**Introduction to the Neurotron**

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## *Abstract*

*This paper introduces the* Neurotron*, a basic neural computing unit to build hierarchical temporal sequence memory as a basis for machine intelligence. Sequence memory is capable of learning sequences, and to predict the next item, when part of a sequence is presented to the se­­quence memory.*

*The Neurotron is a quasi-digital computing unit, which means that the most of signal processing is purely digital, only the learning process is based on analog* permanences*, which represent the development state of the synapse models. The quasi-digital approach in combination with sparse representations of patterns lead to highly efficient algorithms with fast execution time and low memory need.*

*As the concept of the Neurotron has been strongly influenced by Numenta’s architecture of Hierarchical Temporal Memory (HTM), the underlying HTM key concepts have been incorporated into the Neurotron.*

*Finally, we benchmark the performance of a Neurotron based sequence memory layer with the performance of transformer based neural network alternatives, when both systems predict the continuation of text sequences from "Tiny Shakes­peare".*

# Introduction

The neuron model used in most artificial neural networks is known as the perceptron [1], proposed by Frank Rosenblatt in 1957. It has few synapses and no modelling of dendrites (figure 1/A). The perceptron attempts to model the proximal area of the neuron and represents de-facto a mapping of input information arriving at the proximal synapses to the neuron’s *output.* There is evidence that the proximal synapses, those closest to the soma (cell body), have a large effect on the likelihood of the cell generating an *action potential,* which is bursting along the neuron’s axon to synapses of other neurons of the network. It is important to realize that the perceptron model is a pure mapping model without any memory.

In contrast, an excitation of a distal (non-proximal) synapse has little effect at the soma (cell body). For this reason, it was hard to understand how the thousands of distal synapses can play a role for the cell’s responses [2].

Powered by advanced research methods [3], researchers could, however, unlock the secrets, that an activation of neighbored distal synapses within a short time interval leads to a local dendritic NMDA (N-methyl-D-aspartate) spike, which causes a depolarization of the soma. Such depolarization subsequently enables the neuron to fire earlier than neighbor neurons with comparable proximal excitation, which gives such neuron the benefit to inhibit neighbored neurons.

Since NMDA spikes have a much longer duration than the action potential spikes of soma and axon, this mechanism acts like memory based on a depolarized soma state. In addition to the (perceptron like) mapping functionality of the neuron, the availability of memory in neural models offers capabilities like *prediction* and *bursting*, introduced in [4].

Inspired by a better knowledge of distal dendrite functionality Jeff Hawkins and Subutai Ahmad developed their HTM approach [4,5]. Their approach is not targeting a detailed model of a biological model simulating the spatial and temporal behavior of electrical quantities and ion concentrations.

With today’s understanding biological neurons are very com­plex systems with complex sub-functionalities in the soma, axon, trunk and dendritic segments. Understanding of any given neuron requires the right level of abstraction [2]. In this sense the approach in [4] is a powerful conceptual abstraction of an enhanced neural layer functionality serving as an abstract building block for truly intelligent algorithms, which can run efficiently on computers. It incorporates the following concepts:

* *Quasi-digital processing*: most of the signals have binary values (0 or 1) with the exception of permanence values and synaptic thresholds (analog values in the interval [0,1]).
* Sparse representation of patterns
* Binary synaptic weights, depending on the develop­ment state (called permanence, an analog value ranging from 0 to 1) of the synapse. A synaptic weight equals 1 if the permanence exceeds a synaptic threshold, and 0 otherwise)
* *Minicolumn*: a kind of collaborative group of neurons with a mechanism for “*voluntary neuron*” selection, which take over a representation role in case of new patterns.
* *Predictive neuron states:* enable those neurons of a minicolumn to fire, which on base of the processed input pattern are expected to fire, while other neurons of the minicolumn stay quiet.
* *Bursting:* a state being activated in case of an unknown pattern with the purpose that all neurons of a minicolumn vote for role ownership to represent an unknown sequence part.
* *Local learning law:* the learning rule is Hebbian like, but in contrast to Hebbian learning where weight adjustment is proportional to the product of pre-synaptic input and neuron output, HTM neurons are only empowered for learning, if they were able to make a correct prediction.

There are some notable remarks: The quasi-digital nature of the algorithms in combination with sparse pattern representations allows very fast and efficient processing on digital computing hardware.

The capability of *bursting* in case of a new pattern presentation increases the probability of finding voluntary neurons to take over role ownership for co-representation of a new pattern and contributes to learning efficiency.

The HTM learning rule where synapses are only credited for learning, if the neuron made a correct prediction, provides strong learning focus on neurons with actual involved in the prediction process. It helps to avoid ruinous overwriting of well-established memory contents, which is a key challenge in the formulation of suitable synaptic plasticity hypothesis [6].

The HTM algorithm presented in [4] is motivated by new discoveries of biological neuron functionality, but the formulation of the algorithm is for a neuron-layer comprised of Minicolumns. Due to this approach a minicolumn is an array of “HTM-neurons”, leaving it fuzzy how the algorithm might be implemented in detail on a neurobiological level. Such approach is reasonable, when we accept, that the HTM-algorithm is an abstract model of an essential functionality, serving as a building block for truly intelligent behavior.

While in [4] the “molecular” computing unit is the *minicolumn*, we are going to introduce an “atomic” computing unit, the *Neurotron*, which can be used to build *Minicolumns* on the next higher level, but also to build other next level building blocks like a *Spatial Pooler*, another functional unit consulted in the HTM approach.

# The Neurotron

The *Neurotron* is a basic neural computing unit, serving as an atomic building block for HTM sequence memory (in the sense of [4]), for a *spatial pooler (*which is an autonomous learning map to maintain given sparsity by preserving semantics), and possibly for other higher-level functionality which has not yet been defined. The *Neuro­tron* is a well-defined abstraction of a pyramidal neuron, assigned with a well-defined algorithm deter­mining its computing process.

# References

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# Algorithm

Appendix

configure ***K,*** init ***X*** *0,* ***P*** *0*

repeat for *t = 0,1,2,3*

input ***u*** *t*

***Y*** *t = (****X*** *t + (1**-* ***1****m****max****(****X*** *t))° (****1****m* ***⋅ u*** *tT)*

***Q*** *t* = *σ(****D*** *t - η)* ° ***y*** *t(****K****)*

***X*** *t+1* = ([*||****Q****1 t ||1*, *||****Q****2 t ||1*, ... , *||****Q****n t||1*]*T ≥ θ)(m×n)*

***D*** *t+1* ***=*** ***D*** *t* ***+*** *(δ +****Q*** *t - δ -)* ° ***Y*** *t*