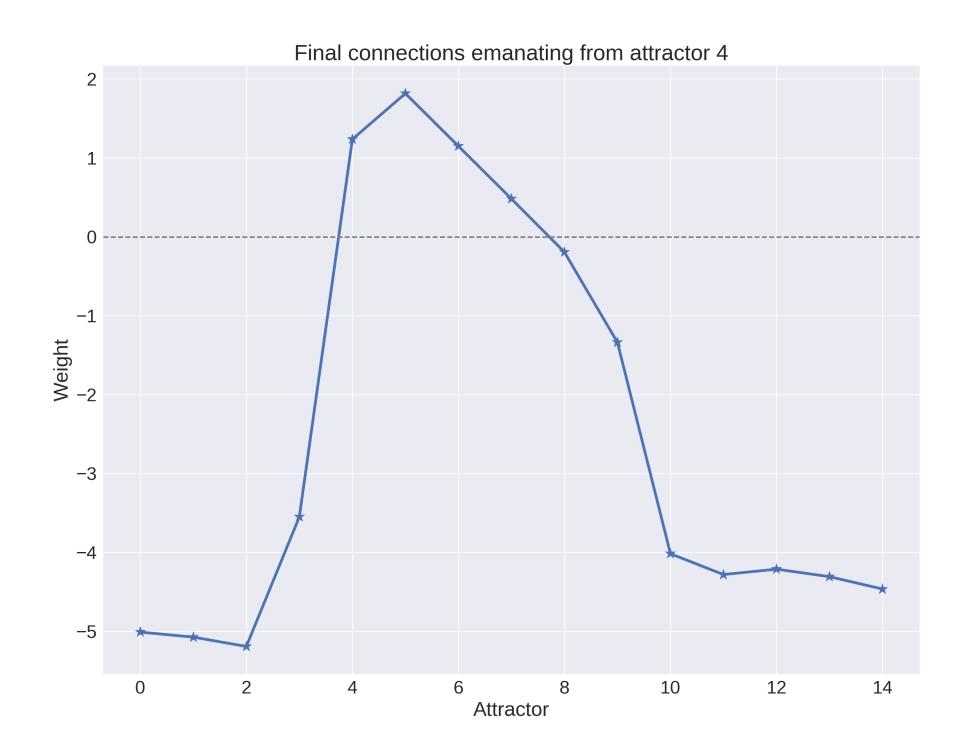
How can neocortical microcircuits encode sequences of activity? How can a stable sequential dynamics self-organize withing the bounds of the biological constrains? As early as 1950 Karl Lashley [3] advocated that the ability to sequence actions is the essential cognitive ability of human, how can we account for it? Since then we have found sequential population bursts in the activity related to the following behaviors:

- Motor
- Sensory
- Memory
- Decision Making

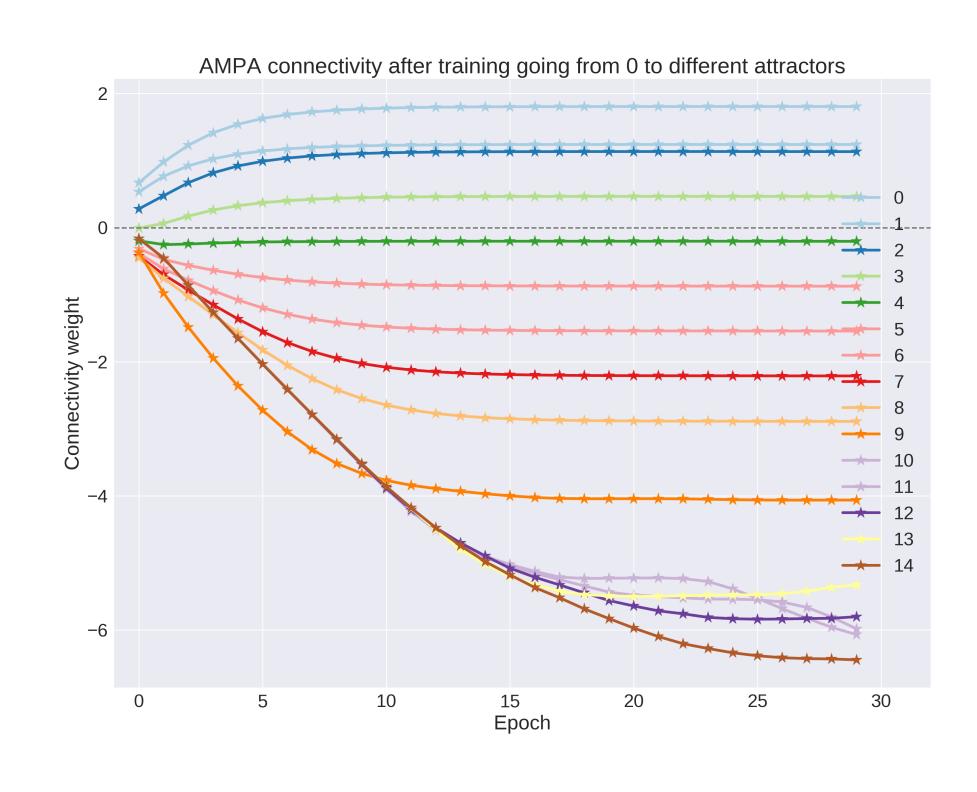
Here we propose a generic model that is a step ahead in solving the riddles above.

## Connectivity I

Here we have the outgoing connectivity from a particular attractor to the rest of them. The asymmetry of the curve implies the direction of the sequence.



# Stability across training

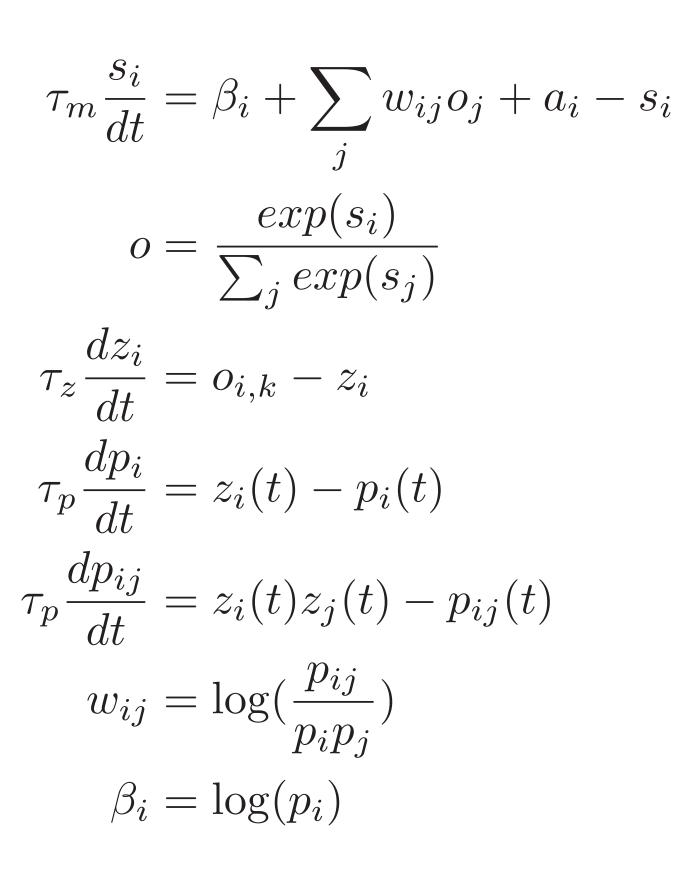


## References

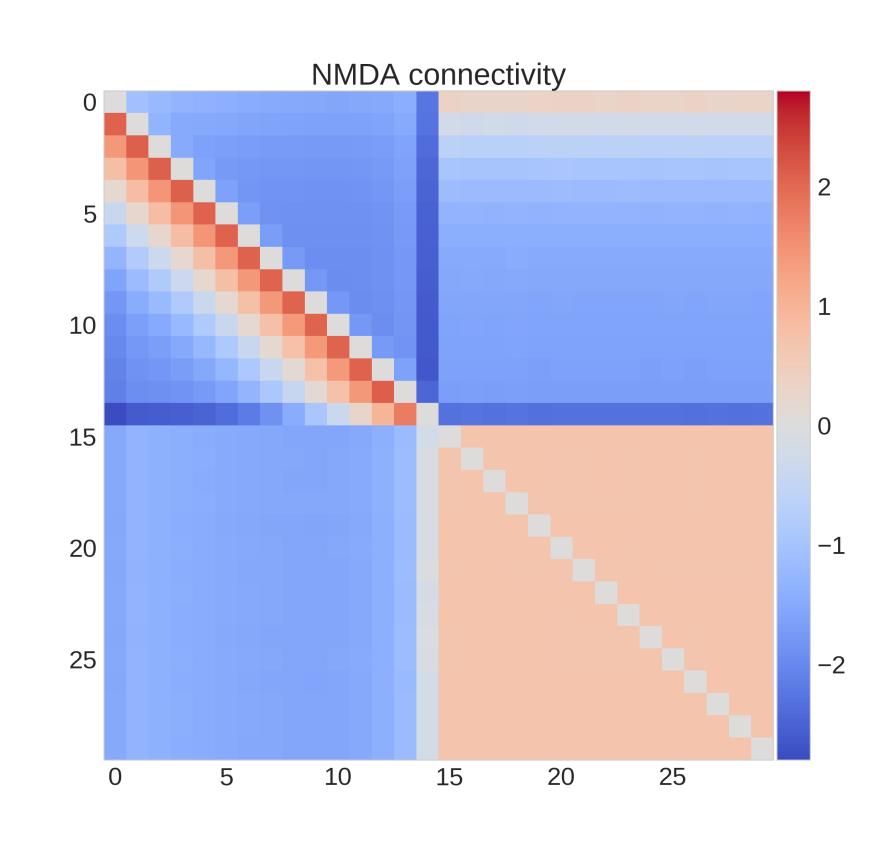
- [1] Tully, Philip J., Henrik Lindén, Matthias H. Hennig, and Anders Lansner. e1004954. *PLoS Comput Biol* 12, no. 5 (2016)
- [2] Sandberg, Anders, Anders Lansner, Karl Magnus Petersson, and Ekeberg 371(1):179-194 *Network: Computation in neural systems* 13, no. 2 (2002))
- [3] Lashley, Karl Spencer pp. 112-136 Cerebral mechanisms in behavior. 1951

#### The Model

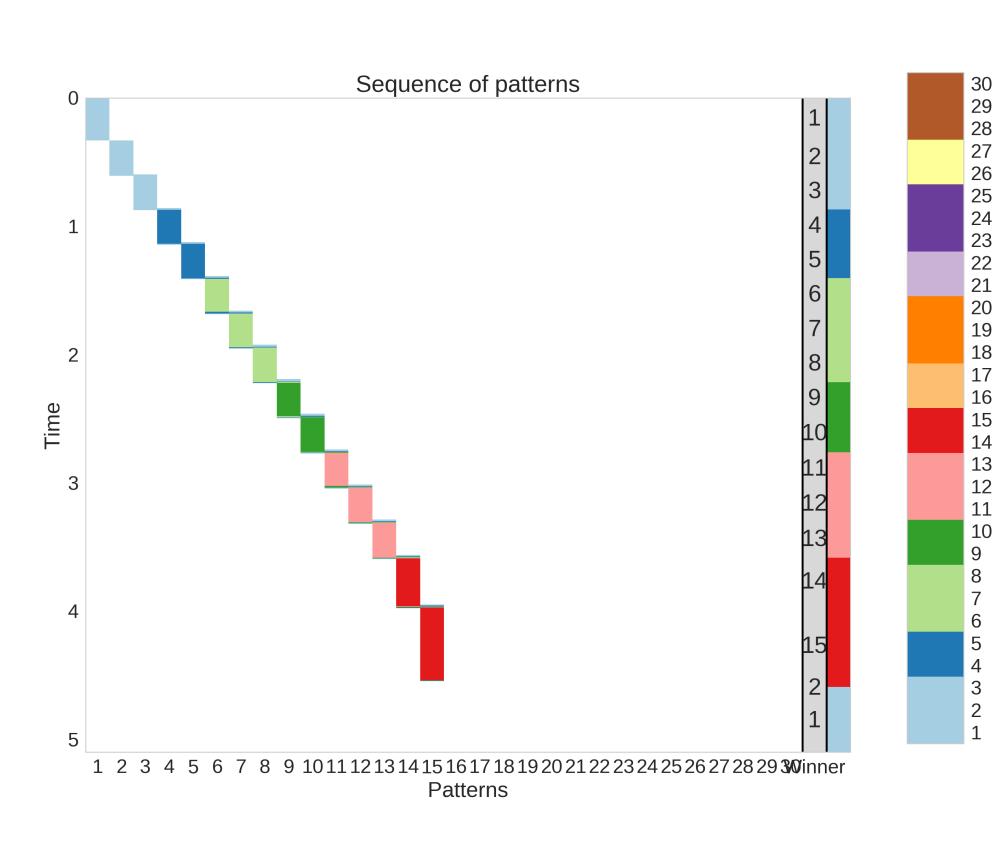
- Previous work has shown that the BCPNN rule can learn sequences in a spike based attractor model with modular structure [1].
- Using the firing-rate version of the model [2] with both fast (AMPA) and slow (NMDA) connectivity we study the capabilities of the system for pattern and sequence storage. [2].



Due to a local transition rule in absence of noise this model can recall a sequence of arbitrary length. Moreover, it can perform within certain limitations both sequence completion and sequence disambiguation.



Example of AMPA connectivity matrix and a successful recall of a sequence.



### Chains

- We stored more complicated sequences in order to probe how effective is our system at retrieving them.
- Two relevant parameters to parametrize the space of all the possible sequences are **overlap** and **overload**.

