Neural mechanisms of information processing and transmission

Dissertation

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> by **Johannes Leugering** Osnabrück, January 2021

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Abstract

English

This (cumulative) dissertation is concerned with mechanisms and models of information processing and transmission by individual neurons and small neural assemblies. In this document, I first provide historical context for these ideas and highlight similarities and differences to related concepts from machine learning and neuromorphic engineering. With this background, I then discuss the four main themes of my work, namely dendritic filtering and delays, homeostatic plasticity and adaptation, rate-coding with spiking neurons, and spike-timing based alternatives to rate-coding. The content of this discussion is in large part derived from several of my own publications, but it has been extended and revised to provide a more accessible and broad explanation of the main ideas, as well as to show their inherent connections. I conclude that fundamental differences remain between our understanding of information processing and transmission in machine learning on the one hand and theoretical neuroscience on the other, which should provide a strong incentive for further interdisciplinary work on the domain boundaries between neuroscience, machine learning and neuromorphic engineering.

Deutsch

Diese (kumulative) Dissertation behandelt Mechanismen und Modelle der Informationsverarbeitung und -übertragung durch einzelne Neuronen sowie kleine neuronale Assemblies. In diesem Dokument stelle ich erst den historischen Kontext dieser Ideen dar, und zeige Gemeinsamkeiten und Unterschiede zu verwandten Ansätzen beim maschinellen Lernen und Neuromorphic Engineering auf. Vor diesem Hintergrund entwickele ich im Anschluss die vier Kernthemen meiner Arbeit: dendritische Filterung und Delays, homeostatische Plastizität und Adaption, Ratencodierung durch gepulste Neuronen, sowie spike-timing-basierte Alternativen zur Ratencodierung. Der Inhalt dieser Darstellung basiert im Wesentlichen auf mehreren eigenen Publikationen,er wurde allerdings weiterentwickelt und ergänzt um die Kernideen einfacher zugänglich zu machen, umfassender zu erklären und ihre inhaltlichen Verbindungen herauszustellen. Ich schließe die Diskussion mit der Schlussfolgerung ab, dass nach wie vor fundamentale Unterschiede in unserem Verständnis von Informationsverarbeitung und -übertragung bei maschinellem Lernen auf der einen, und theoretischen Neurowissenschaften auf der anderen Seite bestehen, die einen starken Anreiz für weitere interdisziplinäre Arbeiten im Grenzbereich zwischen Neurowissenschaften, maschinellem Lernen und Neuromorphic Engineering bieten sollten.

Preface

What this thesis is about

In this dissertation, I talk about several aspects of neural information processing that I believe to be very important for biological systems, but which are often overlooked or under-appreciated in models of (artificial) neurons. The topics of this thesis are therefore situated between the fields of theoretical neuroscience, machine learning and neuromorphic hardware. In order to explain the similarities and differences between these fields, the first two chapters offer a brief historical perspective of how they came to be (chapter 1), and how each of them understands and uses (artificial) neurons and networks today (chapter 2). Chapter 3 gives a brief tour of the field of *neuromorphic hardware* — my application domain for concepts from theoretical neuroscience. In each of the subsequent chapters, I then address one important aspect of neural computation, i.e. computing with dendritic filters and delays in chapter 4, improving computation with homeostatic plasticity in chapter 5, rate-coding with spiking neurons in chapter 6, and finally spike-timing and event-based computation in chapter 7.

What this thesis is not about

It is impossible for me to give a full account of all the topics related to neural information processing in one thesis, and even for the topics that I want to discuss, there is a large host of prior work that is better summarized elsewhere. For those who are interested in a deeper discussion of these topics as well as the historical context, I can highly recommend the books by Rosenblatt [1], Ashby [2], Maass and Bishop [3], Turing and Copeland [4], Eliasmith and Anderson [5], Laughlin [6], and Stone [7].

During my time in the *Neuroinformatics* lab, I also worked on other topics in machine learning and statistical modeling that I have decided to not incorporate into this thesis, since they are thematically disconnected. These include:

- Joint work with Olivera Stojanovic and the Robert-Koch-Institute on a Bayesian spatiotemporal model of the spread of infectious diseases [8].
- Joint work with Kristoffer Appel, among others, on the creation of the TRAUMSCHREIBER, a low-power mobile EOG/ECG/EMG/EEG device for polysomnography [9], as well as a software-stack to go with it and a block-course on wearable electronics.

- Joint work with Pascal Nieters, the German Meteorological Service and others on a model to predict precipitation using deep learning [10].
- The contents of a lecture series on *Ensemble methods for machine learning*, developed and held with Olivera Stojanovic in the summer term of 2017.
- The supervision of 16 Bachelor's and 7 Master's theses and several student projects on various topics.

What are the main scientific contributions within this thesis?

Most of the chapters in this thesis summarize ideas that are explored in depth in some corresponding publication(s). These contributions are the following three journal papers¹, two peer-reviewed conference papers, one book chapter, one patent², one non-peer-reviewed article and one conference poster, each of which is introduced in more detail in the corresponding chapter(s):

- 1. P. Nieters, **J. Leugering**, and G. Pipa, "Neuromorphic computation in multi-delay coupled models," *IBM Journal of Research and Development*, vol. 61, no. 2/3, 8:7–8:9, 1, 2017, ISSN: 0018-8646, 0018-8646. DOI: 10.1147/JRD.2017.2664698.
- 2. **J. Leugering** and G. Pipa, "A Unifying Framework of Synaptic and Intrinsic Plasticity in Neural Populations," *Neural Computation*, vol. 30, no. 4, pp. 945–986, 17, 2018, ISSN: 0899-7667. DOI: 10.1162/neco_a_01057.
- 3. **J. Leugering**, P. Nieters, and G. Pipa, "Event-based pattern detection in active dendrites," *bioRxiv*, 17, 2020. DOI: 10.1101/690792v3.
- 4. F. Meyer zu Driehausen, R. Busche, **J. Leugering**, and G. Pipa, "Bistable Perception in Conceptor Networks," in *Artificial Neural Networks and Machine Learning ICANN 2019: Workshop and Special Sessions*, 2019, ISBN: 978-3-030-30493-5. DOI: 10.1007/978-3-030-30493-5_3.
- 5. **J. Leugering**, "Making spiking neurons more succinct with multi-compartment models," in *Proceedings of the Neuro-Inspired Computational Elements Workshop*, 17, 2020, ISBN: 978-1-4503-7718-8. DOI: 10.1145/3381755.3381763.
- 6. **J. Leugering**, P. Nieters, and G. Pipa, "Computational Elements of Circuits," in *The Neocortex*, W. Singer, T. J. Sejnowski, and P. Rakic, eds., red. by J. Lupp, vol. 27, The MIT Press, 2019, pp. 195–209, ISBN: 978-0-262-04324-3. DOI: 10.7551/mitpress/12593.003.0016.
- 7. **J. Leugering**, P. Nieters, and G. Pipa, "Neuromorpher Musterdetektor und neuromorphe Schaltkreisanordnung hiermit," patent application DE 10 2019 134 044 A1.
- 8. **J. Leugering**, "A visit to the neuromorphic zoo," in *Embedded World Conference 2020 Proceedings*, 2020, ISBN: 978-3-645-50186-6.
- 9. P. Nieters, **J. Leugering**, and G. Pipa, "Neuromorphic Adaptive Filters for event detection, trained with a gradient free online learning rule," presented at the Machine Learning Summer School (MLSS-Africa 2019), 1, 2019.

- ¹ The first two are published in peerreviewed journals, the third has only been published as a pre-print and submitted for review.
- ² The patent has been filed and is currently pending.

But some content is also new, or at least not covered by my own publications. In particular, chapters 4 and 6 contain work that motivated me to pursue the ideas of chapter 7, but ultimately did not directly appear in any of my publications yet. I have therefore decided to include some of this additional content in appendices A and B, respectively, in the hope that it will help to keep the rest of the text concise.

The main body of this thesis is intended to provide a more accessible summary of these publications, to highlight the links between various topics, and to embed them into the bigger picture that has motivated my work. Since I have compiled this thesis over a long time-span, some of my views have also evolved, and I chose to introduce some of these older ideas in a new, hopefully clearer way. In some places, this has revealed some new interesting connections that were not explored in the original work.

Who should read this thesis?

Naturally, I hope the PhD committee will like this text, but I'm writing this with a different audience in mind, as well. Over the last few years, I have recognized more and more underappreciated similarities between theoretical neuroscience on the one hand, and engineering fields like electronics, signal processing and communication systems on the other — both in terms of what questions are asked ("How much information can be transmitted over this kind of channel? Is a pulse-based code effective? How can I realize this computation with these components?"), and in terms of the tools and models used to answer these questions (information theory, signal processing, dynamical systems, control theory, etc.). Similarly, I think that a lot of the early results of cybernetics and connectionism are often overlooked today; but reading papers and books by Minsky and Papert, Ashby, Turing, von Neumann, Rosenblatt and others shows how many of the seemingly revolutionary ideas of the last few years are already implied there! In particular at the fringes where these different fields meet, namely neuromorphic hardware, the close connection and shared history between theoretical neuroscience, computer science, machine learning and engineering becomes obvious. It is therefore not a coincidence that some of the most inspiring books I have read during my time as a PhD student are actually rooted in engineering disciplines. I have tried to follow their example, giving this text a bit of an engineering flavor.

Hopefully, the high-level descriptions given here make these results more accessible than the original publications (which were rather specifically written for other neuroscientists) and thus also prove useful for scientists and engineers from different fields, e.g. neuromorphic hardware designers or machine learning researchers, who are interested in abstract models of neural information processing mechanisms.

Johannes Leugering Nürnberg, 2020

Acknowledgments

During my time in the Prof. Pipa's Neuroinformatics group I was given an unusual amount of freedom to work on diverse topics that really interested me, most of which did not find a way into this thesis. But these apparently "unproductive" activities, too, played an important role for me, because they offered new perspectives and insights that came in handy in completely unforeseen circumstances. Today, I probably wouldn't be working on neuromorphic hardware, were it not for all the hours I had to invest into the design of the TRAUMSCHREIBER! This transition would not have been possible in an environment that singularly values publication metrics rather than curiosity.

I am therefore grateful to our dean of studies at the time, Prof. Achim Stephan, for fostering such a liberating studying environment, to my colleagues and forebears for creating a pleasant and inspiring working environment, to Anna Rushing-Jungeilges for ironing out any and all tensions, and to my supervisor Prof. Gordon "Entropy" Pipa for caring more about content than form and leaving me the freedom to find my own way. Last but not least, I'm thankful to my family and my girlfriend Olivera for being at my side through the ups and downs during these years!

− A Tale of Two Cities by Charles Dickens

The issues that give rise to excitement today seem much the same as those that were responsible for previous rounds of excitement. The issues that were then obscure remain obscure today because no one yet knows how to tell which of the present discoveries are fundamental and which are superficial.

- Perceptrons - Expanded Edition by Marvin Minsky and Seymour Papert

The computer and the brain

When photons hit the retina and cause a neuron to emit a spike, a physical effect becomes information. How is this information represented and processed by the neural network that constitutes the brain? How does it extract structure from its sensory inputs, and learn to adapt to its environment?

These are fundamental questions that have kept generations of scientists and philosophers busy. To answer them, we'll need to thoroughly understand the basic mechanisms at play in neural information processing. The objective of theoretical neuroscience is therefore to identify these principles, from the level of individual neurons and synapses all the way up to networks and brain areas, and to abstract them into theoretical (i.e. mathematical) models, which can be understood without all the overwhelming complexity that has developed over hundreds of millions of years of evolutionary history.

The sudden and rapid development of Deep Learning in the last couple of years might have given many people the impression that we have now finally "cracked the code" of how neural networks work, and that we are on the verge of solving the mystery of the brain and (artificial) intelligence. While this is certainly an exciting perspective, it's important not to forget, that similar claims have been made multiple times before, and the celebration has always turned out to be premature. For example, consider the following brutal assessment by Marvin Minsky and Seymour Papert from the year 1988 and mentally substitute the older term "Connectionism" with its modern counterpart "Deep Learning":

[...] [L]ittle of significance had changed since 1969, when the book was first published[...]. One reason why progress has been so slow in this field is that researchers unfamiliar with its history have continued to make many of the same mistakes that others have made before them. Some readers may be shocked to hear it said that little of significance has happened in this field. Have not perceptron-like networks — under the new name connectionism — become a major subject of discussion at gatherings of psychologists and computer scientists? Has not there been a "connectionist revolution?" Certainly yes, in that there is a great deal of interest and discussion. Possibly yes, in the sense that discoveries have been made that may, in time, turn out to be of fundamental importance. But certainly no, in that there has been little clear-cut change in the conceptual basis of the field. The issues that give rise to excitement today seem much the same as those that were responsible for previous rounds of excitement. The issues that were then obscure remain obscure today because no one yet knows how to tell which of the present discoveries are fundamental and which are superficial. Our position remains what it was when we wrote the book: We believe this realm of work to be immensely important and rich, but we expect its growth to require a degree of critical analysis that its more romantic advocates have always been reluctant to pursue – perhaps because the spirit of connectionism seems itself to go somewhat against the grain of analytic rigor. [20]

So what has changed since then? Are we about to make the same mistakes again? To get a better understanding of where we stand today, I'd like to start with a bit of historical background of the field(s).

1.1 The origins of computational (neuro-)science and machine learning

The medical study of the central nervous system can be traced back for more than three millennia [21], but the mechanism by which it operates has remained a mystery throughout most of this history. It was only after a series of remarkable scientific discoveries in the 19th century, notably the observation of so-called "animal electricity" [22], advances in microscopy and histology [23], the theory of evolution [24] and the popularization of cell theory [25] that the *neuron doctrine* took root [26] and modern scientific theories of the brain's function began to emerge. In 1943, in the middle of World War II, Warren McCulloch and Walter Pitts wrote a landmark paper *A Logical Calculus of Ideas Immanent in Nervous Activity* [27], in which they first proposed that networks of interconnected nerve cells could implement a powerful symbolic logic calculus. A sufficiently large network of neurons, endowed with the necessary periphery and memory, could therefore satisfy the conditions of a universal machine as outlined just seven years prior by Alan Turing [28]. They write [27]:

It is easily shown: first, that every net, if furnished with a tape, scanners connected to afferents, and suitable efferents to perform the necessary motor-operations, can compute only such numbers as can a Turing machine; second, that each of the latter numbers can be computed by such a net; and that nets with circles can be computed by such a net; and that nets with circles can compute, without scanners and a tape, some of the numbers the machine can, but no others, and not all of them. This is of interest as affording a psychological justification of the Turing definition of computability and its equivalents, Church's A-definability and Kleene's primitive recursiveness: if any number can be computed by an organism, it is computable by these definitions, and conversely.

This connection between the biological connectivity of neurons and an abstract, mathematical notion of *computation* created a theoretical foundation for the field of *computational neuroscience*. But the concept of computability did not merely provide a language for neuroscientists to *describe* the operation of the brain — it also made it conceivable to *simulate* neural behavior, and therefore *intelligent* behavior, on any appropriate universal machine. Turing became fascinated by this idea and in 1948 wrote a visionary publication entitled *Intelligent Machinery* [29] that today reads like a prescient outline for many subsequent developments in machine learning.¹

In 1949, Donald Hebb provided the first mechanistically plausible theory of (unsupervised) learning in neural networks, the now famous *Hebbian learning rule*, which in its most explicit form stated that "[w]hen one cell repeatedly assists in firing another, the axon of the first cell develops synaptic knobs (or enlarges them if they already exist) in contact with the soma of the second cell." [33] Thus the study of synaptic plasticity and learning in neural networks was born. Ross Ashby extended this view of self-organization as an essential property of the brain (and life in general), and ultimately proposed in his highly influential 1954 book *Design for a Brain* [2] the "Homeostat", a self-regulating machine, as an example of artificial life.

But since these algorithmic mechanisms could also be simulated by a Turing machine, it now seemed conceivable to simulate intelligent behavior, and, even more interestingly, learning. As Turing himself suggested in private correspondence to Ross Ashby, his *Automatic Computing Engine* (ACE) could be used to that end:

¹ In it, he discussed, for example, not just recurrently connected neural networks, but also proposed randomly initialized networks, which are then trained through reward and punishment, as a reasonable analogy for (some parts of) cortex — a view that anticipated some recently resurfaced ideas in the field of reservoir computing [30]. His B-Type networks furthermore bear some resemblance to gated recurrent units (GRUs) [31] which have been popularized recently by the LSTM model [32].

It would be quite possible for the machine to try out variations of behavior and accept or reject them in the manner you describe and I have been hoping to make the machine do this. [...] Thus, although the brain may in fact operate by changing its neuron circuits by the growth of axons and dendrites, we could nevertheless make a model, within the ACE, in which this possibility was allowed for, but in which the actual construction of the ACE did not alter, but only the remembered data, describing the mode of behavior applicable at any time. I feel that you would be well advised to take advantage of this principle, and do your experiments on the ACE, instead of building a special machine. I should be very glad to help you over this. [4]

In his later publications and talks, Turing pursued the idea of intelligent and learning machines (or rather software programs?) further, and in his 1950 essay Computing Machinery and Intelligence [4] presented the Imitation Game, today known as the Turing Test, which was meant to illustrate how sufficiently powerful computing machines could be considered to be as intelligent (or more so) than their human counterpart. He was quite outspoken about this conviction:

The original question, "Can machines think?" I believe to be too meaningless to deserve discussion. Nevertheless, I believe that at the end of the century the use of words and general educated opinion will have altered so much that one will be able to speak of machines thinking without expecting to be contradicted. I believe further that no useful purpose is served by concealing these beliefs.

These ideas set in motion the development of ever more powerful computer architectures, which in turn enabled generations of increasingly complex artificial neural network models and learning methods. This progress continues well into the present era of deep learning, which owes part of its success to the highly parallelized computing architectures that have emerged in recent decades. But it's worth keeping in mind that this transition from serial "von-Neumann" to parallel "non-von-Neumann" computer architectures is less of a revolutionary new idea than it is a return to the roots of computer science and neuromorphic hardware. In fact, John von Neumann himself had both studied models of biological systems and developed artificial computers like the ENIAC [34], and therefore understood the respective strengths and weaknesses of both approaches. But in a time when computers were still excessively large, expensive and memory a limited resource, he concluded in his tragically incomplete lecture notes The Computer and the Brain [35], from which I have stolen the title of this chapter:

That is, large and efficient natural automata are likely to be highly parallel, while large and efficient artificial automata will tend to be less so, and rather to be serial. [...] More specifically, not everything serial can be immediately paralleled – certain operations can only be performed after certain others, and not simultaneously with them (i.e. they must use the results of the latter). In such a case, the transition from a serial scheme to a parallel one may be impossible, or it may be possible but only concurrently with a change in the logical approach and organization of the procedure. Conversely, the desire to serialize a parallel procedure may impose new requirements on the automaton. Specifically, it will almost always create new memory requirements, since the results of the operations that are performed first must be stored while the operations that come after these are performed. Hence, the logical approach and structure in natural automata may be expected to differ widely from those in artificial automata.

Half a century later and with new materials and manufacturing processes at hand, neuromorphic hardware might finally be able to bridge this gap between natural and artificial automata.

1.2 From Perceptrons to Deep Neural Networks

The theoretical study of artificial neural networks as (simulated) learning machines continued, first under the label of *cybernetics*, then *connectionism*, into the modern field of *deep learning*. First, Frank Rosenblatt's original Perceptron [1] demonstrated that even a simple feed-forward network model, composed of one layer of (random) feature detectors followed by a single McCulloch-Pitts neuron, could solve many perceptual problems. Minsky and Papert [20] thoroughly analyzed the capabilities and limitations of this and similar kinds of network with their corresponding learning rules mathematically, and provided sound arguments why these networks were still impractical for many relevant problems. Despite the fact that they explicitly limited this critique to perceptrons with a single trainable layer ², this may have had an adverse impact on the amount of research and funding dedicated to the study of perceptrons at the time — a period that is sometimes referred to, a bit melodramatically, as the *first AI winter*.

Over the course of a few years, multi-layer perceptrons [36] gradually became more powerful and offered a first flavor of the abstract artificial neural networks (ANNs) still in use today: a hierarchy of (affine) linear combinations of inputs followed by non-linear transformations (in this case a step-function) with coefficients that could all be chosen or learned. Kunihiko Fukushima's Cognitron [37] made use of a deep hierarchy of neural network layers to solve a complex computer-vision problem, and could therefore be considered one of the first deep neural networks - although its weights were not optimized through end-to-end supervised learning, but partly derived from expert models, partly trained through a competitive form of unsupervized learning. A later extension, the neocognitron [38], even introduced shift-invariant features and could be considered an early form of convolutional neural network [39]. A series of proofs, e.g. in [40], finally extended the analysis of the computational power of perceptrons by Minsky and Papert to multi-layered networks and showed that different kinds of feed-forward neural networks are capable of uniformly approximating arbitrary real-valued functions. These proofs of universal function approximation capabilities didn't require particularly deep neural networks – a single hidden layer suffices in principle, so many practitioners questioned whether stacking many layers of neurons into deep neural networks would serve any practical "computational" purpose at all. In a curious repetition of history, Minsky and Papert reaffirmed their skepticism of neural networks in a practically unchanged revision of their influential Perceptron book [20], and to similar effect. They wrote:

The perceptron has shown itself worthy of study despite (and even because of!) its severe limitations. It has many features to attract attention: its linearity; its intriguing learning theorem; its clear paradigmatic simplicity as a kind of parallel computation. There is no reason to suppose that any of these virtues carry over to the many-layered version. Nevertheless, we consider it to be an important research problem to elucidate (or reject) our intuitive judgment that the extension is sterile. Perhaps some powerful convergence theorem will be discovered, or some profound reason for the failure to produce an interesting "learning theorem" for the multilayered machine will be found.

And in some sense their words became a self-fulfilling prophecy, with many researchers opting for the simpler to train and to use shallow network architectures (which had stiff competition from more sophisticated machine learning methods). The following "second AI winter" spelled the end of this *connectionist* era, even though the 'interesting "learning theorem" ', as Minsky and Papert had asked for, already existed unbeknownst to many in the form of the backpropagation algorithm, which was repeatedly re-discovered over the preceding and the following decades [41].

² Their critique also went well beyond the often mentioned inability of individual threshold-linear functions to solve the XOR problem, and included questions of learning speed, complexity of the required networks and even the information content required for specifying all coefficients. Despite significant advances in the field, all of these questions are still relevant today.

In parallel to these studies of feed-forward networks, recurrent neural network (RNN) models were developed, to endow networks with some form of memory and/or allow them to process temporally varying information. Jeffrey Elman introduced context units, i.e. hidden neurons that receive the network's previous outputs as additional inputs, into an otherwise feed-forward network, thus retaining previous activity in a form of "active" working memory [42]. John Hopfield took inspiration from Ising models [43], which were being developed in statistical physics to model the dynamics of the spins of electromagnetically coupled atoms, and provided an alternative account of memory, where each "memory" is associated with a stable fixed-point of a recurrently connected network's dynamics. These two novel perspectives on memory, a form of volatile memory realized by the networks momentary state and a persistent memory encoded in the network's connectivity, inextricably linked the concepts of memory and computation. In Elman's words:

In this account, memory is neither passive nor a separate subsystem. One cannot properly speak of a memory for sequences; that memory is inextricably bound up with the rest of the processing mechanism. [42]

1.3 The "Deep Learning Revolution"

After a phase of relative tranquility, (feed-forward) neural networks entered the spotlight for a third time after several convolutional neural network architectures [44–46] won several computer vision challenges, most famously the network nicknamed AlexNet by Krizhevsky, Sutskever, and Hinton, which severely out-performed the competing machine learning methods and thus proved the impressive capabilities of deep neural networks to a wider audience. The real reason for the breakthrough success of deep learning has since been debated intensely. But besides scientific reasons, which we shall look at in chapter 2, the success of deep learning can be attributed at least in part to the availability of "big data", i.e. large, unstructured datasets, which are ideally suited as training material for (deep) neural networks with their large number of parameters. Another factor is certainly the rapid improvement of computer hardware, graphic cards and dedicated accelerators, and a corresponding surge in optimized software tools for simulating large networks such as TensorFlow [47] and PyTorch [48], which enabled many researchers to develop and test countless variations of network architectures.

But the most compelling explanation, in my opinion, is neither better data, software or hardware, nor better performance of deep networks *per se.* Instead, deep learning owes much of its success to the surprising³ efficiency of the gradient-based optimization of neural networks. This only works because deep neural networks, despite being complex nonlinear models, can be easily differentiated with respect to all their parameters and optimized using *stochastic gradient descent*, also called *(error-)backpropagation* in Deep Learning [39]. The same optimization tools can also be applied to train recurrent networks in discrete time, by a procedure called *backpropagation through time* [49]. If a task can be expressed by a differentiable *loss* function, as it is often the case in machine learning problems, we can therefore use variations of the greedy (stochastic) gradient descent algorithm to iteratively reduce the loss. This offers a very simple interface towards applications, because it only requires specifying the goal of a task in terms of a differentiable loss function and providing some data — little domain knowledge required! So, deep learning really is all about *learning*, albeit in the narrow context of optimization, rather than biology or psychology.

³ We will see in chapter 2 why this is surprising.

1.4 The state of the field(s) today

Of course, how neural networks "learn" has been a critical question not just in machine learning, but also for theoretical/computational neuroscience. However, neuroscientific models of learning naturally have to work within the confines set by biologically plausible mechanisms, and are thus primarily concerned with questions about how *unsupervised* (possibly modulated by other factors) local plasticity mechanisms might interact, what kind of top-down error signals may be provided by the nervous system, or how they might be propagated. Machine learning, on the other hand, does not have to play by the same rules, and instead application-driven questions of reliability, speed, performance and efficient usage of limited labelled training data take center stage there.

Today, half a century after the conception of the multi-layer perceptron, deep neural networks are the dominant method throughout many application areas of machine learning, where they have displaced other approaches such as kernel methods and decision trees from the leaderboards of most competitions.

However, just as machine learning has advanced over the last decades, so has neuroscience, and the early models, such as the logic calculus proposed by McCulloch and Pitts, from which *artificial* neural networks were derived, no longer reflect our best current understanding of *biological* neural networks. Since the first full, mechanistic, dynamic model of a biological neuron by Hodgkin and Huxley [50], major technological and methodical improvements in experimental neuroscience have revealed more and more about the complex biological mechanisms at play, and our theoretical models of neurons and networks have changed accordingly. Chapters 4 to 7 are about some of these developments.

To make a long story short, what once started as a single research question — how neurons process information — has since split into three distinct areas of research: the study of how abstract (deep) artificial neural networks can be used to implement intelligent or learning machines, which today are major subfields of *artificial intelligence* and *machine learning*, the study of how biological neurons process information, which we now refer to as *computational neuroscience*, and the study of how similar artificial systems could be realized efficiently in hardware, now called *neuromorphic hardware*.

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Connectionists use learning rules in big networks of simple components — loosely inspired by nerves in a brain. Connectionists take pride in not understanding how a network solves a problem. [...] If you just have a single problem to solve, then fine, go ahead and use a neural network. But if you want to do science and understand how to choose architectures, or how to go to a new problem, you have to understand what different architectures can and cannot do.

- Marvin Minsky

Information processing in artificial neural networks

When we talk about information processing by *biological* neurons and networks, the best place to start is probably the highly simplified, biologically inspired model of *artificial* neural networks. But as we already saw in chapter 1, the development of these models for machine learning purposes has since become a science of its own, and has produced models that satisfy the constraints of computer hardware very well, but differ from their biological inspiration in substantial ways. These differences are what I want to explore throughout the rest of this thesis.

2.1 Terminology

Throughout this thesis, I will rely on a lot of terminology and basic concepts from theoretical neuroscience and machine learning. If you are already familiar with these, feel free to skip ahead to section 2.2; otherwise the following section offers a brief and high-level summary. For a more in-depth or rigorous definition, any current review or book concerned with neural networks should do, e.g. Goodfellow, Bengio, and Courville [39], Schmidhuber [41], and Strang [51]. In abstract terms, an artificial neural network (ANN) is a graph of nodes and directed, weighted edges. The nodes represent neurons, the edges represent synaptic connections between neurons. Each synapse connects its pre-synaptic neuron to its post-synaptic neuron. In turn, the synapse is one of its pre-synaptic neuron's outgoing, and one of its post-synaptic neuron's incoming connections. Each edge can be assigned a synaptic weight or efficacy, which either represents the gain that the synapse applies to its signal or the probability with which an individual spike (also called never impulse or action potential) is transmitted by the synapse. These weights of a network can be collected in a so-called weight matrix, also called the connectome of the network. In a machine learning context, neurons are typically grouped into an ordered list of k > 0 layers of neurons with typically no connections within each layer. For the most common kind in machine learning, feed-forward networks, the graph is acyclic2, meaning that its directed synapses define a partial ordering of the neurons from the input all the way "forward" to output neurons (hence the name). If the graph describing the network is cyclic, it is referred to as a recurrent network.3 Biological neural networks can also be stratified into layers, but here the definition is based on the neurons' cell-bodies' locations, rather than their connectivity, and hence cannot be used interchangeably.

Some neurons or layers (for feed-forward networks typically the first layer(s) in the hierarchy) receive external input signals and are called the *input neurons* or *layers*, whereas the neurons or layer(s) whose states are taken as the output of the network (for feed-forward

2

¹ The corresponding graph is *k*-partite, and the synaptic weights of such a network form a block-matrix.

² The weight matrix of such a network is (block) triangular without diagonal entries

³ Sometimes, a bit inconsistently, also groups of neurons with mutual connections that are embedded in an otherwise feed-forward structure are referred to as *recurrent layers* of a feed-forward network. The corresponding synaptic weights would then form a block-triangular matrix *with* super-diagonal entries in the diagonal blocks.

networks typically found higher up in the hierarchy) are called the *output neurons* or *layer(s)*. Neurons or layers that are neither input nor output are called *hidden*. Following a rather vague heuristic, a network is typically called a *deep neural network* if it is composed of particularly many layers and its weights are inferred from data — but exceptions to this rule exist, e.g. the Long-Short-Term-Memory [32] architecture is commonly counted among the deep neural network architectures despite it being a recurrent network. Conversely, networks with only very few hidden layers are sometimes called *shallow*. Since the concept of depth is not applicable at all in cyclic graphs, the distinction between *deep*, *shallow* and other networks, is therefore blurry in practice and often determined by the historical context.

The graph structure defines the *topology* of the network, and together with a choice of *neuron* and *synapse models* determines the *network architecture*. The precise behavior of the network depends on the values of its *parameters* such as *synaptic weights*, *gain* or *bias terms*.

Artificial neural networks are often used to approximate functions that are only partially specified by some *training dataset* of input-output pairs. The act of optimizing the network parameters for this task is then referred to as *training*. Using a trained neural network to map (previously unseen) inputs onto corresponding outputs is referred to as *inference*.

When simulating neural networks, time can be represented in different ways. For feed-forward networks in a Machine Learning context, time is typically quantized into *discrete update steps*, where the information is assumed to propagate through the entire network from the input layer to the output layer within one time-step. For recurrent networks in discrete time, one time-step resembles one simultaneous update to each neuron's output, and the time-varying outputs of the network are represented by *real-valued sequences*. In a computational neuroscience context, especially for analog and spiking networks, time is often represented *continuously*. The signals emitted by the neurons are then modeled as *real-valued functions*, *stochastic processes*, or *spike trains*. While it is in principle possible to mix these different kinds of signals within one network, most architectures assume signals to be either all continuous, discrete, or spiking.

2.2 Artificial neural networks are function approximators

For machine learning applications, the utility of neural networks derives solely from their remarkable ability to approximate highly non-linear functions. For example, if a neural network is used to distinguish images of cats from images of dogs, it is implicitly assumed that the network is in principle capable, once the correct parameters are chosen, to solve the problem — at least to a sufficiently good degree of approximation. In other words, we expect the network, a complex mathematical model with lots of parameters, to be able to approximate some (unknown) function that maps from the space of photographs to the discrete set {cat, dog}. This may seem trivial now, but proving that this holds for the kind of networks we use today turned out to be a quite challenging task. It has long been known that networks of simple McCulloch-Pitts neurons with a step-function as nonlinearity can represent any boolean function [27], but showing that this concept can be generalized to arbitrary continuous functions for neurons with other nonlinearities such as the rectified linear or the hyperbolic tangent function is anything but trivial. Cybenko [40] finally provided the very general proofs, that any continuous function on the n-dimensional unit cube or any indicator function of a finite measurable subset on the n-dimensional unit cube can be uniformly approximated by an expression of the form

$$y(x) = W^{out} f(W^{in} x + b),$$

with parameters W^{out} , W^{in} and b if f is continuous and discriminatory 4 (e.g. any sigmoid

⁴ A function is defined to be discriminatory if its integral w.r.t. a nonnegative measure is zero only if the measure is identically zero. The most popular choice of activation function in deep learning, the rectified linear activation functions of the form $ReLU(x) = \max(x, 0)$ for example is not discriminatory, since it is constant on the negative half-plain.

function would do). Hornik [52] showed that the same in fact holds true for any arbitrary function that is continuous, bounded and non-constant. 5 A more recent insight about the approximation capabilities of deep neural networks comes from the study of splines: a feed-forward network with rectified-linear activation functions — a very popular choice in machine learning — provides a parameter-efficient way to construct a continuous, piece-wise linear (CPWL) function or spline [53-55].

In simple terms this means, that both regression problems (i.e. approximating a continuous function) and classification problems (i.e. approximating an indicator function over some subset) can be solved arbitrarily well by a neural network with one or more hidden layers, if there are enough neurons in the hidden layers.

The addition of recurrent connections allows these models to retain information for extended periods of time in the network's state, endowing the system with a form of memory. This was observed by Elman [42], who argued that by including context units, which feed back the system's current output as an additional input, the network can "memorize" and distinguish sequences of input. This idea has been generalized by Funahashi and Nakamura [56], proving that in fact any continuous function of time F(t), including of course the state of an autonomous dynamical system, can be uniformly approximated arbitrary well on a (finite) time-interval by some recurrent neural network with sufficiently many hidden neurons. The argument was further extended by Chow and Li [57] to non-autonomous, i.e. input-driven, systems under few generous additional conditions.

The function/system approximation paradigm therefore also covers recurrent artificial neural networks. Similar arguments can be made for both feed-forward and recurrent spiking neural networks (see chapter 6), which are universal computers as well, given sufficiently many neurons [58].

To summarize the summary, these existence proofs show that feed-forward and recurrent artificial neural networks, spiking or not, can in principle (i.e. if they are sufficiently large, which depends on the task at hand) approximate both instantaneous functions of the input and input-driven dynamical systems arbitrarily well. This makes artificial neural networks extremely powerful tools for machine learning and, as we shall see, neuromorphic hardware.

A bird's eye view of artificial neural networks

The existence proofs above only promise that neural networks are in principle capable of approximating functions, but give no indications of how such networks can be constructed. A large part of neural network research since the Connectionists' era has therefore been concerned with finding different architectures that are good at solving different tasks, as well as smart, systematic ways to combine them. As Minsky and Papert [20] already wrote:

Different kinds of networks lend themselves best to different kinds of representations and to different sorts of generalizations. [...] This is why we maintain that the scientific future of connectionism is tied not to the search for some single, universal scheme to solve all problems at once but to the evolution of a many-faceted technology of "brain design" that encompasses good technical theories about the analysis of learning procedures, of useful architectures, and of organizational principles to use when assembling those components into larger systems.

Today, there are at least four major conceptual frameworks that provide tools for constructing useful, large neural networks. We'll have a look at each of them in the following, and see why neither of them is suitable for studying information processing on the lower level of individual biological neurons that I am interested in.

5 This proof still does not directly cover the ReLU function, which is not bounded, but since a linear combination of two ReLUs can be used to construct a bounded function that satisfied the requirements, this indirectly proves the same power for ReLU functions as well [51].

2.3.1 Deep Learning

Currently, the most popular framework by far for constructing large artificial neural networks is *deep learning*. As the name implies, it typically involves stacking many layers of neurons into deep hierarchies, and training the entire network through gradient-based optimization methods.

Despite its apparent success, this approach was (and still is) met with some skepticism: Why would stacking more than one hidden layer have any *qualitative* benefit for the network's computational power, if the proofs above show that a single hidden layer is apparently enough? Also, given their typically rather large number of parameters, the numerics of optimizing these networks alone are a serious challenge that requires capable linear algebra tools [47]. And more fundamentally, according to basic results from statistical learning theory [59], these networks should either require an infeasibly large amount of training data and/or strong regularization of the parameters, ⁶ or generalize poorly to new data. This is apparent from the large amount of information content that the network weights store, as already pointed out by Minsky and Papert [20]⁷. Another concern from the perspective of nonlinear optimization of such large models should be local minimal of the loss function — i.e. mediocre solutions of the problem, which are difficult to improve for greedy gradient based methods and would require more sophisticated optimization methods.

Nevertheless, the training of large deep neural networks with gradient methods appears to stubbornly defy these intuitions, and has been surprisingly effective in practice. One argument in favor of building such deeper hierarchies is, that this "compositionality" allows for a much more efficient approximation of high-dimensional functions by comparatively few neurons. For the example of networks with rectified-linear activation functions, which correspond to piece-wise linear splines as we already saw above, the number of neurons required to approximate a given function can in some cases be exponentially reduced as we increase the number of layers in the network!8 Another argument is based on the surprising observation that the higher number of parameters in deep neural networks appears to make the optimization easier! Even deep neural networks that have parameters in excess of training data samples, make no explicit use of regularization, and perfectly interpolate all the training data have been shown to generalize well to previously unseen data [62]. This point has been debated a lot, but the answer might come from two rather unexpected directions: First, in the high dimensional parameter-space of neural networks, local minima of the loss function are comparatively less common in relation to e.g. saddle points, which pose much less of a problem for gradient methods [63]. Second, it appears that the commonly used stochastic gradient descent algorithm itself has an implicit regularizing effect on the learned coefficients, and leads the network coefficients towards a minimum norm solution [62, 64, 65]. A different, albeit controversial, explanation of the same phenomenon can be made in terms of the information bottleneck principle [66], which we'll return to in a different context in chapter 5. The lottery ticket hypothesis [67] goes one step further and suggests, that much smaller sub-networks (the winning tickets) that perform equally well can typically be extracted from deep neural networks by eliminating most of the synapses, neurons or even layers. Sometimes, the learned weights of entire layers within a deep neural network can be completely irrelevant for the task, as can be demonstrated by randomizing them with only negligible impact on performance [68]. However, finding such a sparser or shallower structure directly, e.g. by optimizing a smaller neural network to begin with, fails in practice. This, the argument goes, is because the parameter-rich deep neural networks offer a large search space, or a scaffold of sorts, in which the much smaller networks (which are sufficient to solve the task) can be constructed efficiently using gradient based optimization methods.

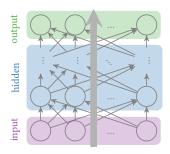


Figure 2.1. A feed-forward network with input, hidden, and output layer(s). Information flows only from input towards output layers.

⁶ There are a lot of explicit and heuristic options for regularization in deep learning, such as Nesterov optimizers, dropout, weight decay, data augmentation and early stopping [39].

⁷ For example, the MegatronLM GPT2 model with its 345 million parameters comes at a compressed size of about 650MB [60], its bigger cousin with 8.3 billion parameters [61] is correspondingly larger.

⁸ To be precise: there are functions f for every integer k such that a deep network with k^3 neurons distributed over k^2 layers can approximate f at least as well as any shallower network with ≤ k layers and $\frac{1}{2}k^{k+1} - 1$ or more neurons! [54].

Therefore, gradient descent and deep learning are so intimately coupled, that one of the most prominent figures in the field, Yann LeCun, even advocated for altogether switching to the more accurate name "differentiable programming" instead of "deep learning", since the optimization of neural networks by gradient-based methods is the defining feature, rather than the depth of the networks, as the term "deep learning" would imply [69].

An increasingly important practical consideration of *deep learning* research today is also the search of parameter efficient network architectures for different domains, i.e. by trying to reduce the number of neurons or synapses [70, 71], efficiently reusing synaptic weights [72] or even lowering the precision of weights [73, 74]. Over time, deep neural network architectures have thus evolved, much like their biological counterparts, and in the process incorporated a lot of domain-specific optimizations and inductive biases that make them well suited to specific tasks, but also pose a risk as they are typically not well understood (see also the note below).

Note: *Nature* or *nurture* in deep learning?

One important question in neuroscience is, how much of behavior is genetically predetermined (i.e. by nature), and how much is learned (i.e. by nurture). A very similar question could in fact be asked for deep learning! On first sight, it may seem obvious that all network coefficients are learned from data, and the amount of domain expertise explicitly build into the network is also minimal compared to early work by Connectionists, hence learning should play the major part. However, this is not true if we look at the discipline of deep learning as a whole: Deep neural networks are typically evaluated and compared to each other by training and testing them on the same benchmark tasks with the same datasets. Naturally, the more effective solutions are more likely to be picked up and developed further. What is kept and modified from one implementation to the next is typically not the weights (although pre-trained networks exist), but the *network architecture* - i.e. the types and numbers of layers, activation functions, etc. But, if we now evaluate the next generation of networks on the same dataset(s), we have committed a cardinal sin of statistical modeling by peeking at the test dataset (i.e. starting from an architecture, a meta-parameter of the model, selected on the same test dataset) before training the model! We could avoid this by testing each model on entirely new testing data, but even in that case merely choosing the best-performing model as a starting-point effectively bakes domain "knowledge" (or rather, information) into the network architecture. The result of this dynamic, which happens not on the level of the individual researcher but across the entire field, could be viewed as an evolutionary algorithm that produces ever more powerful, complex and specialized network architectures for these benchmarks. This is partially deliberate, since we want to find better and more useful models, partially unwittingly, since we might underestimate how much of bias this can introduce. A funny and slightly worrying example of this are the weight agnostic neural networks by Gaier and Ha [75], which show that some network architectures are so specifically designed for a certain task that they can solve it reasonably well even if all of their coefficients are fully randomized – reducing the role of *learning* in deep learning ad absurdum!

In a nutshell, the deep learning framework uses large, mostly feed-forward artificial neural networks with weights optimized by gradient descent to approximate functions that minimize some differentiable measure of "loss" or "cost". Its surprising effectiveness is at least in part due to the quantitative improvements that this optimization approach can offer, such as the reduction of local minima and implicit regularization, but this can come at the expense of over-complete networks, where many neurons or synapses may be entirely redundant. The choice of network architectures plays an important part, as well, but is often based on heuristics and incremental improvements of prior work. Despite interesting attempts to address the issue [76, 77], this "black-box" character of deep neural networks remains a contentious topic of debate [78] to date.

2.3.2 Attractor Networks

Attractor networks offer a completely different explanation of information processing by artificial neural networks. As we have seen above, recurrent neural networks can, if large enough, implement arbitrary dynamical systems. The computation realized by a neural network could therefore also be attributed to the long-term dynamics of the network as a whole, such as convergence of the network's state to some fixed-point, rather than the instantaneous output of a network in response to input. This insight underlies the model proposed by Hopfield [43]. He studied the linearized dynamics of recurrently connected neural networks and showed, that under certain constraints on the connectivity between neurons (e.g. symmetric and bounded connection weights), the network state must —without external input- converge to one of possibly many stable equilibria, depending on the network's initial state. Starting from a perturbed state that is similar but not identical to one of these stable attractors brings the network activity towards the attractor, and thereby "restores" the unperturbed stable state. By converging to a fixed-point, the network thus retrieves and reconstructs a perfect "memory" from an incomplete or corrupted version of that memory — a form of memory that he called *content addressable memory*. Rather than a function, the network thus implements an iterative algorithm! Since each memory is implemented by a fixed-point of the network dynamics, it imposes a constraint on the weight matrix of the network, and the total number of memories that can be stored this ways (and the robustness with which each memory can be recovered) depends on the size of the network. While Hopfield networks themselves are barely used in machine learning, they have been very influential in theoretical neuroscience. Similar ideas can be found today also in reservoir computing approaches.

2.3.3 Reservoir Computing and Conceptors

Reservoir computing (also known as echo state networks or liquid state machines) [30] represents a more radical approach to using recurrent network dynamics. It uses the transient dynamics of randomly connected recurrent neural networks, which are continuously driven by the network's inputs. The time-varying state of the networks' neurons thus provides a random, non-linear embedding of the network's input history into a high-dimensional vector space. For this embedding to be useful, it only needs to satisfy a few basic and easy to satisfy conditions [58]. The recurrent network is then called a reservoir computer, and a linear transformation of the network's high-dimensional state vector can be used to approximate a time-varying target signal, i.e. some function of the network's recent input history. This transformation is also called the network's linear readout, and its coefficients are the only parameters of the reservoir computer that are optimized for a specific task. 9 The conceptor framework [80] combines this approach with ideas from attractor networks. By controlling the attractors and limit-cycle dynamics of the reservoir (like a Hopfield network), a conceptor network can shape the dynamics to produce or resonate with certain stable time-varying patterns of activity, which a linear readout (like in a reservoir computer) can shape into a desired time-varying output signal. For an example of a network architecture built from such conceptor networks, see contribution 1.

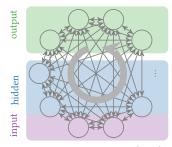


Figure 2.2. A recurrent network with input, hidden, and output neurons. Information is actively maintained in the network by recurrent activity.

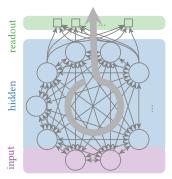


Figure 2.3. A reservoir computer with input and hidden neurons and a linear readout layer. Information is actively maintained in the network by recurrent activity and accessed through the readout layer.

⁹ This combination of a random high-dimensional, non-linear feature-expansion with a simple linear regression model is reminiscent of the *kernel trick* popularized by support vector machines [79] and the original Perceptron [1].

Contribution 1: Bistable Perception in Conceptor Networks

This conference paper explores, how a hierarchy of conceptor networks, which can act as generative models for time-series signals, can be used to actively suppress noise and minimize prediction errors. The idea of such a hierarchical predictive coding scheme is in line with biological observations and provides an appealing model of perception. When presented with ambiguous superposition of two stimuli, this architecture reproduces the well known psychological phenomenon of bi-stable perception, where either of the two pure stimuli is perceived in isolation for a period of time, before the percept switches to the other. It matches empirical results with surprising fidelity, including the distribution of the time-spans for which either of the stimuli is perceived! This paper extends ideas developed within Felix Meyer zu Driehausen's thesis, which I had the pleasure of supervising. Felix and Rüdiger Busche subsequently turned it into a viable model and a nice conference paper, for which they deserve all the credit.

Reference:

F. Meyer zu Driehausen, R. Busche, J. Leugering, and G. Pipa, "Bistable Perception in Conceptor Networks," in Artificial Neural Networks and Machine Learning – ICANN 2019: Workshop and Special Sessions, 2019, ISBN: 978-3-030-30493-5. DOI: 10.1007 /978-3-030-30493-5_3.

The Neural Engineering Framework

The Neural Engineering Framework (NEF) incorporates ideas from both deep learning and reservoir computing, but it offers a rather different perspective on how to construct large neural networks. A great in-depth discussion of this approach can be found in [5]. It focuses on the design of modular neural network architectures from smaller building blocks with welldefined function, rather than end-to-end optimization of network coefficients from data. Each of these building blocks is itself a randomly connected feed-forward network or reservoir computer, whose activation encodes (or represents) some variable, typically low-dimensional. Connections between the blocks are optimized to implement functions (or transformations) of these variables. By stacking and connecting many such modules together, large and complex networks with well-defined behavior can thus be constructed. This approach is very general and works well for different kinds of neuron models, from the simple linear-nonlinear neurons to more complex dynamic neuron models like leaky integrate-and-fire neurons, because it doesn't rely on the backpropagation of gradient information through deeply nested hierarchies of neural network layers. Figure 2.4 illustrates this modular approach.

A practical limitation of this approach is that it requires the problem to be analytically decomposed into sub-problems that are each simple enough to be efficiently solved by a single module, and we need either an analytical solution of each of these sub-problems or sufficient data to train them. But when this is possible, the neural engineering framework provides a straight-forward way to compose smaller feed-forward and recurrent neural networks into very large networks with well-understood, highly complex functionality.¹⁰

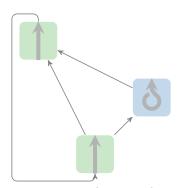


Figure 2.4. A network constructed out of mutually connected feed-forward and reservoir modules according to the neural engineering framework.

 $^{\rm 10}$ The Nengo software tool [81] offers a very fast and entertaining way to construct neural networks using the NEF.

2.4 Where artificial and biological neural networks diverge

We now looked at four popular frameworks for constructing large artificial neural networks, each of which offers a different interpretation of 'neural information processing'. But do these machine learning frameworks explain information processing in the brain? I believe not, because despite their shared history, terminology and many conceptual connections, the artificial neurons and networks shown above are extreme simplifications, idealizations and modifications of their biological inspirations. To name just of a few of the fundamental differences that have emerged between the artificial neural networks models form machine learning and those from theoretical neuroscience:

- 1. Artificial neural network models employ a single kind of neuron and a single kind of synapse, the behavior of which is fully parameterized by a single bias and weight coefficient each. In the brain, however, neurons are morphologically and behaviorally so diverse that even a classification into distinct classes can be challenging. But with around 60 different types of neurons in the retina alone, the number of different classes of neurons in the human brain may well be in the hundreds [82] not to even speak of other cell types like glia cells, which are entirely absent from artificial neural network models. Similarly for synapses, whose inhibitory and excitatory effects are not adequately expressed by positive and negative synaptic weights alone [83]. Also, electrical gap junctions, which directly and bi-directionally couple neurons and play an important role in some biological neural networks [84], are not modeled at all in artificial neural networks. In addition to these neural structures, the vast variety of neurotransmitters or -modulators is ignored in artificial neural networks, despite their critical influence on behavior in biology [85].
- 2. A lot of biological structures are genetically pre-determined, rather than learned in an end-to-end fashion as is popular in machine learning. For instance, in simple multicellular organisms such as *C. elegans* [86] or tadpole larvae [87], neurons are assembled in very specific, genetically determined patterns; so specific in fact, that the entire connectomes can be described on the level of individual neurons. Neurons can also form specific *motives* [88] of mutual connectivity patterns, that are reproduced many times throughout the nervous system, or larger homogenous groups of strongly coupled neurons called *cell assemblies* [89], or even larger structures called *canonical microcircuits*, possibly arranged in a columnar structure called *cortical (micro-)columns* but this latter point remains a contentious topic [90]. These innate, highly specific structures don't fit any of our machine learning paradigms above, yet they "compute" nonetheless.
 - This is particularly relevant if one considers the fact that neurons and networks must have *evolved* from such and even simpler structures. Studies of the evolutionary origins of neurons and nervous systems point to simple sensory cells arranged in homogenous *nerve nets* [91] that made use of available electrical or chemical messaging processes [92] to broadcast sensory stimuli or motor commands across the animal's body. None of these mechanisms seem to fit the frameworks discussed above or even the function approximation paradigm in general!
- 3. Conversely, many biological neural networks also rely heavily on local synaptic, intrinsic and structural plasticity, and thus continuously adjust to changing circumstances (see chapter 5). The weights, biases and nonlinearities of deep networks, on the other hand, are typically assumed to remain fixed after initial training.
- 4. The transmission of signals also incurs delays, which are ignored in most artificial network models, yet they have significant and unavoidable implications on the behavior, in particular for real-time or recurrently connected neural networks (see chapter 4).

- 5. The neuron and synapse models in artificial neural networks are also typically modeled as memory-less functions, that instantaneously map an input onto an output. Biological neurons, on the other hand, integrate information in time (see chapter 4), adapt (see chapter 5) and can have a rather complex internal state and memory (see chapter 7).
- 6. For the majority of neurons in the brain, the output is not communicated as a real-valued signal, but via distinct spike events, which requires a fundamentally different mathematical framework (see sec. 6 and chapter 7) of event-based detection of spike patterns, rather than rate-based function approximation.
- 7. The complex morphology of biological neurons and their dendritic arbors is typically ignored in favor of point-neuron models — despite ample evidence of the relevance of that morphology for behavior (see chapter 7).

Given the remarkable complexity of biological neurons, it seems like a fool's errand to attempt to give a single model of neural computation on the same level of abstraction as current artificial neural networks and to then expect that it applies to "the brain" in general — let alone across different species. In the following chapters, I will therefore only attempt to describe a few of these points in some more detail. For a deep dive into the intricacies of many of these neural mechanisms, I'd refer to Singer, Sejnowski, and Rakic [93], which also contains contribution 2, or Laughlin [6] for a more "bottom-up" perspective.

Contribution 2: Computational Elements of Circuits

The book "The Neocortex", published by the Ernst Strüngmann Forum, compiles the current state of knowledge about the basic principles of operation of the neocortex. In our contribution to this work, the book chapter entitled "Computational Elements of Circuits", we discuss several fundamental properties of neural computation — from homeostasis to delayed interactions, synchronization, random feature expansion and reservoir computing. Within this book chapter, my own largest contribution can be found in the section "Information Processing in Single Neurons and Populations", which elaborates and generalizes ideas from contribution 6.

Reference :

J. Leugering, P. Nieters, and G. Pipa, "Computational Elements of Circuits," in The Neocortex, W. Singer, T. J. Sejnowski, and P. Rakic, eds., red. by J. Lupp, vol. 27, The MIT Press, 2019, pp. 195-209, ISBN: 978-0-262-04324-3. DOI: 10.7551/mitpress /12593.003.0016.

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In particular, this requirement [of a physical implementation] will help to prevent the solution from being a mere verbalistic 'explanation', for in the background will be the demand that we build a machine to do these things.

- W. Ross Ashby, Design for a Brain

Neuromorphic computing — a bridge between engineering and neuroscience

As we have seen in chapters 1 and 2, machine learning can produce very large artificial neural networks that require correspondingly large data sets and a lot of power to train. Since the available compute resources are a major bottleneck for the deployment of deep neural networks in many real-world applications, the numerical efficiency of artificial neural networks has become a major concern of deep learning research. This optimization of neural network models for the available hardware has certainly produced impressive results, but it also limits the scope of research to just those kinds of models that *can* be efficiently simulated on current hardware. Due to this compromise, modern deep learning on the one hand makes use of tools and algorithms that are not available to biological neurons, and on the other hand it cannot use many of the interesting biological mechanisms studied in theoretical neuroscience.

The field of *neuromorphic hardware* approaches this issue from the other side: If a neural mechanism seems promising for improving computation, but it lacks efficient hardware-support, then we should develop custom hardware rather than compromise our models! The design constraints of neuromorphic hardware are therefore determined only by what can be efficiently realized by analog and digital (or even ionic and photonic) circuits. A large range of biological mechanism that are difficult to integrate into a classical machine learning setting, such as spiking neural networks, are thus commonly used in the neuromorphic computing community. At the same time, the design constraints imposed by the hardware development raise other important questions that can help to challenge and improve theoretical models. As we shall see, many of the constraints faced by engineers are in fact quite similar to the limitations that biological neurons have to overcome. Hence, I believe that neuromorphic hardware can build a bridge between the two disciplines by providing engineers with biological inspirations, and neuroscientists with tools and measures to evaluate their models.

3.1 The neuromorphic zoo

In the wake of the impressive success of deep learning and the foreseeable end of Moore's Law, the research of new, alternative computing architectures and technologies on which to efficiently execute these neural network models attracts considerable interest from both academia and industry [95]. This has led to a revival of research on *neuromorphic hardware*, which promises to convert the theoretical insights from neuroscience into tangible benefits

3

Contribution 3: A Visit to the Neuromorphic Zoo

In this paper, which accompanies a public talk held at the Embedded World conference, I provide a brief overview over current concepts and academic as well as commercial developments in the field of AI-hardware acceleration in general, and neuromorphic hardware in particular. This paper was selected by WEKA Fachmedien for a republication in the magazine DESIGN&ELEKTRONIK, where it appeared in German translation under the title "Neuromorphe Hardware". Since the Embedded World conference accompanies an industry fair, the proceedings are targeted towards an engineering audience and are intended to offer an accessible high-level perspective. (These papers passed an editorial process, but no scientific peer-review.)

Reference:

J. Leugering, "A visit to the neuromorphic zoo," in *Embedded World Conference 2020 – Proceedings*, 2020, ISBN: 978-3-645-50186-6.

A German translation of this article appeared also in:

J. Leugering, "Neuromorphe Hardware," *DESIGN&ELEKTRONIK*, no. 7/2020, pp. 41–47, 2020.

for the development of biologically inspired, highly efficient computing hardware. While this idea is not entirely new¹, there are several good reasons for the renewed interest:

The first is *economical*: The stunning success of neural networks in recent years has revealed many new potential application areas for neural networks, from sensor and image processing and voice control all the way to autonomous robots and vehicles. Since many of these applications are "at the edge" [96], i.e. they do not have direct access to high performance computing infrastructure, they require on-board hardware capable of executing specific neural network architectures. Neuromorphic hardware can address this new and growing market.

Second, *technological* breakthroughs in the development of new materials such as various memristive devices [97], carbon nanotubes, in-silicon photonics, spintronics and much more [98], as well as improved procedures of lithography provide new freedom to implement neuromorphic architectures efficiently in hardware.

Third, there are promising new *theoretical* concepts in the field of neural networks as well as in electronics design. For example, an increasing emphasis is placed on non-volatile memory that can persist even when power is switched off. This is often paired with (analog) *in-memory* computing [99], which brings simple processing elements such as logic gates directly together with storage elements. That is great news for neuromorphic hardware, which can leverage this for an efficient implementation of synaptic connections [100, 101]. Similarly, a lot of theoretical models like spiking neural networks, that play only a minor role in conventional machine learning, are being actively explored in the neuromorphic computing domain [102].

All of these factors combined explain the current resurgence of neuromorphic hardware as one leg of the so-called *next generation computing* (NGC, the other leg being quantum computing). Recently, a large variety of hardware implementations, using analog, digital, mixed signal and even photonic circuits have been developed, and the research of neuro-

¹ In fact, neuromorphic hardware is as old as neural networks, since they both pre-date the emergence of powerful general purpose computers. An early example is the Perceptron Mark 1, a physical implementation of the perceptron model by Rosenblatt [1].

morphic computing has since developed into an independent discipline in academia and industry alike. For a brief review of the current state of the neuromorphic hardware field, see contribution 3.

A signal processing view of neuron models

In chapter 2 we saw how machine learning frameworks build large networks from individual neurons. Each neuron in that context is really just a function of the form $x_i(t+1)$ $f_i(\sum_i w_{i,i}x_i(t) + b_i)$, which becomes a seamless part of the larger function that describes the network as a whole. It makes little sense to ask, how much energy this neuron consumes, how much memory it has, or what it's latency is. But biological neurons have to have a physical realization of some sort, and this kind of question becomes critical. I therefore think the analogy between neurons and electronic components can be much more illustrative and satisfying than the mathematical abstraction of neural networks as function approximators if we want to better understand the behavior of real neurons. In the following, we will look at a few common neuron models, and we will explore how each of them resembles a well-known electronic component with a specific application in computer science or signal processing. These analogies will allow us to transfer some intuitions from engineering disciplines to neuroscience, and will thus help us better understand the computational capabilities and limitations of various neuron models. Figure 3.1 shows an overview of neuron models and closely related electric circuits.

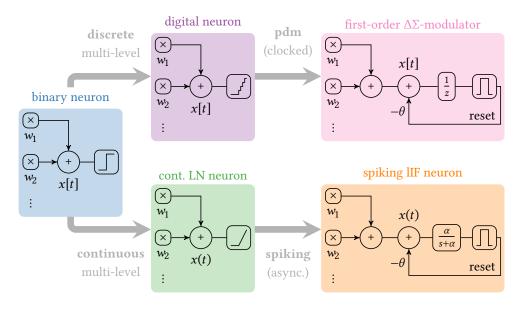


Figure 3.1. Neuron models, electronics components and their relationships. Boolean logic gates are equivalent to binary neurons (left). They can be extended by allowing multi-valued outputs, resulting in the standard linear-nonlinear neuron model in either digital (top center) or continuous (bottom center) form. Transmitting these multi-valued outputs by a pulse-density modulated code yields the first-order $\Delta\Sigma$ modulator (top right) for discrete time models or the leaky integrate-and-fire spiking neuron model (bottom right) for continuous time.

Binary neurons are logic gates 3.2.1

To get started, let's consider the most basic neuron model and logic calculus that McCulloch and Pitts [27] proposed. It states that the neuron's binary output y[t] within some brief time-interval t is 1 (the neuron emits a spike) if its membrane potential exceeds a critical threshold and 0 otherwise. Formally, $y[t] = f(\sum_{i=1}^{N} w_i s_i[t]) - b$, where $f(x) = \mathbb{1}_{[0,\infty)}(x)$ is the Heaviside step-function, w_i are the synaptic weights, b is a bias term and $s_i[t]$ are the values of the input signals at time-interval t. See figure 3.2 for an illustration. The neuron thus maps its N binary input signals onto a single binary output signal, and hence implements a Boolean function. If we limit ourselves to ternary weights as McCulloch and Pitts did, i.e. $w_i \in \{-1, 0, 1\}$, each input is either inverted (an inhibitory input), absent, or left unchanged (an excitatory input). We can then view the neuron as a *logic gate* that calculates whether *at least b* of its N weighted inputs are equal to 1. This "gate" includes as special cases the AND gate (b = N) and the OR gate (b = 1), negations thereof, the constant true gate (b = 0) as well as the constant false gate (b > N); it is therefore a *functionally complete set* of first-order logic. ²

Figure 3.2 shows a schematic of this model.

By assigning each input with an appropriate weight, we can thus construct arbitrary logic circuits, or *binary neural networks with ternary synaptic weights* [103]. From the perspective of deep learning, such *extremely quantized* networks may seem tedious, since they are difficult to train using gradient based-methods. In fact, logic circuits are traditionally optimized using numerically slower discrete optimization methods such as the Quine-McCluskey algorithm [104]. But recent advances in deep compression techniques are leveraging the powerful gradient-based optimization methods from deep learning for the optimization of ternary neural networks (or *XNOR networks*) and have produced highly promising results [74].

3.2.2 Linear-nonlinear neurons are summing amplifiers

A natural extension of these binary neurons to real-valued signals leads to the typical (discrete-time continuously-valued) *linear-nonlinear* neuron model that is typically used in deep learning. Here, the hard threshold is simply replaced with a continuous, typically monotonic non-linear function, e.g. a sigmoid function like $f(x) = \tanh(x)$ or the rectified linear function $f(x) = \max(0, x)$. Figure 3.3 shows a schematic. This reflects the observation, that the *firing rate* of a biological neuron increases as a continuous function³ of its membrane potential (see e.g. chapter 6). We already saw in chapter 2 that a network of such continuous *linear-nonlinear neurons* can be used to approximate arbitrary continuous functions, rather than just Boolean functions. But more importantly, the nonlinear function f that replaces the step-function of the McCulloch-Pitts neuron can be chosen to be (piece-wise) differentiable, which makes the entire network differentiable with respect to its parameters! Deep learning makes use of this fact and relies on gradient-based optimization methods to fix the synaptic weights.

Of course, a similar device is also useful for countless signal processing applications, in particular if we choose the rectified-linear function f as the nonlinearity. The behavior of this neuron could then, in engineering terms, be described as an ideal *summing unity-gain amplifier* or *buffer* that clips off negative values — a common component e.g. in analog audio circuits. See figure 3.3 for a schematic.

This linear-nonlinear model can be realized in hardware either by a digital or a fully analog electronic circuit, and both options are used in practice. Computing directly with analog voltages and currents is an extremely appealing concept and the cornerstone of many *neuromorphic hardware* designs, because it can result in very high energy efficiency and low latency. But analog computation comes with its own drawbacks: for one, analog signals in continuous time are difficult to buffer or route and hence require dedicated physical connections between the neurons, the number of which grows quadratically with the number of neurons. This arrangement may work well in the three-dimensional brain, ⁴ but poses a serious challenge for neuromorphic hardware that must be laid out in two dimensions. In addition to that, transmitting analog signals over large distances makes them susceptible to noise. A digital implementation of the same model (see figure 3.4) can alleviate these problems, but possibly at the cost of reduced accuracy due to quantization, increased circuit size and complexity, and increased power consumption. This discrete-time, discrete-value implementation is used in many digital neuromorphic circuits and is emulated by most DNN software-models.

² The *exclusive* OR (XOR) gate, however, was famously shown by Minsky and Papert [20] to be not representable by only a single layer of such neurons.

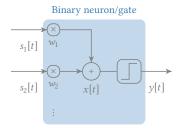


Figure 3.2. A binary neuron is a special boolean logic gate.

³ Strictly speaking, this only holds for the so called *class-I* or *type-I* class of neuron models; *type-II* neurons have a discontinuous jump in their firingrate response [105].

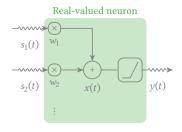


Figure 3.3. The linear-nonlinear neuron can be implemented in analog electronics.

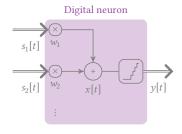


Figure 3.4. The neuron model in figure 3.3 can be discretized into a digital neuron model.

⁴ Even in the brain, the longer-ranging synaptic connections of the white-matter alone can make up half of the cortex by volume [106].

(Leaky)-Integrate-and-Fire neurons are $\Delta\Sigma$ -modulators

Spiking neurons combine some benefits of analog computation (namely energy efficiency and speed) with the benefits of binary transmission (namely noise robustness)5, because they process signals in the analog domain (whether in a biological neuron's dendrite or a neuromorphic circuit) while sending out only a series of binary pulses (see also chapter 6)!

⁵ For a neuromorphic hardware designer, there is the additional benefit of being able to route the discrete spikes through a bus system.

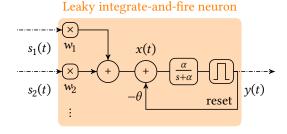


Figure 3.5. A leaky integrate-and-fire neuron converts analog or spiking input signals into a continuous-time spike-train. A negative feed-back loop resets the neuron after each pulse. The leaky integrator with leak-rate α is represented here by its Laplacetransform, $\alpha/s+\alpha$.

The simplest example of this is the well known integrate-and-fire neuron. In essence, it integrates its input(s) over time and fires a pulse whenever the integral x[t] exceeds a threshold. A negative feed-back loop then resets the system, and the process begins anew. Instead of a perfect integrator, a first-order exponentially decaying filter is often used to describe the response of biological neurons, which accounts for the fact that absent any input, the neuron's membrane potential tends to return to its resting potential over time. This model is called the *leaky integrate-and-fire neuron* (LIF neuron, see figure 3.5). Instead of the exponential filter, other filters could be used as well, which affects the neuron's response in interesting ways that we'll discuss in chapter 4.

The firing-rate of such a neuron encodes the input in a very similar way to the linearnonlinear neuron, it only uses a pulse-based code to transmit its output. For band-width limited signals, this encoding can be entirely lossless [107], but it may require rather large firing rates (see chapter 6 for a discussion).

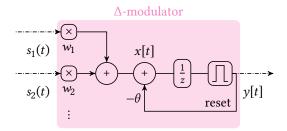


Figure 3.6. A first-order Δ -modulator converts analog input signals into a clocked sequence of binary pulses. A negative feed-back loop resets the component after each pulse. The discrete-time integrator is represented here by its z-transform, 1/z+1.

Such pulse-based communication schemes are also popular in electronics. In fact, the LIF neuron directly resembles a very popular electronic circuit, the so-called $\Delta\Sigma$ -modulator, which is an integral part of $\Delta\Sigma$ analog-to-digital converters [108] and pulse-width-modulators (PWM). In this comparison, the dendrite of the LIF neuron corresponds to a *demodulator*, which converts, sums and integrates the pulse-based input signals into a single analog signal, while the spike-generation mechanism at the soma uses Δ -modulation to encode whenever this signal has increased beyond a fixed threshold. Figure 3.6 shows a schematic.

To improve the noise characteristics of analog-to-digital converters further, another feedback can be added that subtracts the recent average output signal from the input and thus prevents the accumulation of quantization errors over time. This, too, has a direct counterpart in biological neuron models, namely the adaptive exponential integrate-and-fire neuron (AdExp, [109]), which also happens to be a more faithful representation of biological spiking neurons than the simpler LIF neuron [110]. We will discuss similar adaptation mechanisms in

chapter 5. By replacing the first-order integrator by higher-order filters, this can be improved further - an idea that we will also return to in chapter 4.

In an electronics context, the pulse-train generated by such a circuit constitutes an "over-sampled" digital *pulse-density modulated* (PDM) signal, which can then either be *decimated*, i.e. converted into a higher bit-precision signal at a reduced sampling rate, or directly transmitted over a digital connection. From this perspective, the (time-varying) density of the binary pulses (or *firing rate*) encodes the (time-varying) value of the analog signal. This view mirrors the *rate-coding* perspective, which we will look at in chapter 6.

A different perspective would be to treat the circuit as an event-detector, which emits a spike once it has accumulated enough input. We will discuss this alternative in chapter 7.

3.3 Closing the gap

For machine learning applications of artificial neural networks, the physical implementation of the individual neuron is of little concern — it represents an abstract mathematical function that is viewed as an "atomic operation" inside a larger algorithm. But from both a neuroscience and a neuromorphic computing perspective, the internal mechanisms that *generate* this behavior are of great interest. The two fields can therefore benefit from each other, by using engineering methods to investigate the function of biological neurons, or by taking inspiration from biological mechanisms for the development of a new generation of computing hardware. Ultimately, I believe these two disciplines ought to come together in a single discipline which I'll just refer to as *neuromorphic science* — the study of neuroscience-inspired physical mechanisms of information processing.

In the following, I will therefore occasionally use tools from signal processing and engineering to describe the behavior and information processing capabilities of biological neurons, and focus on the kind of questions that is also relevant for neuromorphic design. In chapter 7, I will then present a neuron model derived entirely from biological observations, along with an efficient neuromorphic circuit to implement it.

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A single neuron in the brain is an incredibly complex machine that even today we don't understand. A single "neuron" in a neural network is an incredibly simple mathematical function that captures a minuscule fraction of the complexity of a biological neuron. So to say neural networks mimic the brain, that is true at the level of loose inspiration, but really artificial neural networks are nothing like what the biological brain does.

— Andrew No

Dendritic filters and delays

In chapters 2 and 3, we discussed information processing in artificial neural networks of simple *point-neurons*. However, while these models offer a very convenient simplification, they don't account for the complex structure and behavior of real dendritic arbors, the behavior of which is better described by *neural cable theory* [111]. If we take the attenuation and delays into account that inevitably occur as membrane potentials are propagated along the dendrite, then even the location of a synapse on the spatially extended dendrite influences the effect of an input signal on the neuron's firing rate [112]! This makes the behavior much more difficult to describe, but it could also increase the computational complexity of the individual neuron considerably by endowing it with a notion of time or *memory*. Among the first to see the serious implications of this arrangement was, once again, von Neumann [35]:

It may well be that certain nerve pulse combinations will stimulate a given neuron not simply by virtue of their number but also by virtue of the spatial relations of the synapses to which they arrive. That is, one may have to face situations in which there are, say, hundreds of synapses on a single nerve cell, and the combinations of stimulations on these that are effective (that generate a response pulse in the last-mentioned neuron) are characterized not only by their number but also by their coverage of certain special regions on that neuron (on its body or on its dendrite system, cf. above), by the spatial relations of such regions to each other, and by even more complicated quantitative and geometrical relationships that might be relevant. [...] Lastly, I would like to mention that systems of nerve cells, which stimulate each other in various possible cyclical ways, also constitute memories. These would be memories made up of active elements (nerve cells).[35].

In a very similar sense, the delayed interaction of neurons by synaptic spikes, which could be seen as an imperfection of an idealized neuron and thus a nuisance to be avoided, also increases the complexity of the neuron's behavior by introducing long-lasting dependencies, i.e. *memory*. These complex nonlinear dynamics and long memory are very appealing for reservoir computing, where neurons with delayed feedback have recently been evaluated as a potential computational substrate. But like many other concepts mentioned in this thesis, the idea to use delayed self-coupling as a form memory and computing device is actually surprisingly old and goes all the way back to Turing and Copeland [4], who proposed building a computer based on the delayed interaction of shock waves in tubes of mercury:

It is proposed to build 'delay line' units consisting of mercury [...] tubes about 5' long and 1" in diameter in contact with a quartz crystal at each end. The velocity of sound in ...mercury [...] is such that the delay will be 1.024ms. The information to be stored may be considered to be a sequence of 1024 'digits' (0 or 1) [...] These digits will be represented by a corresponding sequence of pulses. The digit 0 [...] will be represented by the absence of a pulse at the appropriate time,

the digit 1 [...] by its presence. This series of pulses is impressed on the end of the line by one piezo crystal, it is transmitted down the line in the form of supersonic waves, and is reconverted into a varying voltage by the crystal at the far end. This voltage is amplified sufficiently to give an output of the order of 10 volts peak to peak and is used to gate a standard pulse generated by the clock. The pulse may be again fed into the line by means of the transmitting crystal, or we may feed in some altogether different signal. We also have the possibility of leading the gated pulse to some other part of the calculator, if we have need of that information at the time. Making use of the information does not of course preclude keeping it also [4].

Therefore, two aspects of biological neurons that are often disregarded as a nuisance, dendritic filtering and synaptic transmission delays, could theoretically serve an important purpose for neural computation. But is all this complexity really instrumental, or is it just an inevitable side effect of some biological process, through which nature approximates a much simpler mechanism? In this chapter, we'll make use of some simple tools from signal processing to investigate the role of dendritic filtering and delays for neural information processing.

4.1 Terminology

Before we get started, I'd like to introduce a few terms that are used inconsistently across different disciplines: The general mathematical formalism we'll use is temporal *convolution*, which is also called *filtering* in engineering domains. Since physical systems cannot be retroactively affected by future events, *causal filters* play a special role, which are linear integral operators that only depend on the values of the signal in the recent past. Such a filter operator K can be applied to a continuous-time signal s by the convolution

$$Ks(t) = (s * \kappa)(t) = \int_{-\infty}^{t} s(\tau)\kappa(t - \tau)d\tau$$

where κ is the *kernel* or *impulse-response function* of K. *Delays* are a special case of causal filtering that can be represented by the convolution $(s * \kappa)(t) = s(t - \Delta t)$ with a shifted Dirac- δ -distribution kernel of the form $\kappa(t) = \delta(t - \Delta t)$, where Δt is the duration of the delay. Conversely, we can interpret a continuous filter kernel κ as the limiting case of a linear combination of delay terms (for a derivation, see appendix A.1). This equivalence is used in the (digital) signal processing domain to design filters for *periodically sampled* discrete-time signals [113, p. 67]. Therefore, filtering and delays are really two sides of the same coin!

I will also use the one-sided Laplace transform $\mathcal{L}\{\kappa\}(s) = \int_0^\infty \kappa(t) \exp(-st) dt$, which is closely related to the Fourier transform, to represent a filter with kernel $\kappa(t)$ in the frequency domain [114]. The Laplace transform can simplify the analysis greatly, because it allows us to represent a concatenation of multiple filters in the frequency domain simply by the multiplication $\kappa_1(s) \cdot \kappa_2(s) \cdot \kappa_3(s) \cdot \kappa_4(s) \cdots$ of their individual transformations, rather than by the convolution $\kappa_1(t) * \kappa_2(t) * \kappa_3(t) * \kappa_4(t) \ldots$ in the time domain. If it is clear from context, I will just write $\kappa(t)$ and $\kappa(s) := \mathcal{L}\{\kappa\}(s)$ to denote the filter kernel in the time- or the Laplace-domain, respectively.

4.2 Dendritic filtering improves information transmission

Why should we care about the filtering effect of dendrites, in the first place? For one, because including it greatly increases the accuracy with which the firing-rates of biological neurons can be approximated when compared to simpler point neurons [3, 115]. But more importantly, filtering offers an opportunity to improve the information transmission and

processing capabilities of individual neurons. This same argument can be made in multiple ways, and we'll look at five different perspectives in the following.

4.2.1 Denoising

From the perspective of information theory, the linear-nonlinear neuron is a noisy channel with limited capacity to transmit information (see also chapter 2 of [7] and the later chapter 5). This is formalized in the Shannon-Hartley theorem [7, 113], which states that an analog channel's ability to transmit information (in some frequency band) is limited by the channel capacity $C = B \log_2(1+R)$, where B is the analog channel bandwidth and R is the signalto-noise-ratio, i.e. the ratio of the expected power of the signal to be transmitted relative to the power of the independent noise signal. Since the amount of transmitted information adds up linearly across distinct frequency bands, it follows that a carefully chosen filter can selectively amplify or attenuate different frequency bands to boost the signal while suppressing the noise. Therefore, a key benefit that an appropriately chosen dendritic filter can offer for information transmission is to improve the signal-to-noise-ratio and thus the channel capacity of the neuron. 1 In general, it will be impossible to completely filter out all noise this way, because the spectra of signal and noise are likely to overlap. But the signal-tonoise ratio can always be optimized by the so-called (causal) Wiener filter or matched filter [116], which shapes the spectrum in a way that maximizes the relative power of the signal while minimizing that of the noise.

¹ The simplest example is when the signal and the noise occupy distinct frequency bands altogether, in which case a simple band-pass filter can be used to fully isolate the signal and suppress the noise.

4.2.2 Pattern detection and sparse coding

Denoising can also be understood in a rather different way. Let's consider the special case that the signal s(t) consists only of repetitions of some stereotypical pattern g(t), $0 \le t \le T$ with duration T subject to white noise $\eta(t)$, i.e.

$$s(t) = \sum_{\tau_i \le t} g(t - \tau_i) + \eta(t) = (g * \chi)(t) + \eta(t) \quad \text{with } \chi(t) = \sum_{\tau_i \le t} \delta(t - \tau_i).$$

Then the power spectrum of the noise is flat, and the kernel of the Wiener filter simplifies to the time-reversed pattern h(t) = g(T - t). Denoising the signal s with this filter h yields the signal

$$(h*s)(t) = (r*\chi)(t) + (h*\eta)(t) \quad \text{ where } r(t) = (h*g)(t) = \int_0^t g(\tau)g(\tau + T - t)d\tau \text{ for } t \le t.$$

So whenever the stereotypical pattern g is seen in the input, the filter responds with r, the autocorrelation function of g, which always has a distinct maximum at t=T. These peaks in the signal can be easily detected by an appropriately chosen threshold despite the noise, because filtering the white noise η with the same kernel merely results in colored noise with comparatively lower amplitude. Therefore, a neuron with appropriate dendritic filter can become an efficient pattern detector (with delay T) for the stereotypical pattern g. Detectors of this sort have long been used, for example, in radar systems to detect reflected radio pulses of known shape [116], and the same ideas transfer to other time-series signals. In chapter 7, we'll contrast this to a rather different type of pattern detector.

4.2.3 Deconvolution

The problem of pattern detection can also be approached from a different perspective: Imagine that in the same setting as in section 4.2.2 we'd like the filter h to directly return $(h * s)(t) = \chi(t) + (h * \eta)(t)$ rather than $(r * \chi)(t) + (h * \eta)(t)$. This inverse problem, called

deconvolution, is unfortunately generally ill-conditioned² and outright impossible if we are limited to causal filters. ³ But we could introduce another filter q and try to solve the relaxed problem $h*s = q*\chi + h*\eta$ instead, i.e. h*g = q. If we choose q well, this can be (approximately) solved for h even if g cannot be inverted. Looking back at section 4.2.2, we can e.g. choose q = r, which again gives us the matched filter h(t) = g(T - t). Or we could choose the filter q to approximate a delay-line with delay T' > T, which would allow us to approximate a *delayed* deconvolution (with delay T')!

4.2.4 Equalization

Another way of looking at the same idea is the *equalization* [118] (sometimes also called *whitening*) of signals, which removes temporal correlations from the signal itself and thus leads to an *equalized* or flat power spectrum, resembling that of *white noise* (hence the name). This is done in the context of communication systems with bandwidth-limited communication channels, where the most information can be transmitted if all the available spectral bandwidth is used to convey relevant (i.e. non-predictable) information. A very similar argument can be made for the neuron as well, and we'll return to this idea also in chapter 5 when we talk about optimal firing rate distributions.

4.2.5 Predictive Coding

A more biologically motivated perspective is *predictive coding* (see e.g. chapter 6 of [7]), which argues that the dendritic filter can subtract predicted *future* inputs, leaving only the residual error to be transmitted by the neuron. The benefit of such an encoding is, again, that (given a sufficiently good prediction) these residuals are temporally decorrelated (i.e. equalized), which leads to an information theoretically and metabolically efficient encoding [7, chapter 6]. This has been experimentally observed in visual [119, 120] and auditory neurons [121], as well as in other modalities and animals [6], and is believed to play an important role for neural information processing in general [122].

4.3 Dendritic filtering in the linear-nonlinear model

We saw that, when we view the neuron as a signal-processing device, the ability to implement dendritic filters is extremely attractive. We will now investigate, how this could be implemented on a mechanistic level.

If we are willing to ignore all the non-linear effects that can occur in neural dendrites, such as dendritic plateau potentials,⁴ and instead focus exclusively on the linear effects described by neural cable theory [111], then the behavior of a dendrite can be approximated by assigning a specific impulse response to each synapse, depending on its location on the dendrite. The somatic membrane potential can then be approximated by a linear combination of these differently filtered synaptic inputs. The result is a more general type of linear-nonlinear neuron model, which I'll just call the "filter-nonlinear" model in the following (see figure 4.1).

A very simple special case of this model is Rall's so-called *ball-and-stick model* [123]: Under certain simplifying assumptions (e.g. specific relationships between the thickness of dendritic branches), the effect of a synaptic input onto the soma's membrane potential only depends on the *distance* of the synapse to the soma — the complex tree-shaped topology of the dendrite can be ignored altogether. In this case the dendrite therefore behaves equivalently to a single cylindrical dendrite as shown in the top panel of figure 4.5. This cylinder can be approximated by a chain of multiple *compartments*, each of which receives and filters input from its "up-stream" neighbor compartment as well as synaptic inputs. How many of these

² For a discrete convolution operator, which can be represented by a matrix, this can be done through iterative matrix-inversion methods with strong regularization [117], and it becomes more complex for continuous-time operators.

 $^{^3}$ Just consider, for example, the kernel $\delta(t-\tau)$ with a delay $\tau>0$. Inverting this filter would imply inverting the time-shift, which would require an acausal "negative delay"!

⁴I actually believe that these nonlinear effects are absolutely crucial for dendritic computation, and I argue for this position in chapter 7.

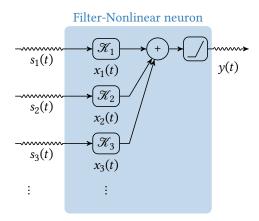


Figure 4.1. A linear-nonlinear neuron, where the dendrite is modeled by a set of filters \mathcal{K}_i . Each synaptic input $s_i(t)$ now produces a different post-synaptic membrane potential $x_i(t)$, which are then summed up and non-linearly transformed at the

compartments a synaptic input has to traverse depends on the distance of the synapse to the soma, therefore the dendrite can be also viewed as a filter bank. See the middle panel of figure 4.5 for an illustration.

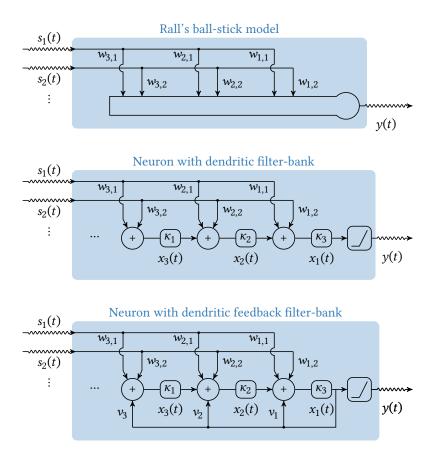


Figure 4.2. Top: The ball-and-stick model [123] abstracts the neuron's dendritic arbor into a single "equivalent cylinder" or cable, on which the propagation of activity can be modeled by a partial differential equation. The impulse response of an input signal depends on the location along the cylinder. Middle: A neuron with a dendrite modeled by a tapped filterbank composed of individual filters κ_i . Each tap of the filter-bank provides a local state-variable $x_i(t)$. The neuron's output $y(t) = f(x_1)$ is then just a non-linear function of the somatic membrane potential x_1 . **Bottom:** By adding a linear feedback term, more complex filters can be constructed.

Just like forward-propagation, the effects of backward-propagation of membrane potential in the retrograde direction, i.e. away from the soma, can be also incorporated by adding linear feedback terms (see the bottom panel of figure 4.5). In this model, the weights of the (feed-forward) input and the weights of the feedback term parameterize a family of dendritic filters that can approximate the filtering effect of a dendritic tree. ⁵

In the following, we'll look at two particularly simple and relevant implementations of such a dendritic filter banks:

⁵ Curiously, this type of dendritic filter with feed-back resembles a well known topology of infinite impulse response filters for electrical signal processing [124]!

4.4 Dendritic filtering in the Gamma Neuron

If we implement the dendrite shown in figure 4.5 as a bank of identical first-order low-pass filters with transfer function $\kappa_i(s) = \frac{\alpha}{s+\alpha}$ and time-constant α , then this much simpler multi-compartment model can be described by a system of *ordinary* differential equations instead of Rall's original *partial* differential equation. The resulting neuron is shown in figure 4.3.

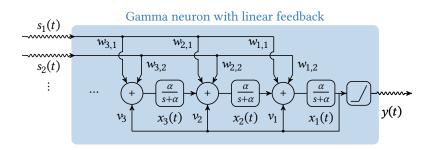


Figure 4.3. The Gamma Neuron uses a filter-bank composed of identical first-order filters with transfer function $\frac{\alpha}{s+\alpha}$ to model the dendrite. Inputs are projected onto the filter taps through multiple synapses with weights $w_{j,i}$. Similarly, feed-back paths can be added with weights v_j . These weights parameterize the dendritic filter. A nonlinear activation function is applied to the filter output.

The low-pass filter response of each tap resembles the simplified sub-threshold dynamics of the leaky integrate-and-fire neuron, and the filter bank can hence be interpreted as a chain of weakly coupled *dendrite compartments*. The result is the versatile *Gamma-neuron* [125], which can be represented by a particularly simple system of ordinary differential equations. This model owes its name to the fact that a synaptic spike arriving at the k^{th} compartment from the soma would be subjected to the transfer function $(\alpha/s+\alpha)^k$, which is the Laplace transform of the density function of some Gamma distribution. The impulse response of the dendrite therefore becomes broader and broader with every additional compartment that a spike has to traverse on its way to the soma (see figure 4.4).

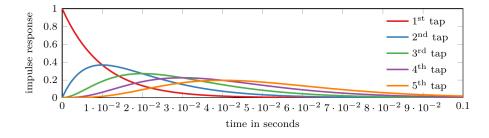


Figure 4.4. Open-loop impulseresponses of the somatic membrane potential in response to synaptic input spikes received at various taps along the dendrite. The impulse responses have the form of the probability density function of Gamma distributions, hence the name *Gamma Neuron*.

Now, if the same input signal s_i arrives at various taps j along the dendrite through multiple synapses with weights $w_{j,i}$, then the total effect of the dendrite on that input is a linear combination of the individual taps' responses. By adding a feedback path with additional coefficients v_i , we can extend this neuron model to allow much more complex filters to be implemented by the dendrite [126]. This family of filters can be formalized nicely (see the note below) and includes a lot of interesting special cases: For example, low-, high-and band-pass filters can be implemented with just two taps, and with increasing order the (open-loop) impulse response of the taps more and more resembles Gaussian filters. Also, for any filter κ that can be implemented by a Gamma Neuron, the derivative κ' can be trivially implemented as well. See appendix A.3 for derivations.

If we put all of this together, a single Gamma neuron could therefore theoretically

- (a) extract relevant frequency-bands from each of its multiple input signals,
- (b) equalize each signal to extract the maximum amount of information,
- (c) calculate derivatives or ("leaky") integrals thereof,
- (d) and linearly combine them into a single signal that is then

Note: The ring of Gamma filters

The filters that can be implemented by the Gamma neuron span a finite-dimensional function space, parameterized by the input and feedback weights. With a bit of algebra, we can make this more precise (see appendix A.2 for a derivation): The space of filters that can be implemented by the Gamma neuron corresponds to exactly those with a proper rational transfer function in the Laplace domain. The addition and convolution (i.e. concatenation) of two implementable filters yields another implementable filter, but the inverse κ^{-1} of an implementable filter κ is in general not a proper rational function (i.e. κ^{-1} is acausal) and thus not implementable. Therefore, the class of filters that can be implemented by such a filter bank forms a commutative ring (or rather pseudo-ring) without multiplicative identity and inverse. This space of filters is extremely general; it contains all analog linear filters that can be implemented by networks of lumped electric elements, i.e. discrete resistors, capacitors and inductors [127], and any transfer function can at least be well approximated by such a rational function, also called a Padé approximant [128].

(e) passed through a nonlinearity.

This also makes the individual neuron at least as powerful as a PID controller [129], a versatile tool from control theory and much more impressive than the simple logic gates we saw in chapter 3!

In contribution 4, we extend this model further and employ a synaptic plasticity rule to train individual (spiking) Gamma neurons to detect specific temporal patterns in their input.

Contribution 4: Training the Gamma Neuron for event detection

We extended the Gamma Neuron to a spike-based temporal pattern detector for a conference poster presented first at the Cognitive Computing 2018 conference in Hannover, Germany, and then again at the Machine Learning Summer-School (MLSS) 2019 held in Cape Town, South Africa. Here we investigated how this type of neuron model could be trained to produce a spike-based classification of temporal patterns through a local, reward-modulated synaptic learning rule.

Reference:

P. Nieters, J. Leugering, and G. Pipa, "Neuromorphic Adaptive Filters for event detection, trained with a gradient free online learning rule," presented at the Machine Learning Summer School (MLSS-Africa 2019), 1, 2019.

4.5 Computing with synaptic delays

We already saw above that delays are just a special case of causal filtering and vice versa, but delays deserve special treatment in the study of neural systems. They are a fact of life, since no physical system can respond *instantaneously* to its input - and of course neurons are no exception. However, even minuscule delays can make otherwise benign dynamical systems difficult to control or even chaotic [130], which is why they are often seen as a nuisance to be avoided by theoreticians and engineers alike. But biological neurons are inherently analog machines that work asynchronously and in real time, so we have no choice but to recognize and understand the effect of delays on their dynamics. As we shall see, this might be a blessing in disguise, since there are even (somewhat surprising) ways in which delays might actually *improve* the computational capabilities of neurons and networks!

To understand *how* delays, e.g. caused by synaptic transmission, could be used constructively by biological neurons, we need to make a brief detour into (digital) signal processing and control theory.

4.5.1 Delay-embeddings, state-estimation and Koopman-control

So far, we looked at continuous-time signals and filters, but a lot of the intuitions about information transmission come from the study of *sampled discrete-time* systems. The connection is established by the *Nyquist sampling theorem* [118, 124], which states that any analog bandwidth limited signal can be fully represented without loss of information by *samples* of the signal, if they are measured at a sufficiently high finite sampling rate. In that context, the same effect that a continuous filter would have on a continuous signal can be achieved by filtering the sampled signal with a discrete-time filter, which can be implemented by a linear combination of the outputs of a *tapped delay-line*. ⁶ Just like in the Gamma neuron, a linear feedback loop can be used to extend these filters, which allows us to also construct *infinite impulse response filters*. See [118, 124, 131] for an introduction into filter design and tapped delay-lines.

⁶ This is analogous to how we used a continuous filter bank above to construct the dendritic filter of the Gamma neuron.

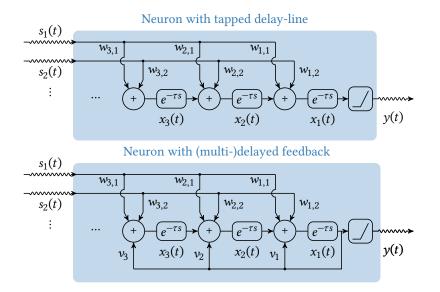


Figure 4.5. **Top:** A neuron with dendritic or synaptic delays modeled by a tapped delay-line composed of individual delay elements with transfer function $e^{-\tau s}$ and identical delay τ . This topology can implement finite impulse-response filters. **Bottom:** By adding a linear feedback term, infinite impulse-response filters can be constructed.

But of course, tapped delay-lines can also be utilized in a continuous-time setting. In that context, we'd say that the outputs of all the taps constitute a *delay embedding* of the signal — a higher dimensional representation of the signal and its recent past. Such an embedding contains a lot of information about the signal that can, for example, be used to estimate derivatives or to forecast the signal into the future. See also the note below. Under the more general heading of *embedding theory*, this has many practical applications for the study of dynamical systems, signal analysis and causality. Schumacher [132] has a great discussion of this subject. One fairly recent application of these ideas has been in control theory, specifically *Koopman control* [133–135], which comprises many state-of-the-art approaches to controlling non-linear systems by the use of delay embeddings.

Note: Derivatives, finite differences and delay embeddings

To give an intuitive example, how delay embeddings can be used to extract relevant information from a continuous-time signal, let's consider the definition of (higherorder) derivatives. The left derivative of a function f can be defined as f'(t) = $\lim_{\Delta t \to 0} \frac{f(t) - f(t - \Delta t)}{\Delta t}$. So if we can produce a delayed signal $\tilde{f}(t) = f(t - \Delta t)$ for a small delay Δt , we can use the signals f and \tilde{f} to continuously estimate the derivative f'. In fact, the same idea can be applied repeatedly: if $f'(t) \approx \frac{f(t) - f(t - \Delta t)}{\Delta t}$ and $f'(t - \Delta t) \approx \frac{f(t - \Delta t) - f(t - 2\Delta t)}{\Delta t}$, then $f''(t) \approx \frac{f'(t) - f'(t - \Delta t)}{\Delta t} \approx \frac{f(t) - 2f(t - \Delta t) + f(t - 2\Delta t)}{\Delta t^2}$, and so on. We can thus estimate the first N derivatives of a signal by a linear combination of N+1 delayed versions with delays $k \cdot \Delta t, k \in \{0,1,2,\ldots,N\}$. I'll let $\delta^k(t)$ denote the delay line $\delta(t - k\Delta t)$. The mapping

$$f(t) \to ((\delta^0 * f)(t) \quad (\delta^1 * f)(t) \dots (\delta^N * f)(t))$$

is then a so-called delay embedding, which embeds the one-dimensional time-varying signal f into an N + 1-dimensional space. This can be implemented by a tapped delay*line* composed of N concatenated delay elements, each with the same delay Δt . From this embedding, approximate derivatives of order $\leq N$ can be trivially read out by linear combinations of different taps' outputs. Therefore, an *n*-tap delay-embedding contains enough information about the signal to approximate a Taylor-approximation of order n-1 at the current point in time!

Delayed nonlinear feedback 4.5.2

So far, we only discussed linear systems with delay, but what if we include nonlinear feedbackloops? The result is a nonlinear delay- or retarded differential equation, which has an infinite dimensional state-space and can exhibit extremely complex, if not chaotic, behavior. The inherent complexity of such systems with nonlinear delayed feedback can introduce very long-lasting memory effects (see also [130] for more examples). 7

Filter- or multi-delay-coupled reservoir computing

The idea to use the long memory of systems with delayed nonlinear feedback also underlies an admittedly weird neuron model called the single node, multi-delay-coupled reservoir computer (SNMDCR). It is an extension of the slightly simpler single node, delay-coupled reservoir computer (SNDCR), which is situated between machine learning, neuroscience and optical neuromorphic hardware [137].8 We extend this model in contribution 5 to use multiple delayed feedback terms (hence the slight change in name), which considerably increases the neuron's ability to learn complex temporal dynamics. The resulting neuron model is summarized in figure 4.6.

Figure 4.6. A single-node multi-delaycoupled reservoir. It resembles the neuron in figure 4.3, but with delay elements instead of exponential filters for all but the first tap. Each tap delays the signal by τ . Note that the feedback-signal is here taken after the nonlinearity, so the behavior of the entire system is no longer of the simple linear-nonlinear form.

SNMDCR neuron $s_1(t)$ $w_{3,1}$ $w_{2,1}$ $w_{1,1}$ $w_{3,2}$ $w_{2,2}$ $s_2(t)$ $w_{1,2}$ α $e^{-\tau s}$ $s+\alpha$ y(t) $x_3(t)$ $x_2(t)$ $x_1(t)$

⁷ One example of such systems are feedback shift registers, whose very long memory is used for the generation of maximally long sequences of non-repeating pseudo-random numbers [136]!

⁸ The original physical realization of the system was implemented by selfcoupled lasers with optical delay elements.

Contribution 5: Neuromorphic computation in multi-delay coupled models

In this paper, we explored how delayed feedback, in particular the interaction between differently delayed feedback-loops, can be exploited to endow a single neuron, which could be implemented in an electrical or photonic circuit, with memory and the capability to compute complex functions of its input history. For a simple single-node multi-delay-coupled reservoir neuron, we show how the relationship between the delay terms leads to different complexity of behavior, and hence different performance of the trained neuron across different time-series regression tasks. Curiously, we can show that — and why — co-prime delays result in the best performance, and thus give some intuition for the complex behavior of delay-coupled systems.

Reference:

P. Nieters, **J. Leugering**, and G. Pipa, "Neuromorphic computation in multi-delay coupled models," *IBM Journal of Research and Development*, vol. 61, no. 2/3, 8:7–8:9, 1, 2017, ISSN: 0018-8646, 0018-8646. DOI: 10.1147/JRD.2017.2664698.

Due to the combination of continuous dynamics and delayed feedback, the SN(M)DCR must be modeled by delay-differential equations, and it shows highly complex if not chaotic behavior, depending on the precise choice of the relative delays. We use this neuron with its complex dynamics as a substrate for reservoir computing, i.e. we inject various taskspecific input signals into the neuron and use a weighted linear combination of the delay embedding of the neuron's output as a readout. As usual, (only) these weights are optimized such that the readout approximates the desired output signal of the task. To investigate the impact of the precise choice of delays on the ability of the SNMDCR to produce interesting behavior, we look at a neuron with just two feedback paths with different delays τ_1 and τ_2 . We systematically vary one of the two delays τ_2 while keeping the other fixed, and for each choice of τ_2 optimize the neuron's weights for some simple task, like estimating the N-bit parity of a binary signal or approximating a fixed NARMA model of a continuously-valued signal. Remarkably, the SNMDCR performs very well on either task, but its performance critically depends on the relative timing of the two delays and deteriorates whenever this ratio approaches a ratio of small integers such as 1:1,1:2,2:3 etc. Since the SNMDCR is described in discrete time and the delays are integer multiples of these time-steps, we can compare the location and magnitude of the performance drops to the greatest common divisor of the two delays and find a clear correspondence.

To get a better mechanistic understanding of how and why the SNMDCR works (and when it fails), we analyze how it integrates and recombines information over time, and conclude that co-prime delays provide the best "mixing" over time with the longest memory, which appears to be the critical factor for performance on these tasks. While these results are specific for an unusual type of neuron model and cannot be directly transferred to others, they nevertheless provide a useful intuition: Dendritic filtering and synaptic delays can be used to not only extract relevant information from time-varying signals, but also to improve information transmission, provide volatile memory, and implement computation within a single neuron!

⁹ For example, in the continuous-time context of biological neurons, the inherently discrete concepts like coprimality and greatest common divisors are not applicable.

Dendritic filtering in the real world 4.6

We have seen for a couple of examples above how a neuron could, in principle, make constructive use of the delays and filtering effects introduced by synaptic transmission and the dynamics of ion currents in the dendrite. A natural question to ask now is: How much, what for and how, if at all, do biological neurons actually use dendritic filtering? A second question is: Should we use dendritic filtering in machine learning models of neural networks?

On the one hand, there has been a lot of biological evidence that shows dendrites using delays and filtering to do much more than just instantaneous linear combinations of incoming signals. For example, the distance-dependent filtering and delaying effect of dendrites has long been proposed as a critical feature for binaural localization of sounds [138]. There is even some evidence that these transmission delays can be fine-tuned by controlling the myelinization of axons [139], which would represent an entirely new form of plasticity mechanism! The huge theoretical potential that dendritic filtering can offer for processing time-series signals also makes it likely that evolution would have found ways to exploit it in some way.

On the other hand, it is also tempting to entirely brush off the intimidating complexity of biological dendrites as functionally irrelevant "implementation details", and there is also biological evidence to support this view. For example, the attenuation along the dendrite appears (in some cases) to be precisely counteracted by some other mechanism like synaptic scaling or "synaptic democracy" [140, 141]. This could ensure that each synaptic spike, regardless of its location on the dendrite, has the same effect on the somatic membrane potential. Through such regulatory mechanisms, an apparently complex nonlinear neuron could produce a rather simple linear behavior that is well described by the point-neuron model, after all. 10 But more critically, the real time-constants of dendritic filtering and synaptic delays (on the order of microseconds to tens of milliseconds) might just not be long enough to implement most useful filters on behaviorally relevant time-scales. And on a more fundamental level, the strongly nonlinear effects that can be observed within small dendritic branches call the assumption into question, that the temporal dynamics can be well approximated by linear filters, at all! Instead, the interaction of active, localized nonlinear processes within the dendrites need to be taken into account [142]. We will return to this point in detail in chapter 7.

It might therefore turn out, that biological neurons make only limited and rather specific use of (linear) dendritic filtering, e.g. for the purpose of adaptation (see chapter 5) and for the processing of spiking inputs (see chapter 6), while relying on different mechanisms on the neuron or network level, such as active dendritic processes (see chapter 7) or recurrent networks (see chapter 2) for more sophisticated temporal integration and processing of

Of course, for machine learning models and neuromorphic hardware these biological constraints do not apply, and dendritic filtering is certainly worth considering. However, the introduction of dendritic filters (in analog or digital hardware as well as in software) complicates the individual neuron substantially, and the additional slow dynamics makes temporal credit assignment difficult. Models like the Gamma neuron are therefore more difficult to simulate and to train using conventional gradient-based methods (see contribution 4). I therefore see the most promising applications of these ideas in conjunction with local learning rules (e.g. for unsupervised equalization of signals) and/or in the context of analog neuromorphic hardware, where filtering is inevitable and can be efficiently realized by simple electronic circuits.

¹⁰ However, such a regulatory mechanism might only affect the amplitude, not the delay, of a synaptic input as a function of its location on the dendrite. In that case, the arguments of this section could still be applied.

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The green reed which bends in the wind is stronger than the mighty oak which breaks in a storm.

Confucius

'[A]daptive' behaviour is equivalent to the behaviour of a stable system

— W. Ross Ashby, Design for a Brain

Homeostatic plasticity

In the previous chapters, I presented networks and neurons as information processing "machines" and likened them to logic gates and other electronic components. But while logic gates are fed a steady diet of ones and zeros, nervous systems are embedded in biological organisms, and they are bombarded by noisy input signals from an ever-changing environment, perceived through sensors that themselves develop or degrade over time. To keep working in this chaotic setting requires the organism to take active counter-measures to maintain itself. This ability to adapt to changes is so critical for survival that early cyberneticists like Ashby [2] saw the concept of homeostasis as the defining feature of life, and one of the main differences that sets life (and nervous systems) apart from dead matter (and logic gates). The term "homeostasis" entails that some attribute(s) of the organism are maintained at a desirable state, and that the system can recover this state from small disturbances through some self-regulating mechanism. In the case of the nervous system, that could mean to remain functional (or to quickly regain functionality) even if certain aspects of the sensory inputs change abruptly. The idea of homeostatic adaptation is therefore quite central in theoretical and computational neuroscience, and it should play an important role for our understanding of (artificial) intelligence as well. In fact, a survey of different definitions of intelligence [143] found that at least 23 out of the 72 definitions see the ability to adjust or adapt to the environment as a defining feature, and many of the others imply it!

But in most current deep learning research, adaptation plays only a minor role, and most of the big data sets on which models are trained and evaluated have been explicitly preprocessed to remove any of the systematic changes or drifts that would require the system to adapt in the first place¹. Since self-regulating, adaptive systems are also typically harder to understand, control and train than static ones, the homeostatic plasticity mechanisms that we know from biological neurons are hence still largely absent from machine learning models.

In this chapter, I'd like to illustrate why homeostatic adaptation is not just a biological necessity, but also a useful mechanism for neural information processing in general. I'll present an abstract framework that unifies two different forms of biological plasticity mechanisms to solve a practical, easily interpretable machine learning problem. Most content of this chapter is directly based on contribution 6, but it offers another, hopefully simpler motivation for the main results, while leaving out a lot of the technicalities here. Nevertheless, this will take us through a number of abstract mathematical concepts, and I will try my best to explain them here on a rather high level.

5

¹ Many image recognition challenges, for example, present a fixed set of training images in randomized order. Often, these images are coloradjusted, scaled to equal size, centered or otherwise prepared. It would be considerably harder, if the training set was allowed to change over time.

Contribution 6: A Unifying Framework of Synaptic and Intrinsic Plasticity in Neural Populations

In this rather long paper, I explore the relationships and interaction of intrinsic and synaptic plasticity for computation. The entire chapter 5 of my thesis is largely based on ideas contained within this publication. I try to motivate the same main results here using a slightly different approach that introduces concepts like optimal transport theory. But for most content of the current chapter, a more in-depth discussion can be found within this original publication.

Reference:

J. Leugering and G. Pipa, "A Unifying Framework of Synaptic and Intrinsic Plasticity in Neural Populations," *Neural Computation*, vol. 30, no. 4, pp. 945–986, 17, 2018, ISSN: 0899-7667. DOI: 10.1162/neco_a_01057.

5.1 The Information Bottleneck Principle

The Information Bottleneck Principle [66, 144] loosely states that in neural networks, the capacity to transmit information from neuron to neuron, or from layer to layer, or from region to region, is often the limiting factor for computation — a bottleneck, so to speak. As such, the capacity should be used providently, and neural computation should be optimized to make efficient use of it. This provides a clear objective, towards which a neuron or network can be optimized: to convey as much information as possible about the input signal through an information channel with limited capacity. A popular and quite literal example of this idea are auto-encoders [145] in deep learning, where a feed-forward network is supposed to transmit its input signals without loss to its output layer, but with one important twist: some intermediate layers of the network contain only few neurons and thus present an information bottleneck. In order to reproduce the network's input signal on its output, the network layers leading up to this bottleneck must find a very low-dimensional, compressed representation of the input (encoding), which the subsequent layer can then decode again. This is illustrated in figure 5.1. By forcing the network to find such a low-dimensional *latent* space representation of its input, we can make sure that the network picks up only on the *most informative* features of its input — or so the story goes.

As we already saw in chapter 4, the same principle can be also applied at a much smaller level, the individual neuron, which needs to reduce its high-dimensional input signal from thousands of incoming synaptic connections to the only one-dimensional output signal with finite bandwidth ([146]; see figure 5.2). Particularly when metabolic constraints and noise are considered, the capacity of individual neurons to transmit information is limited and a loss of information becomes inevitable. If we apply the information bottleneck principle to the single neuron, the neuron should be tuned to ensure that as much (relevant) information as possible about its inputs is preserved in its output. Example 1 gives some intuition for this idea.

5.2 Mutual information and maximum entropy

From an *information theoretical* perspective [7, 113, 147], the neuron represents a *noisy channel*, and the information bottleneck problem is a matter of maximizing the mutual

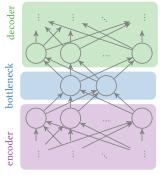


Figure 5.1. An auto-encoder with a narrow bottleneck in between the encoder and decoder layer(s).

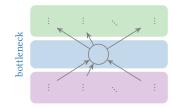
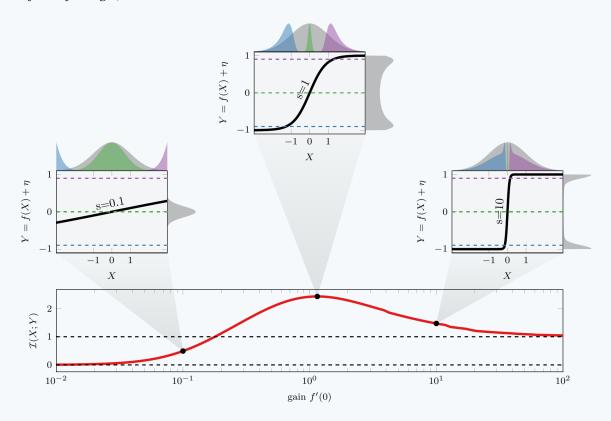


Figure 5.2. A single neuron acts as an information **bottleneck** in between its synaptic inputs and outputs.

Example 1: Gain modulation and the information bottleneck

Consider a linear-nonlinear neuron of the form $f(x) = \tanh(s \cdot x)$ with a single free parameter, the slope s = f'(0) at the origin. We assume that the output signal $Y = f(X) + \eta$ produced by the neuron is corrupted by additive noise η . A perfect recovery of the input signal X from the corrupted output signal is Y impossible, so the neuron becomes a lossy, non-linear noisy channel. How well this channel can transmit information depends on the choice of the gain s. To illustrate this, let's consider three different neurons, one with low, medium and high gain each (see the three insets below, from left to right):



Each neuron receives input with the same prior distribution $X \sim \mathcal{N}(0,1)$ (gray, above each inset), which results in different output distributions P(Y) (gray, right of each inset). If we now observed the three output values $y_1 = 0.9$, $y_2 = 0.0$ and $y_3 = -0.9$ (horizontal dashed lines), we can infer the conditional input probability distributions $P(X|y_i)$ (shown above in corresponding colors). For a neuron with a low gain of $s \approx 0.1$, these conditional input distributions are quite broad and uninformative. For a slope of $s \approx 1$, each of the three outputs encodes a distinct, narrower input distribution. For a very steep slope $s \approx 10$, the observations $y_1 = 0.9$ or $y_3 = -0.9$ again reveal only little about X—mostly just whether it was positive or negative.

The red curve below quantifies this dependence of the neuron's information transmission $\mathcal{F}(X;Y)$ on the gain parameter: As $s \to 0$, f becomes constant, and the transmitted information content approaches 0 bits. In the other extreme, when the nonlinearity approaches a step-function ($s \to \infty$), the output distribution becomes sharply bimodal, and conveys only the sign-bit of the input signal. For some intermediate *optimal slope* ($s \approx 1$), however, the neuron's output yields a maximum of about 2.5 bits of information about its current input. Modulating the gain can therefore help to mitigate the information bottleneck!

information between the channel's in- and output. This critically depends on the statistical properties of the source signals to be transmitted and the noise affecting the channel, as well as the parameters of the channel itself. If we think of the standard linear-nonlinear neuron model with an invertible activation function, then we can express the mutual information as follows:

$$I(X;Y) = I(\bar{Y};Y) = h(\bar{Y}) - h(\bar{Y}|Y),$$

where X is the neuron's input (or membrane potential), $\bar{Y} = f(X)$ is its *noiseless* output, Y is the noisy signal that is ultimately received by the next neuron, $h(\bar{Y})$ is the differential entropy of \bar{Y} and $h(\bar{Y}|Y)$ is the conditional differential entropy of \bar{Y} given that Y has been observed, i.e. the uncertainty of our decoding of \bar{Y} from the noise-corrupted version Y.

In the absence of noise, the channel's capacity to transmit information is only limited by the *source entropy* $h(\bar{Y})$ — this is Shannon's famed first theorem (*source-coding* theorem)[113, 147]. To maximize information transmission by a noiseless channel, we therefore need to use an "*encoding*" $\bar{Y} = f(X)$ that results in a maximum entropy distribution of \bar{Y} .

Note: Subtleties of differential entropy

For the continuously valued case we are interested in, there is one extra caveat to consider [147]: Since the differential entropy $h(\bar{Y})$ can be arbitrarily increased by just scaling \bar{Y} (in fact, $h(\alpha \cdot Y) = h(Y) + \log(|\alpha|)$), the absolute value of the differential entropy is typically meaningless, as it depends on the choice of units and scales of the variables of interest. This is fundamentally different from discrete entropy, which is invariant to *any* invertible transformation![147]. When we talk of maximizing differential entropy $h(\bar{Y})$, we therefore always include direct or indirect constraints on the scale of the random variable and focus not on the absolute value of this maximum, but only on the distribution that achieves it (this only requires comparing differences of differential entropy, in which case the scale-dependent terms cancel.).

In practice, any physical channel is subject to noise that reduces its capacity, and the neuron is of course no exception. This is the core of Shannon's even more famous second theorem (*channel-coding* theorem)[113, 147], which establishes the limit of how much information can be transmitted through the channel in the presence of noise. This upper limit can be increased by improving the *signal-to-noise* ratio, either by allocating more bandwidth to the signal (e.g. scaling up firing rates) and/or by suppressing the noise (e.g. by filtering, see also chapter 4).

Optimizing the encoding while simultaneously taking into account the statistical properties of the source signal and the characteristics of the channel is called *joint source-channel-coding* and can be quite challenging. Luckily for us, the *joint source-channel separation theorem* [147] suggests² that an optimal solution to this problem can be found by separately optimizing the source encoding (which only depends on the distribution of source signals) followed by a channel-specific encoding (which only depends on the characteristics of the channel). In the following, we will therefore limit ourselves to the simpler problem of source-coding, i.e. we'd like to find the distribution of neural outputs \bar{Y} with the largest differential entropy $h(\bar{Y})$ under certain *metabolic constraints* imposed by the channel.

For biological spiking neurons, these constraints could be a finite maximum firing rate of the neuron, an energy constraint on the mean firing rate, or even a constraint that depends nonlinearly on the firing rate. For neuromorphic hardware, they could be finite supply voltages or limits on the energy dissipation.

² The theorem assumes discrete channels, and does not perfectly translate to continuously valued signals [148].

To incorporate such metabolic constraints, we need to find the output distribution P^* with the largest entropy subject to a list of equations or inequalities of the form $E_{P(y)}[g_i(y)] = c_i$ for $i \in I$ and $E_{P(y)}[g_i(y)] \ge c_j$ for $j \in J$. Finding such a measure P^* might seem like a daunting task, but fortunately there is a beautiful solution to this very problem by Jaynes [149], generalizing results attributed to Ludwig Boltzmann. It states that the optimal distribution P^* is always from an exponential family with a probability density p^* that can be expressed directly in terms of the constraints:

 3 To ensure that the result is a valid probability distribution, there is always one additional equality constraint $0 \in I$ with $g_0 \equiv 1$ on the domain of P and $c_0 = 1$.

$$p^*(y) = \exp\left(\sum_{i \in I \cap J} \lambda_i^* g_i(y)\right)$$
 where $\lambda^* = \operatorname{argmax}_{\lambda} \left(\sum_{i \in I \cap J} \lambda_i c_i - \int \exp\left(\sum_i \lambda_i g_i(y)\right) dy\right)$ subject to $\forall j \in J : \lambda_i \geq 0$

In contribution 6, we look at some examples of such maximum entropy distributions. The fact that the resulting distributions are all from some exponential family has a lot of interesting implications and comes in handy for the analysis. We will revisit this in chapter 6 and contrast it to a different approach, which aims to maximize *metabolic efficiency* of information transmission rather than maximizing information transmission under metabolic constraints. Here, we will just continue with the knowledge that we can in principle derive the optimal distribution of the output of a linear-nonlinear neuron under metabolic constraints according to the information bottleneck principle, and that it takes the form of some exponential family distribution.

5.3 Optimal Transport and the Monge Problem

We saw above how the neuron's activation function shapes the neuron's output distribution and thus its ability to transmit information. I then showed how metabolic constraints determine the optimal output distribution. Putting these results together, finding an activation function that produces that desirable output distribution would improve information transmission, and thus constitute a solution to the bottleneck problem. But what would this activation function look like for different input distributions? And what is the best approximation that a neuron could realize?

Again luckily for us, more general versions of this optimization problem, the *Monge-Kantorovich*, *Kantorovich-Rubinstein* or *Optimal Transport Problem* [150], have been studied extensively. It can be loosely paraphrased as the problem to find the 'best' deterministic transformation to map one given probability density onto another given probability density. What 'best' means in this context is precisely defined by a *cost* function $c(x,y): \mathcal{X} \times \mathcal{Y} \to [0,1]$ that penalizes the *transport* of probability mass from x to y. In the historic setting, in which this question was originally posed, c quite literally referred to the cost of moving earth from one spot x to another spot y, which is also where the alternate name *earth mover's distance* originates from [150]. The original Monge problem can be expressed in terms of this cost function as an optimization problem:

$$f^* = \operatorname{arginf}_{f_{\#}(\mu) = \nu} \int c(x, f(x)) d\mu(x),$$

where μ and ν are the source and target the probability distribution, respectively, and $f_{\#}(\mu)$ is the distribution onto which μ is mapped by f (i.e. the push-forward measure of μ under the function f).

In our context, we would like the neuron's activation function to deviate as little as possible from a linear function (in part because this makes the decoding simpler), so we choose c(x,y) to penalize any deviation of y=f(x) from x. The cost function is then some radial function c(||x-y||) that only depends monotonically on the distance between x and y in some norm $||\cdot||$. Under generous assumptions, which our cost function satisfies, ⁴ it turns out that a unique optimum f^* exists and has the following simple form — regardless of the precise choice of c [150, Remark 2.30]:

 4 Both p_X and p_Y must be atom-free, univariate, continuous probability distributions.

$$f^* = F_{Y^*}^{-1} \circ F_X,$$

where F_X and F_{Y^*} are the cumulative probability distribution of X and Y^* , respectively.

Combining this result with the maximum entropy approach above, we therefore know the *optimal activation function* f^* that will maximize the neuron's ability to transmit information for a given input distribution! And since f^* is defined purely in terms of the distributions F_X and F_{Y^*} , any parameter of these distributions becomes a parameter of f^* . Conveniently for us, the optimal function f^* is also an acceptable candidate for an activation function of a neuron, since it is monotonically increasing and continuous if F_X is continuous and F_{Y^*} is injective. ⁵ In contribution 6, I derive the same solution, albeit from a very different perspective, and discuss in more detail the properties of this functional mapping; a related derivation can also be found in [151].

⁵ For every root $p_{Y^*}(y) = 0$, f^* has a discontinuity at $x \in F_X^{-1}(\{y\})$.

Note: Kantorovich's relaxation [150]

The existence of a *deterministic one-to-one* mapping f^* as above is no longer guaranteed if we allow discontinuous probability distributions. But *even if* such a solution exists, the activation function f can become highly nonlinear. If we wish to avoid that while also maintaining the desired output distribution, we can express the optimization problem in a more general form or *relaxation* proposed by Kantorovich [150]. Rather than a deterministic one-to-one function f^* , it defines a probabilistic mapping $\gamma^*: \mathcal{X} \times \mathcal{Y} \to \mathbb{R}^+$, where $\gamma^*(x,y)$ is the relative amount of probability mass to be transferred from x to y.

$$\gamma^* = \operatorname{arginf}_{\gamma \in \Gamma(p_X, p_Y)} \int c(x, y) d\gamma(x, y)$$

Here $\Gamma(p_X, p_Y)$ is the set of all joint probability densities with marginal distributions p_X and p_Y , respectively. Such a map always exists (consider for example the trivial case $\gamma(x,y)=p_X(x)p_Y(y)$), but the optimal map is not (necessarily) the graph of a deterministic function from x to y, i.e. such that $\gamma(x,y)=\delta(x-f(y))$, as the stricter Monge-problem would require. In this non-deterministic case, the same input x is instead "distributed" probabilistically over possible outputs y with distribution $p_{y|x}=\gamma(x,y)$. This means that in order to enforce a desired output distribution, we could also include randomness into the neuron model itself and control it in an input dependent way. But whether this is useful or not is a separate topic I will not discuss here.

Let me summarize these results: in order to be an efficient information channel, a neuron should tune its activation function f to map its input $X \sim P_X$ to an output $Y^* = f(X)$ with maximum entropy distribution $Y^* \sim P^*$. The optimal way to achieve that is to set $f \leftarrow f^* = F_{Y^*}^{-1} \circ F_X$. This is also extremely useful if we are interested in optimizing spikebased information transmission, which we will return to in chapter 6.

5.4 Intrinsic homeostatic plasticity

These derivations all describe the mapping of a single random variable, X, onto a single random variably *Y* by the neuron's activation function *f*, but a real neuron is faced with a continuously varying input signal X(t) which must be expressed as a stochastic process. ⁶ And what if the probability distribution of X(t) were to suddenly change? In order to achieve and maintain the optimal output distribution, we'd expect the neuron to adjust to any changes in its input distribution by homeostatically regulating its nonlinear activation function in real time. This finally brings us to the main topic of this chapter, intrinsic homeostatic plasticity.

If we take for granted that a neuron can approximate the information-theoretically optimal input-output mapping in principle, the challenge for homeostatic plasticity is to keep the mapping optimal, i.e. to constantly adjust the coefficients of the activation function to changes in the environment that affect the input distribution. However, since the current probability distribution over input values is determined by external factors, it must be constantly inferred by the neuron from the recent history of its own input signals. In order to do this with a finite number of variables, the input distribution has to be approximated by some parameterized family of distributions, the time-varying parameters of which have to be estimated online, e.g. by the concentrations of some chemicals or voltage traces.

In contribution 6, I model the neuron's fast-changing membrane potential X(t) by a continuous stochastic process with a given stationary probability distribution from some exponential family, e.g. a Gaussian. Changes due to environmental factors are assumed to occur sporadically on a much slower time-scale, which I model as sudden changes in the stationary distribution of the process. I show that despite the much more complicated mathematics involved with stochastic processes, the intuitions derived above for the probability distributions of in- and output can in fact be applied directly to the stationary distribution of the stochastic in- and output processes. Thus, by transforming the stochastic process that describes the neuron's membrane potential through some nonlinear function f, we can produce a stochastic process with any desired stationary distribution as the neuron's output, including the maximum entropy distribution that solves the information bottleneck problem.

By assuming a parameterized family of both the stationary input and output distributions, the optimal activation function also becomes parameterized. Since we are working with a stationary membrane potential distribution from an exponential family, these parameters are determined by the distribution's so-called sufficient statistics.

These sufficient statistics all take the form of an expected value of some nonlinear function of the process, which we can therefore estimate by filtering, e.g. with an exponentially weighted continuously running average. 7 I prove in contribution 6 that as the running estimates of the sufficient statistics approach the true values⁸, the realized output distribution also approaches the desired output distribution. Since the filtering of the sufficient statistics constitutes a form of running average, the neuron will thus quickly recover from a perturbation to its input distribution! Example 2 illustrates this mechanism for the simple example of Gaussian inputs. While this description has focused on continuous linear-nonlinear models, the same arguments can be extended to spiking neurons, which are known to adapt to their input distributions, e.g. by regulating the spike-threshold [153]. Stabilizing a neuron's output by adjusting to the statistical properties of its input can also promote sparsity and might help explain the emergence of complex cell receptive fields in visual cortex [154]. This form of dynamic re-scaling has been directly observed in biological neurons in vision [155, 156], olfaction [157], audition [121], and might play an important role for neural information processing, in general [6].

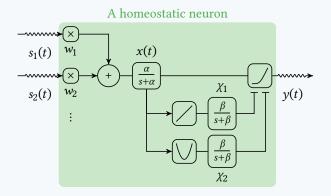
⁶ I only look at *drift-diffusion processes*, which have a stationary distribution from some exponential family [152].

⁷ If the mean value itself is one of these sufficient statistic (as e.g. for the Gaussian distribution), the estimation and homeostatic regulation of that parameter could be realized entirely by dendritic filtering, as discussed in chapter 4.

⁸ They do this in an unbiased way, but with some residual uncertainty that depends on the estimator's timeconstant. The longer the time constant is, the better the approximation becomes at the cost of a longer latency.

Example 2: A homeostatic neuron for Gaussian inputs

With the popular choice of an Ornstein-Uhlenbeck process [152] as a model of the membrane potential X, the stationary distribution is Gaussian with sufficient statistics $s_1 = \mathbb{E}[X]$ and $s_2 = \mathbb{E}[X^2]$. Two internal state variables χ_1 and χ_2 provide a running estimate of s_1 and s_2 , respectively. In order to produce an output with cumulative distribution function F_Y , the neuron nonlinearly transforms its membrane potential through the function f.



The traces χ_1 and χ_2 are used to parameterize the activation function

$$f(x) := F_Y^{-1}(F_X(x)) \approx F_Y^{-1} \left(1 + \operatorname{erf}\left(\frac{x - \chi_1}{\sqrt{2(\chi_2 - \chi_1^2)}}\right) \right)$$

The neuron operates on two time-scales defined by the fast time-constant α of the membrane potential dynamics, and the slower time-constant β of the adaptation process. Any shift in mean or variance of the input distribution is counteracted by the neuron on the slower timescale — the neuron exhibits homeostatic self-regulation.

5.5 The complex interactions of synaptic and intrinsic plasticity

Of course, intrinsic plasticity mechanisms that adjust the neuron's response are not the only form of neural plasticity. The most critical mechanisms for learning appear to be structural and synaptic plasticity [158], which lead to the (dis-)appearance of synaptic connections (or dendritic spines) and an adjustment of the synaptic efficacy, respectively. Each of these forms of plasticity, intrinsic to the neuron or occurring within each synapse, only have access to different information and can thus influence the behavior of neurons in different ways.

Each synapse can, in principle, modulate its transmission strength (or transmission probability) based on the activity of the two neurons it connects, while each neuron, through *intrinsic plasticity*, can only adjust its nonlinearity based on the neuron's membrane potential. To see where a combination of both rules leads, we investigate the dynamics of a neuron's membrane potential and its synaptic weights under the effect of both intrinsic and synaptic plasticity in contribution 6.

5.5.1 Principal Component Analysis

Consider as an example a neuron with two synaptic inputs, which evolve according to a (non-linear) Hebbian rule with weight decay of the form $\frac{1}{\eta}\frac{\mathrm{d}w_{j,i}(t)}{\mathrm{d}t}=f(y_i(t))g(y_j(t))-w_{j,i}(t)$, where f and g are increasing functions, $w_{j,i}$ is the weight of the synapse connecting neuron i

to j and y_i , y_i are the corresponding neurons' activations. In this model, if the outputs of the pre- and post-synaptic neurons stayed constant, the weight $w_{i,i}$ would approach the expected value $\mathbb{E}[f(y_i(t))g(y_i(t))]$ over time. But there is in fact a positive feedback-loop, since an increase in the synaptic weight leads to an increase in the post-synaptic activation, which in turn leads to a further increase of the synaptic weight, and so on. This could potentially lead to unstable runaway dynamics, where the weights all either converge to 0 or diverge to $\pm \infty$. Stable variations of this rule exist for that reason, such as the popular BCM rule [159], which includes a term that adjusts for the neuron's mean activity. Instead, I use the homeostatic intrinsic plasticity of the post-synaptic neuron to the same end, i.e. to maintain a fixed distribution of the neuron's output.

Note: Beyond linear Hebbian learning

The most commonly used synaptic learning rules are (bi-)linear Hebbian learning rules, where the rate of change of the weights is a product of a linear function of pre- and post-synaptic activation. But non-linear dependencies on the pre- and post-synaptic activations are of course conceivable, as well! Such non-linear Hebbian learning rules make it possible to further decouple the effects of synaptic and intrinsic plasticity, e.g. choosing an activation function purely to maximize information transmission in combination with a learning rule to realize principal or independent component analysis. Non-linear Hebbian learning rules therefore open countless more opportunities for synaptic learning rules that could be studied in this framework.

So what happens when we drive an assembly of multiple neurons with a multi-variate stationary input process, and let the synaptic connections and intrinsic parameters evolve according to these synaptic and intrinsic plasticity rules? As we show analytically in contribution 6, without any stabilizing homeostatic plasticity, the weights do in fact diverge. But under the effect of intrinsic plasticity, the weights follow a gradient field and settle in stable fixed-points. For multi-variate Gaussian inputs, these fixed-points correspond exactly to the principal component directions. More accurately, the linear Hebbian synaptic learning rule finds a projection of the multi-dimensional input space onto the one-dimensional membrane potential, for which the expected activation of the post-synaptic neuron is maximized, whereas intrinsic plasticity normalizes its expected activation.

As the input distribution (and thus its principal component directions) changes, the weight vector re-aligns itself and thus counteracts this transformation, thereby realizing a special form of homeostasis.

Independent component analysis 5.5.2

In the previous section, the combination of intrinsic and synaptic plasticity lead to the discovery of principal components, because for that choice of input distribution and activation function, the variance of the input had the largest effect on the expected output of the neuron. But what if we were to choose a different input distribution than Gaussian, where higherorder moments carry important information?

Due to the nonlinear activation function, the expected value of the neuron's output also depends on the higher-order moments of its membrane potential (see the appendix of contribution 6). For example, with a monomial activation function $f(x) = x^n$, the neuron's mean output measures the n-th moment of the input distribution. With n = 2, such a neuron would be sensitive to the *variance* of the input, and to the *curtosis* for n = 3. In general, it depends on the Taylor expansion of the activation function, how much each moment of the input distribution influences the neuron's mean output. If we choose a different activation function or input distribution, the neuron can therefore discover other subspaces that maximize higher-order moments of the input distribution, instead. This can be used to disentangle signals that are *uncorrelated*, but not *independent*, because they do share higher-order correlations. In analogy to principal component analysis, this procedure is therefore called *independent component analysis* (ICA). Our combination of intrinsic and synaptic plasticity mechanisms produces either principal or independent component analysis or a mixture thereof, depending on the input distributions and/or activation function!

⁹ The original work by [160] introduced independent component analysis using a similar, neuro-inspired motivation with the activation function $f(x) = x^3$ to maximize curtosis.

5.6 Applying the information bottleneck to neural assemblies

We can generalize these ideas from individual neurons to neural assemblies: By selecting and scaling the inputs into the neurons, synaptic plasticity determines how the neurons' inputs are related to each other, whereas intrinsic plasticity independently controls the marginal distribution of each individual neuron's outputs.

This raises an interesting question: how much control over its *joint output distribution* could an assembly of neurons theoretically exert, if the only free parameters are each neuron's nonlinearity and the incoming synaptic connections? Can an assembly of neurons map an arbitrary multi-variate input distribution onto an arbitrary multi-variate output distribution? The general answer is no¹⁰, but to make this more precise, we have to disentangle the effects of synaptic and intrinsic plasticity. In contribution 6, we do this by introducing a concept from probability theory called *copula* ([161], see also the note below).

¹⁰ Consider e.g. that a uni-variate input signal cannot be transformed into multiple independent output signals.

Note: Copulas describe the coupling of random variables

For an assembly of N neurons with individual inputs $X_i \sim P_{X_i}$ with joint distribution P_X and outputs $Y_i \sim P_{Y_i}$, we define the intermediate random variables $U_i = F_{X_i}(X_i)$, called the *ranks* or *quantiles* of X_i , each of which is marginally uniformly distributed. The joint distributions of these rank-variables is the copula

$$C(u) := F_X(F_{X_1}^{-1}(u_1), F_{X_2}^{-1}(u_2), \dots, F_{X_N}^{-1}(u_N)),$$

a probability distribution in the N-dimensional unit cube, that captures how the random variables X_i are related — regardless of their marginal distributions! A common application of this rank-transformation in statistics is when random variables need to be compared across different scales. In that case, the correlation between the ranks of the variables can be used, which is just the correlation of the copula. The copula has many more interesting theoretical properties, e.g. [162], but most importantly for us, it is invariant under any invertible univariate transformations of the individual random variables X_i . Therefore, the copula C of the assembly's multi-variate input and its multi-variate output distribution coincide and we can just talk of the copula of the assembly. The copula is only a property of the synaptic connections and the input distribution, and unaffected by the neuron's nonlinearity.

Using the copula C, we can factorize the stationary joint probability distributions of the membrane potentials X and activations $Y = f^*(X)$ as follows:

$$F_X(x) = C(F_{X_1}(x_1), F_{X_2}(x_2), \dots, F_{X_N}(x_N))$$

$$F_Y(x) = C(F_{Y_1}(y_1), F_{Y_2}(y_2), \dots, F_{Y_N}(y_N))$$

The last factorization specifies the population's joint output distribution in terms of the desired individual marginal distributions F_{Y_i} of each neuron's output, which can be enforced by intrinsic plasticity, and the copula function C, which captures the co-dependency between the neurons' activity. The copula is invariant under element-wise invertible transformations, and therefore only depends on the synaptic connections — not the activation function. In other words: No matter what activation functions we choose, we can only modify the marginal distributions of each neuron's output —but not the assembly's copula— through intrinsic plasticity! Synaptic plasticity, on the other hand, can shape how different signals are combined by the individual neurons, and thus it can influence the assembly's copula, but not the activation function.

In general, C can be arbitrarily complex, and there is little hope that it can be fully controlled by setting the synaptic weights alone. 11 For copulas that are parameterized by more than one parameter per synapse, for example, synaptic plasticity alone is obviously insufficient to fully control the copula. But e.g. for jointly Gaussian inputs, the distribution is fully parameterized by the covariance matrix (and the mean), which can be shaped arbitrarily by an appropriate choice of synaptic weights (and intrinsic plasticity).

If we apply the information bottleneck argument now to an entire assembly of multiple neurons, the neurons should jointly maximize information transmission. This can be achieved if the neurons' outputs are i.i.d. with a marginal maximum entropy distribution.

It therefore seems reasonable from an information bottleneck perspective, that the individual neurons should encode different independent (or principal) components. One way to ensure this is mutual decorrelation of the neurons within an assembly by lateral inhibition in order to enforce the learning of different weights. We demonstrate that this leads to the unsupervised clustering of the MNIST handwritten digits dataset by the extraction of independent components. For a second control dataset composed of random image patches no such independent components should exist, and indeed the same setup leads to the discovery of the dominant principal components instead. The same idea, sometimes called blind source separation, can be generalized to other types of input signals as well, in particular to separate heavy-tailed12 source signals (see also example 3).

There is plenty of biological evidence to prove that such a decorrelation of signals by PCA/ICA also occurs in nature. For example, this can be observed on a very low level of the mammalian visual system (see [7, chapter 5] for a great summary of this topic). There, colors are sensed by receptors tuned to different (but overlapping) spectra of visual light, but the signals that are transmitted by ganglion cells appear to be linear combinations of these "raw signals": instead of a 'red', a 'green' and a 'blue' channel, the spike-trains transmitted over the optic nerve seem to represent a 'blue-(red+green)' difference, a 'red+green' sum and a 'red-green' difference channel! This representation decorrelates the highly correlated responses of the individual color channels, and results in a more information theoretically and metabolically efficient code. Just as in our hypothetical example, this requires an appropriate rotation of the synaptic weight vectors and an appropriate scaling of the neuron's nonlinear activation function, although that may be genetically predetermined, rather than learned, in this specific case. A similar observation can be made in the olfactory bulb of zebrafish larvae [164], where appropriate lateral inhibition decorrelates the neurons' responses.

Plasticity is information processing

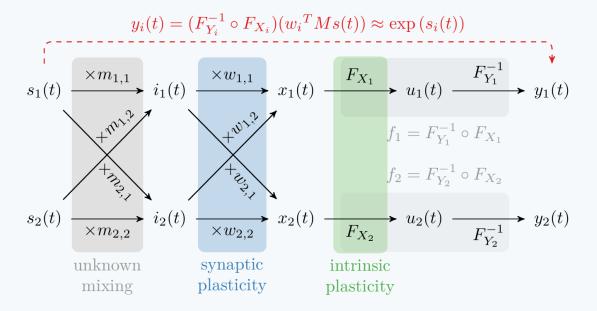
In the field of machine learning, we think of neural networks in terms of a training phase, where the network is optimized, and an inference phase, where the trained network is used to process information. As I tried to show in this rather long chapter, this perspective completely misses the important role that plasticity mechanisms play in information processing, in

¹¹ Of course, it may still be possible to control the copula through synaptic weights by using multiple layers of neurons, as is done e.g. for so-called normalizing flows [163].

 $^{^{12}\,\}mathrm{I}$ use this term to refer to random variables with larger higher-order moments than a normally distributed variables with equal variance.

Example 3: Blind-source separation with synaptic- & intrinsic plasticity

Suppose we want to "de-mix" two independent source signals $s_1(t)$ and $s_2(t)$ from two different mixtures $i_1(t)$ and $i_2(t)$. This could be two independent sound sources that reach our two ears with different attenuation, or it could merely be two correlated outputs of neurons in a previous layer — in either case, the mixture coefficients $m_{i,j}$ are not explicitly known. These two input signals are then transmitted through synaptic connections with weights $w_{i,j}$ to the two neurons, where they are integrated into the membrane potential $x_1(t)$ or $x_2(t)$, respectively (we ignore the temporal filtering of the membrane potential here). Each neuron i then applies its activation function f_i to produce the output $y_i(t)$. We'd like each neuron to reproduce a nonlinear function $\exp(s_i(t))$ of just *one* of the input signals. This is called *blind-source separation* or independent component analysis.



The interaction of synaptic plasticity and intrinsic plasticity can solve this problem in an unsupervised manner. To ensure that $f_i(x) = (F_{Y_i}^{-1} \circ F_{X_i})(x) = \exp(x)$, we assume X_i has a probability distribution from the same family as S_i , and set $P_{Y_i} = \exp_\#(P_{S_i})$. The greedy mechanism of synaptic plasticity then finds a weight matrix W that inverts the unknown mixing matrix M, while intrinsic plasticity stabilizes this process and ensures that the outputs have the desired distributions F_{Y_i} . The intermediate variable $U(t) = (u_1(t) \ u_2(t))^T$ is marginally uniform, and its joint distribution is the *copula* of X_1 and X_2 or Y_1 and Y_2 , respectively.

particular if we consider the dynamics of online learning that has to happen in real-time, such as homeostatic intrinsic plasticity and synaptic plasticity. Since the interaction of intrinsic and synaptic plasticity can help not just to stabilize the behavior of neurons and networks, but also to extract, compress and track relevant information like principal or independent components in high-dimensional signals, I'd consider them to be information processing mechanisms in and of themselves. While the high-level discussion above was focused on continuous linear-nonlinear neuron models, we will apply these concepts to spiking neurons, as well, in chapter 6.

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Dr. Ian Malcolm in Jurassic Park

Rate-coding with spiking neurons

The neuron models we looked at so far all use real-valued, continuous output signals as a proxy of the neuron's current firing rate. But — electric gap-junctions aside — biological neurons in the human brain communicate via chemical synapses that actually have to generate individual *spikes* to communicate. This mode of communication has been known for almost two centuries [165], but to this day it plays only a minor role in machine learning models of neural networks. Why is it, that biological neurons send spikes, rather than continuous signals? And why haven't we seen more applications of spiking neural networks in machine learning?

There are two fundamentally different schools of thought on this issue. The *rate-coding* paradigm assumes, that the only relevant information conveyed by a spike-train is the time-varying rate at which the spikes are generated, while the *spike-(time-)coding* paradigm treats each individual spike as a symbol, the timings of which convey individual pieces of information. Unsurprisingly, how and how well a neuron can represent its input and whether rate-coding is a viable model for that, has been one of the oldest research questions in theoretical neuroscience [166], and has been revisited many times (see e.g. [167, 168]). Surprisingly, there is still a lack of conclusive biological evidence one way or another and the distinction between the two is not always clearly cut [169], so this apparently simple question hasn't been settled even after decades of intense debate. To better understand what rate-coding entails, I will therefore stick to an entirely theoretical view of rate-coding in the spirit of [166], and analyze its capacity to encode and transmit information for two of the most common neuron models. In chapter 7, I will then try to account for more recent biological evidence, which will lead us to spike-time coding, or rather *event-coding*.

6.1 Why do (only) biological neurons spike?

For proponents of rate-coding, the additional complexity of spike-based communication can be understood as a biological "implementation detail" of sorts: Pulse-based communication offers a noise-robust and energy-efficient means to approximately convey a continuous, real-valued signal (the *firing rate*) under metabolic constraints over what is essentially a binary channel (the neuron's axon with its chemical synapses). The continuous signal is then represented by the *rate* or *density* of the pulses per unit time-interval. This form of encoding is simple to implement and very reliable, which is why variations of this scheme are also used in digital electronics to transmit inherently analog signals (e.g. audio signals or servo-motor controls) over a digital connection. ¹

 1 In electronics, there is in fact a corresponding pulse-based encoding scheme for each of the spike-based communication paradigms (see also [3, chapter 3]): pulse-density modulation (PDM) and its variants, which go by the names pulse-frequency modulation (PFM) and pulse-code modulation (PCM) correspond to rate coding. Pulse-position modulation (PPM) corresponds to spike-time coding, which we will discuss in chapter 7. Asynchronous $\Delta\Sigma$ modulation is directly related to integrate-and-fire neurons and time-encoding-machines [107].

6

Spike-based communication also shares another benefit with digital electronics: While analog signals strongly attenuate as they propagate along the neural membrane (see also chapter 4), binary spikes can be detected over long distances and regenerated to their full amplitude by error-correcting mechanisms. In cortical neurons, this happens at distinct locations along the axon called *Ranvier nodes* [6], ² which are separated by highly myelinated (i.e. electrically insulated) stretches of the axon. This insulation not only reduces leakage but also greatly increases conduction velocity. The resulting *saltatory propagation* of action potentials allows for an extremely fast, reliable and energy efficient communication over long distances without degradation, which is critical for coordinating motor activity in limbs far away from cortex [7, 139] and plays a crucial role in the consolidation of memory [139].

In short, from the rate-coding perspective, spike-based communication is a very useful adjustment to bio-physical constraints. But as long as a spiking neuron's firing rate can be well approximated by a nonlinear function of its input, the precise mode of communication makes little conceptual difference; the overarching framework is still function approximation by linear-nonlinear neurons, and spiking neurons are merely a hardware-efficient implementation (or approximation) thereof. In fact, as we shall see below, there is a direct correspondence between continuous linear-nonlinear neurons and simple spiking neuron models that makes it trivial to convert back-and-forth (as long as we are only concerned with mean firing rates).

6.2 Encoding continuous signals into rate-coded spike-trains

For the rate-coding paradigm, any benefit of spiking neurons has to come not from increased computational power, but rather from an increased *metabolic efficiency*, i.e. the amount of information transmitted per Joule of energy spent. To discuss the capabilities and limitations of rate-coding, it's therefore important to understand how and how well time-varying signals can be encoded into a series of spikes, in the first place.

Since a neuron will have to be able to decode the relevant signal again from the spike train, typical requirements for spike-based codes are as follows: The current firing rate of a neuron is only a function of its recent input 3 ; a neuron's response is time-equivariant, i.e. a temporal shift in the input results in a corresponding shift in the output; each spike is represented by a brief stereotypical pulse with identical mass⁴; and despite the *non-linear encoding*, a *linear decoder*, i.e. a filter, must be sufficient to decode the continuous rate from the spiking signal⁵. This linear 'decodability' imposes a hard constraint on the sort of spike-patterns that can be used to convey information, but there is biological evidence to support this simplifying assumption⁶. Since the spike-encoding mechanism might induce filtering effects that are irreversible, as we have seen in chapter 4, we will be satisfied if we can encode a signal s(t) into a spike-train z(t) and are then able to recover $(\kappa * z)(t) = f((\psi * s)(t))$ for some kernel ψ and some invertible function f from the spike-train by the linear decoder with kernel κ . These constraints still leave room for many different mechanisms to encode a continuous signal into a discrete sequence of spikes, but I will only consider three particularly interesting types of rate-based encodings.

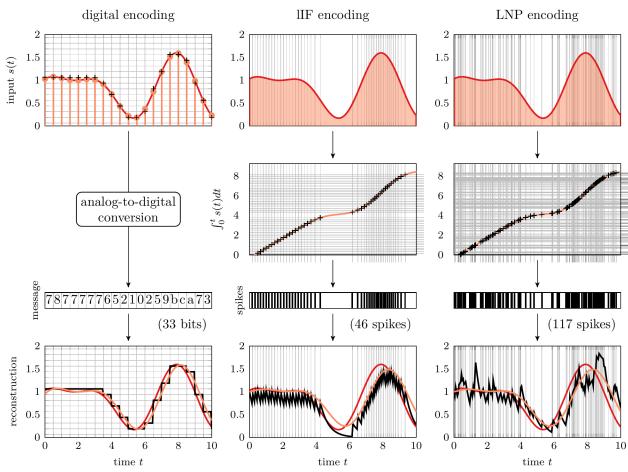
Since each spike requires energy to generate and transmit, we can compare these different approaches by the number of spikes they tend to generate, and how well we can decode the underlying continuous signal from the spike-train.

6.2.1 Periodic sampling and digital transmission

To put the encoding capabilities of spiking neurons into perspective, let's compare them to a well-known reference: digital encoding schemes. In the signal processing domain,

- ³ Adaptation effects as in chapter 5 are sometimes included as well, but on a much slower timescale.
- ⁴ Typically, a Dirac $\delta(t)$ is chosen for continuous time and a Kronecker δ_{t} /rectangular pulse for discrete-time models.
- ⁵ This requirement is the *first principle* of the *neural engineering framework* [5]; see also chapter 2.
- ⁶ Pairwise correlations between spikes (which can be assessed with a linear filter) carry most of the information content in individual neurons' spike-trains [170].

² Again, there is a rich analogy to manmade electrical communication systems: while the strong attenuation of analog signals along wires originally made the transmission of analog signals (such as the telephone) over large distances difficult, the digital nature of the telegraph allowed for a regeneration of the signal at periodically spaced relay stations, much alike Ranvier nodes, and thus enabled fast long range communication. A nice account of this development can be found in [113].



continuous real-valued signals are typically measured at periodic time-intervals, which produces a discrete sequence of real-valued samples. According to the Nyquist-Shannon sampling theorem [113], any bandwidth-limited signal can be losslessly represented this way if the *sampling rate* is sufficiently high. Each real-valued sample can then be approximated by *n*-bit binary number, and the active bits can be transmitted as brief pulses via *n* parallel wires. If we scaled each of these pulses (or bits) by the corresponding power of two, summed and filtered them appropriately, ⁷ we'd recover the continuous signal. While this binary encoding would hence (with a little stretch of the imagination) satisfy our requirements of a rate-based encoding, it is of course not actually a viable model of neural spike-based communication. But it does provide a theoretically optimal reference implementation for the pulse-based transmission of bandwidth-limited continuous signals, against which we can measure the performance of other, more plausible rate-coding schemes.

6.2.2 Rate-coding with (leaky-)integrate-and-fire neurons

Biological neurons generate spikes through a cascade of opening and closing ion channels, which modulate in- and outgoing ion currents that in turn drive the neural membrane potential. This complex biological mechanism is described by the famed Hodgkin-Huxley model [50], but much simpler models suffice if we are only interested in capturing the encoding of continuous signals into spikes. The simplest of these is the ubiquitous *integrate-and-fire* model [3, 171, 172], which operates by integrating the input signal up to a critical threshold, where it resets and fires a spike.

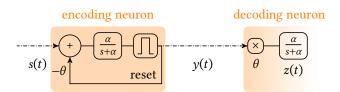
Figure 6.1. The same input signal (top row, solid red) is encoded by a 4-bit digital code (left column), an LIF neuron (middle column) and an LNP neuron (right column) over a 10s interval. The LIF neuron produces spikes at (almost) even increments of the signal's integral, while the LNP neuron fires at uniformly distributed signal integrals (second row). The digital code results in a parsimonious representation that here requires only 33 active bits. The LIF neuron requires 46 spikes for a slightly worse reconstruction (bottom row, black line) of the filtered input signal (orange line), while the LNP neuron gives a bad approximation even for 117 spikes.

⁷ The optimal kernel, a sinc function scaled and stretched according to the sampling frequency, is acausal, but an approximate solution can also be obtained with a causal kernel. See also chapter 4.

This can be theoretically "justified" as follows: For a continuous signal s(t), the firing rate of the neuron should encode the signal in a way that can be linearly decoded by filtering (see also chapters 2 and 5). Therefore, the firing rate ought to be proportional to s(t), i.e. the average number of spikes in an interval $[t_1, t_2]$ should be proportional to $\int_{t_1}^{t_2} s(t) dt$.8 For the neuron to fire exactly one more spike, the average time-interval between the previous spike at time t_1 and the new spike at time t_2 should therefore be $c \int_{t_1}^{t_2} s(t) dt = 1 \Leftrightarrow S(t_2) = S(t_1) + 1/c$, where $S(t) = \int_{-\infty}^{t} s(\tau) d\tau$ and c is a constant of proportionality. In other words, the neuron should fire a spike whenever S(t), the integral of the signal s(t), increases by more than the threshold $\theta = 1/c$ over its value at the previous spike. Of course, this is exactly the mechanism implemented by the standard integrate-and-fire neuron, which integrates its input s(t) up to the critical threshold θ , where the opening of voltage-gated channels resets the neuron back to its resting potential (here chosen as 0 for the sake of simplicity), and the process begins anew. Such an encoding is also called a send-on-delta scheme, as a spike is emitted whenever there is a significant change ("delta") in the (integral) of the signal.

This allows us to say something about the timing of the spikes in relation to the signal: If we assume a positive input signal s, then S(t) is a monotonically increasing (and hence invertible) function. Whenever $S(t) = S(t_i) + 1/c$, a new spike t_{i+1} is generated, therefore the spike times t_k satisfy $S(t_k) = k\theta$ and thus $t_k = S^{-1}(k\theta)$. This spike-train is linearly decodable by construction, and we can easily verify that this also meets our other requirements of a rate-code, since a delay in s leads to an equal delay in s and thus in t_k .

The ideal integral operator in this construction would require the membrane potential to remain constant in the absence of external inputs. Not only do inevitable leakage currents make this implausible for both biology and neuromorphic hardware, but it also has the undesirable theoretical implication, that the output of the neuron depends on a potentially infinitely long history of inputs. 9 To remedy this, the integral operator, whose impulse response is a step-function, is often replaced by a low-pass filter with an exponentially decaying impulse response. This results in the more biologically plausible and very popular leaky integrate-and-fire (LIF) neuron [3]. If the time-scale of the exponential filter is very short, it approaches a Dirac- δ kernel and the neuron acts like a coincidence detector of nearly simultaneous spikes, to which we will return in chapter 7. For a very long time-scale, on the other hand, the exponential filter approaches a step-function and the model converges to the pure integrate-and-fire neuron. As theoretical considerations and biological observations show (see e.g. chapter 3 of [7]), the optimal trade off between these two extremes in terms of metabolic efficiency (i.e. how many bits are transmitted per Joule spent) seems to be achieved when the filter's time-scale roughly equals the expected inter-spike interval. Other filters than the exponential could be used as well (see also the node below), but will not be further discussed here.



Let's consider the example shown in figure 6.2 of a pair of LIF neurons with the dendritic filter $\kappa_{\alpha}(s) = \frac{\alpha}{s+\alpha}$. Suppose we'd like to recover the filtered signal $(\kappa_{\alpha} * s)(t)$ from the decoding neuron's membrane potential z(t). We know that the residual $(\kappa_{\alpha} * s)(t) - (\kappa_{\alpha} * y)(t)$ between the filtered input signal and the filtered spike-train is bounded between 0 and θ , because whenever the error exceeds that bound, another spike is generated, resetting the error back to 0. Under a few simplifying assumptions (see appendix B.1), this residual has a mean value of

 8 The average number of spikes in a small interval $[t-\Delta t/2,t+\Delta t/2]$ of length Δt should be approximately proportional to $s(t)\cdot \Delta t$. Partitioning the interval $[t_1,t_2]$ into strips of width Δt and calculating the Riemann sum for $\Delta t \to 0$ gives this result.

Figure 6.2. Two simple leaky integrate-and-fire neurons as used in figure 6.1, one of which receives a time-varying continuous signal s(t) as its input and encodes it into a spike-train y(t). The second neuron decodes the spike-train into piece-wise continuous membrane potential trace z(t). Both use the same exponential kernel $\kappa_{\alpha} = \frac{\alpha}{s+\alpha}$ with rate α .

⁹ This would e.g. violate the fadingmemory assumption of reservoir computing [30].

 $\approx \theta/2$ and a root-mean-squared error (RMSE) of $\approx \alpha\theta/\sqrt{12}$. By reducing θ (and thus increasing the firing rate) or α (and thus increasing the filter's time-constant), we can therefore reduce the error bound arbitrarily and get uniform convergence $\lim_{\theta \to 0} z = s * \kappa_{\alpha}$.

For a constant signal $s(t) = c \ge \theta$, the neuron's firing rate thus scales almost ¹⁰ linearly with c, whereas the expected RMSE remains constant across almost the entire input range of the neuron. No spikes at all are generated for $c \leq \theta$, so the mean firing rate as a function of the constant input *c* approximates the rectified-linear unit (ReLU) $f(c - \theta/2) \approx \max(0, c - \theta/2)$. See also appendix B.1 for a derivation.

As figure 6.1 shows, the LIF neuron is capable of reliably encoding a time-varying signal into a single pulse-train. Because of its simplicity, a very similar mechanism is also commonly used in signal processing under the name $\Sigma\Delta$ or $\Delta\Sigma$ modulator [108] or just integrate-and-fire sampling [173] to convert continuous signals into pulse-trains¹¹.

Note: Filter-and-fire neurons

The (leaky) integrator represents only one specific type of filter that a neural dendrite could implement (see chapter 4). By substituting in various other kernels, the integrate-and-fire model can thus be generalized to a very interesting class of filterand-fire models [3]. The leaky-integrate-and-fire (LIF) model with an exponential kernel shown here is a particularly popular example, since it can be motivated from biological first principles and can be implemented very efficiently by a single firstorder low-pass filter. But also second-order filters like the α -kernel, a convolution of two exponential kernels, are used to model the combined effect of filtering by the chemical synapse as well as the neuron's membrane potential [3]. Such a higherorder filter could help remove the high-frequency noise otherwise introduced by the discontinuous jumps that result from filtering a spike-train with a first-order filter. Naturally, the choice of kernel has strong implications for the behavior of the neuron, and all the arguments from chapter 4 apply to spiking neurons just as well.

6.2.3 Stochastic encoding

A rather different approach to rate-coding utilizes stochasticity. A linear-nonlinear-Poisson (LNP) spiking neuron [174] fires spikes according to an inhomogeneous Poisson process [175] that uses the input signal s(t) as its time-varying *rate*. The resulting spike-times are stochastic, but the expected number of spikes per time-interval $[t_1, t_2]$ is proportional to the integral $\int_{t}^{t_2} s(t) dt$, just like for the LIF neuron above. But in contrast to the LIF neuron, the spike-times in that interval are independently and identically distributed with cumulative distribution function S, i.e. $t_k \sim S^{-1}(u_k)$ where u_k is a uniform random variable. ¹² In fact, this property allows us to elucidate the key difference between the deterministic integrate-and-fire and the stochastic LNP model: While the spike-times of the IF neuron contain no information besides the signal s (i.e. the spike-times t_k are deterministic given s), the spike-times of the LNP neuron also encode noise (i.e. the spike-times t_k are randomly distributed with a cumulative distribution function proportional to S). A different way of looking at the same phenomenon is to view the LNP neuron as equivalent to an IF neuron with exponentially distributed random threshold¹³, or subject to a corresponding distribution of noise on the membrane potential. All other things being equal, the LNP neuron is therefore likely to achieve a much worse signal-to-noise ratio, as we shall also see below.

Let's look at an example of LNP neurons in figure 6.1. Just like in the case of the integrateand-fire neuron above, the signal can be linearly decoded by filtering the spike-train with an

¹⁰ This approximation of the mean and RMSE fail when the mean input to the neuron goes below θ .

¹¹ The main difference is, that a $\Delta\Sigma$ modulator encodes both positive and negative changes of the signal into the rising and falling edges of a binary pulse-width-modulated signal, whereas the LIF mechanism encodes only positive changes into spikes and relies on the passive leakage for decreasing the signal.

¹² Recall that in (L)IF neurons, the latent variable u_k would instead be (almost) regularly spaced at $u_k = k/N$ for N spikes.

¹³ Since spikes are generated at uniformly distributed levels of S, the increments from one spike to the next are exponentially distributed.

exponential kernel. One can show (see appendix B.2), that exponentially filtering the spike-train provides an unbiased estimate of the signal with an RMSE that approaches $\sqrt{\alpha c/(2\lambda)}$. Like for the LIF neuron, the expected firing rate response of the LNP neuron to constant input is therefore given by a rectified-linear function, but unlike the LIF neuron, the RMSE actually grows with c.

Key benefits of the stochastic approach and the main reason for its popularity are the possibility to incorporate noise and its conceptual simplicity, which allows it to be trivially extended to assemblies of multiple neurons. Deterministic LIF neurons, for contrast, can show phase-locking and other specific dynamics, or may fail to fire all together if the input is sub-threshold, whereas LNP neurons respond linearly across the entire input range. The stochasticity of the LNP neuron could thus actually *enhance* information transmission in some specific cases, but it generally comes at considerable expense in others, as we'll quantify in section 6.4.

6.3 Rate-coding neurons are linear-nonlinear neurons

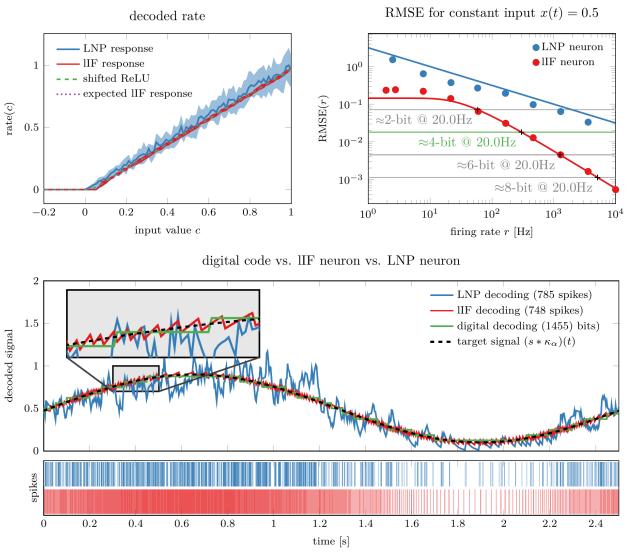
We already saw above that the mean firing rate of spiking neurons can be modeled as a function of their (constant) input signals. Conveniently, this function takes the rectified-linear form for both LIF and LNP neurons, one of the most popular choices of activation function in current deep neural network architectures. ¹⁴ By making the firing rates sufficiently large, ¹⁵ an arbitrary accuracy (i.e. an arbitrarily low RMSE) can be achieved. If we also choose the dendritic filters' time-constants appropriately, a trained deep neural network can be trivially translated into a spiking neural network simply by replacing each continuous neuron with one accordingly configured spiking neuron — $et\ voil\dot{a}$, we have a trained deep spiking neural network! A direct conversion of this sort has been shown to work even for very large, state-of-the-art network models [176, 177]. The same idea can be applied to recurrent networks and reservoir computers, as well (see also [5]). As the success of this one-to-one conversion confirms, rate-coding really is merely a different implementation of the continuous function approximation paradigm discussed in chapter 2.

6.4 How good is rate-coding for transmitting information?

If we adopt the rate-coding perspective, the single purpose of spike-based communication is to transmit analog signals reliably under biological constraints. But which mechanism works best? How does it fare in comparison to a purely analog implementation? And how does rate-coding with spikes compare to conventional *digital* sampling schemes that are used to simulate deep neural networks? Are rate-coding spiking neural networks a viable machine-learning alternative to conventional deep neural networks?

I'll attempt to (partially) answer these questions here, starting with the example shown in figure 6.1. There, we saw three fundamentally different coding schemes that all represent a continuous signal by discrete series of binary pulses, but they yield rather different results. Using the digital "neuron" as a reference, I'll compare, how efficiently the LIF and the LNP neuron can encode information into a spike-train. Figure 6.3 shows the RMSE of both neuron models when encoding the constant signal s(t)=c=0.5. As we systematically vary the neuron's firing rate by varying the LIF neuron's threshold θ and the LNP neuron's gain λ , we can observe a consistently and substantially lower error for the LIF neuron than for the LNP neuron. Besides the fact that both the LIF and LNP neuron follow the rate-coding approach and use the same exponential kernel with rate $\alpha=40Hz$ for decoding, the LIF neuron makes much better use of the precise timing of each spike, and encodes the continuous input signal more effectively into a spike train.

- ¹⁴ This is of course hardly a coincidence, since the rectified-linear activation function was in fact modeled after its (spiking) biological inspiration.
- 15 For the (L)IF neuron, this can be achieved by lowering the threshold $\theta,$ for the LNP neuron by raising the gain $\lambda.$



We can quantify this more accurately with the help of information theory (see appendix B.3 for a derivation), which shows that encoding a (constant) signal with the same specified accuracy ϵ (defined by the differential entropy of the residual) will require a quadratically larger number of spikes for the LNP neuron than for the LIF neuron as we increase ϵ !

But how do these simple rate-coding spiking neural network models compare against an analog implementation or their digital counterparts from deep learning? Not too well, unfortunately: As figure 6.3 already shows, even for firing rates as high as 10,000Hz, the LIF neuron in this setup only reaches an accuracy less than that of a 10bit signal sampled at 20Hz. At a more reasonable 250Hz firing rate, the rate-coding LIF neuron barely matches the accuracy of a 4-bit signal sampled at a rate of 20Hz, resulting in an effective information content of less than a quarter bit per spike. The LNP neuron fares much worse than that.

As shown above, this disparity is in large part due to the poor scaling of the accuracy with the number of spikes, which is linear in the firing rate for the LIF neuron and only scales with the square root of the firing rate for the LNP neuron. For contrast, the accuracy of a digital code grows *exponentially* with the bit-depth of the signal! This is bad news for rate-coding with spiking neurons, since most deep learning models currently make use of the much more accurate half-precision (16-bit) or quarter-precision (8-bit) floats or integers, with only few networks quantized to precisions as low as 4-bit or below [73]. In the rate-coding

Figure 6.3. Top left: The normalized mean firing rate as a function of the constant input *c* matches the (shifted) ReLU activation function max(0, x b) closely, where b = 0 for the LNP neuron and $b = \theta/2$ for the LIF neuron. The interquartile range of the LNP neuron's firing rates (in blue), estimated over 50 trials, grows with \boldsymbol{c} and is much larger than for an LIF neuron (in red). Top right: The RMSE as a function of the firing rate for constant input c = 0.5 shows, that the LIF neuron achieves much lower errors at equal firing rates than the LNP neuron. Bottom: Spike-trains and decoded signals in response to a slow varying sine-wave (black dashed line). For reference, a 4-bit signal sampled at 20Hz is included (green).

context, SNNs also offer no qualitative benefits in terms of raw computational power; to the contrary, they are merely used to approximate the behavior of DNNs and can therefore not be expected to surpass their performance. ¹⁶ Any benefit of rate-coding in biology, machine learning or neuromorphic hardware must therefore come from a more efficient *physical implementation*, rather than an information theoretical argument.

¹⁶ It is however possible that spiking neural networks have intrinsic biases that prove beneficial, e.g. if they had a regularizing effect of sorts, but I'm not aware of any proof of that.

6.5 Optimal rate-coding under metabolic constraints

So far, we focused on the encoding accuracy of spiking neurons (measured by the RMSE) for some given constant signal. In the language of information theory, this corresponds to the problem of channel coding. But if we want to fully assess the neuron's ability to transmit information, we need to consider source-coding, as well. In section 5.2, we approached this from the information bottleneck perspective: to make the neuron's output as informative as possible, we opted to maximize its entropy (measured in bits/second) while respecting the metabolic constraints imposed on the neuron. This leads to the most powerful neuron that the energy budget allows, but in a realistic setting that may not be the best solution overall. Here, we will instead try to make the best possible use of the energy by optimizing the metabolic efficiency $\varepsilon := h/\cos t$ instead, i.e. the entropy of the neuron's output in relation to the required power cost (measured in bits/Joule). This alternative approach leads to a less powerful but more parsimonious neuron, which is a worthwhile trade-off if energy, rather than the number of neurons, is the most critically limited resource. In general, there is a four-way trade-off for spiking neurons between raw performance on the one hand, and firing rates ('paying with spikes'), the complexity and size of neurons, synapses and circuits ('paying with hardware'), and metabolic costs ('paying with power') on the other [7]. If we take into account, that the human brain demands almost 20% of the body's entire energy budget [178] and almost 80% of that energy is directly spent on the firing of spikes [178, 179], and scales linearly with the mean firing rate [178, 180], the enormous evolutionary benefit of increased metabolic efficiency becomes obvious. And indeed, the general tendency for biological neurons seems to be optimization of metabolic efficiency [6, 7]. The idea of such an economy of impulses goes back at least to [181] and was formalized by [182] for populations of neurons. We'll briefly look at what this implies for the single rate-coding neuron.

One important observation is, that under rather mild assumptions (see example 4) the metabolic efficiency is maximized for a unique optimal firing rate μ^* that strikes a good balance between the inevitable *static* power consumption of the neuron, which occurs regardless of the neuron's firing rate, and the *dynamic* power consumption due to the generation of spikes. A brief back-of-the-envelope calculation in example 4 using parameter estimates from real neurons puts this optimal rate at a surprisingly low mean firing rate of around 1.41 spikes/s — much lower than what cortical neurons are capable of, but very well in line with the distribution of firing rates observed *in vivo*. ¹⁷

Which ever firing rate distribution offers the best trade-off, a neuron could achieve and maintain it with an appropriate activation function and homeostatic plasticity, as I argued in chapter 5. For the LNP neuron, this can be implemented by *explicitly* making the instantaneous firing rate a nonlinear function of the membrane potential. This is more complicated for deterministic (L)IF neurons, because their effective nonlinearity is only *implicitly* defined by the neural dynamics, but the effective firing rate function can be influenced by various indirect means such as filtering (see chapter 4) or nonlinear dependencies between the membrane potential and the input [183] or between the threshold and the membrane potential [184]. In either case, a homeostatic mechanism like in chapter 5 could help achieve and

 $^{^{17}}$ The mean firing rate of human cortical neurons is estimated at 1.15Hz with a range from 0.5 – 2.0Hz; see [179] and references within.

Example 4: Low firing rates optimize metabolic efficiency

Let's assume that the static power consumption of the neuron is a constant cost_{static}, whereas the dynamic power consumption scales linearly with the number of spikes $\mu = \mathbb{E}[Y]$ at a fixed cost of e_{spike} per spike. The total power consumption of the neuron is then $\mu e_{\text{spike}} + \text{cost}_{\text{static}}$. If we further assume that the firing-rate Y of the neuron is only subject to additive noise and exponentially distributed, which, as we saw already in chapter 5, maximizes the neuron's capacity for a certain mean firing rate μ , the neuron's capacity to transmit information is $1 + \log(c\mu) + c$, where c is a unit-dependent scaling factor to make $c\mu$ unit-free. Under these (mild) assumptions, the efficiency is a function of the mean firing rate

$$\epsilon(\mu) \propto \frac{1 + \log(c\mu) + c}{\mu e_{\rm spike} + {\rm cost}_{\rm static}},$$

which has a unique maximum for

$$\mu = \gamma/cW(\gamma)$$
, where $\gamma := \frac{\cos t_{\text{static}}}{e_{\text{spike}}}$ and *W* is Lambert's function.

If we take biological measurements from rodents [178] to estimate static and dynamic power ($e_{\text{spike}} \approx 7.12 \times 10^8 \text{ATP}$, $\text{cost}_{\text{static}} = 3.42 \times 10^8 \text{ATP}/\text{s}$, both measured in terms of the consumed number of adenosine triphosphate (ATP) molecules, c = 1s), we get a factor of $\gamma \approx 0.48$ and thus an optimal firing rate $\mu^* \approx 1.41$ Hz.

maintain this optimal encoding in the face of changing or unpredictable input distributions

Rate-coding spiking neural networks and machine learning

Spiking neuron models are certainly worth studying for both biologists and neuromorphic hardware designers, but since they are more difficult to simulate in software and rate-coding offers no apparent qualitative computational benefits over deep learning, they are currently of little relevance for machine learning. Hence, the viability of SNNs hinges on how well they can encode continuous signals into spikes and back, and how efficiently this can be physically implemented. Rather than representational power of the neuron, metabolic efficiency might therefore be the decisive factor behind the evolutionary success of spiking neural networks.

In the following chapter 7, we'll pursue this idea to it natural conclusion by looking at even more parsimonious, event-based alternatives to the rate-coding paradigm itself.

¹⁸ Work to apply these ideas to the design of neuromorphic hardware is currently ongoing, but not yet completed at the time of writing this thesis.

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What we have found is that at least a comparable information capacity is potentially available in respect of impulse timing [...] and it seems unlikely that the nervous system functions in such a way as to utilize none of this.

- MacKay and McCulloch in [166]

Spike-timing and event based computation

In chapter 6 we came to the sobering conclusion that in order to match the accuracy of linearnonlinear neurons, rate-coding neurons would have to fire at excessive firing rates. The high metabolic cost associated with the generation of spikes makes such "naive" rate-coding theoretically unappealing for biological and artificial neurons. We addressed this issue by directly optimizing the firing rate distribution for metabolic efficiency instead, which resulted in a much more efficient encoding with firing rates as low as 1Hz. While such low rates are in line with experimental observations, they pose a theoretical conundrum for rate-coding: We assumed that a rate-code should be linearly decodable by the dendritic filter of another neuron i.e. on a timescale on the order of tens of milliseconds — but a rate of 1Hz is orders of magnitude too slow to be smoothed by a dendritic filter! One way to make sense of this is to consider that each neuron receives input from not just a single other neuron, but from thousands, and it could be their combined input that rate-codes a signal. But this explanation raises another question: If all of these neurons encode the same signal through their firing rates, this redundancy increases the energy cost again — destroying any gains in metabolic efficiency due to the individually lower firing rates. Wouldn't a single neuron at a higher firing rate be more efficient? 1 On the other hand, if all of these neurons encode different signals, each of these signals is represented by a too low firing rate to be interpretable in a rate-coding setting, and we're back to square one.

In this chapter, I'd therefore like to ask a more fundamental question: Is rate-coding already the best we can do, or is there a more metabolically efficient code for spike-based communication?

An early study [166] applied tools from information theory to establish limits for how much information a single spike could, in principle, transmit in a realistic setting. Their estimate put this capacity at an astonishing 9 bits per spike — orders of magnitude larger than what we saw for rate-coding neurons in chapter 6! Early empirical studies have yielded more conservative estimates for the amount of information *actually* transmitted per spike *in vivo* (around 1 bit per spike, see [186] for a (dated) review), but some more recent experiments do come surprisingly close to this theoretical limit, demonstrating transmission of around 5.6-7 bits per spike [7, 187]! In order to achieve such a high information content per spike, merely counting the average rate of spikes per second is not sufficient — the timing must be taken into account, as well. In the following, we'll therefore look at *spike-timing*-based codes.

7

¹ In chapter 6 we saw that the power consumption scales almost linearly with firing rates with a per-neuron overhead due to static power consumption. Such a redundant population code would therefore almost certainly be less efficient.

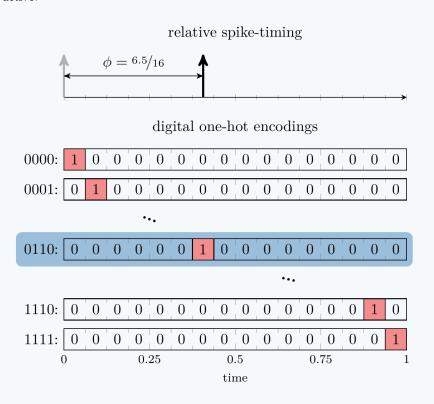
² For example, we saw an LIF neuron firing around 300 spikes per second to encode a signal with an accuracy and speed roughly comparable to a 4 bit signal sampled at 20Hz, i.e. at a rate of only about 0.27bits per spike.

7.1 Spike-time coding

Depending on the definition, there might be infinitely many different spike-trains with the same (time-varying) firing rate, so there is a lot of potentially relevant information encoded in the precise spike-times in addition to the mere firing rate (see also example 5). But how could a biological neuron extract such timing information?



To understand how spike timing allows a single spike to convey multiple bits of information, we can make a simple analogy to a digital serial code that uses a "one-hot"-encoding with 2^n bits to convey n bits of information. In the case of n = 4, this means that within a time-interval discretized into $2^4 = 16$ time-steps, exactly one bit is active:



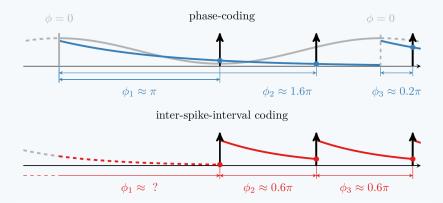
By looking at *which* bit is active within a given interval, we can thus recover 4 bits of information encoded by the corresponding sequence (here, 0110). A single *active* bit (or spike) in this example therefore transmits 4 bits of information! Note however, that this requires a reference signal to indicate the start of the interval and a precise clock signal against which the relative timing of the spike can be measured. Therefore, phase coding is just the time-continuous counterpart of *pulse-position-modulation*, a popular digital encoding scheme in signal processing!

Let's look at two popular models of spike-time coding that can be implemented by integrate-and-fire neurons (see also example 6):

The first makes use of the timing-delay between a neuron's spikes and a separate reference signal to encode a real-valued number. The reference signal could either be some particular event such as the onset of a stimulus or a saccade (then also called "time-to-first-spike" coding [188]), or it could be some change in the electrical potential of the neuron. If the reference

Example 6: Implementing phase and ISI-codes

Information can be encoded by the timing of spikes relative to some reference signal. Here, the reference signal could be either a spike from another neuron or a specific phase of a background oscillation (phase coding), or it could be the previous spike from the neuron itself (ISI coding).



In either case, the relative timing of each spike provides one real valued sample. Using a simple mechanism like dendritic filtering with a slowly decaying exponential filter that is triggered by the reference signal, the relative timing delay of a spike to the reference is a (nonlinear) function of the remaining trace at spike-time.

signal is periodic, such as theta oscillations in hippocampus [189], the spike-train thus encodes a periodically sampled signal. Each sample is then encoded by the relative phasedelay between the spike and the reference signal, which is why this code is also referred to as phase-coding. For a comparatively slow reference signal, such a code results in a very sparse spike-train with, in the extreme case, only a single spike per cycle that encodes a multi-bit measurement! Just like in the linear-nonlinear neuron model, synaptic weights can be used to change the timing of spikes, and the real-valued samples encoded by the relative spike times can be used for universal computation (see e.g. chapter 2 of [3]). Such a code also offers some computational advantages over the rate-codes from chapter 6: Consider, for example, an assembly of multiple neurons, each of which represents a different feature of an input signal and is decorrelated from its neighbors by inhibitory lateral connections, e.g. via some inhibitory inter-neuron. The neuron with the strongest response then fires first, and thus disables the others before they can fire. Such an assembly would compute the maximum operation over multiple signals with only a single spike fired! This mechanism is also extremely robust to changes in scale: If all neurons' responses were scaled down, the time of each spike might be delayed, but the order would be preserved, leading to the same result. This is consistent with the observations that cortical neurons typically fire at much lower frequencies and respond faster than the rate-coding perspective would require (see chapter 6), and that most information about a novel stimulus can often be decoded from just the first few spikes [188].

The second kind of spike-timing code is inter-spike-interval (ISI) coding [3], and it assumes that information is conveyed by the precise time-interval between two consecutive spikes. Just like with periodic sampling, a bandwidth-limited analog signal can in principle be encoded, transmitted and decoded without loss [107] using such an encoding³. A biological neuron might implement this e.g. by an exponentially decaying trace of the membrane

³ However, optimal decoding of such a signal might require a more complex mechanism than the linear decoder we required in chapter 6.

potential or some chemical, which is reset to a fixed value by each spike. The value of the trace at any point in time then (nonlinearly) encodes the time since the previous spike, and affects the likelihood of the neuron to fire once it receives a new input. One might argue that this is quite similar to what the leaky-integrate-and-fire model already does in the limit of extremely low firing rates, and the key to explain its superior performance over the LNP model (see chapter 6).

Both of these spike-timing based encoding-mechanisms integrate nicely with the theoretical framework of *spike-timing dependent synaptic plasticity* (STDP) [190], which not only consider the simultaneous firing rates of the pre- and postsynaptic neuron (like rate-based Hebbian rules), but also the relative timing of their spikes.

7.2 Event coding

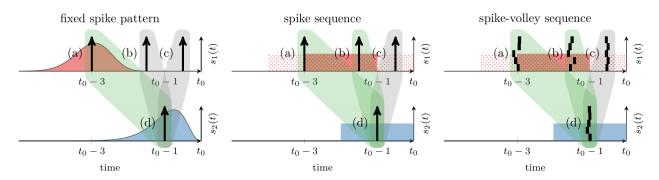
Both of the spike-timing-based approaches above rely on the same basic assumption as rate coding, that a spike-train ultimately encodes some time-varying, continuous signal or samples thereof. This provides a nice mathematical framework in which to compare various encoding schemes, and it integrates perfectly with the prevalent machine-learning perspective of neural networks as continuous function approximators (see chapter 2). But in some situations this might be an overly convoluted way of explaining a much simpler phenomenon: the neuron just fires a spike, whenever it receives a "relevant" stimulus! I'll call this simpler view the *event-coding* paradigm, in which each spike (or volley of spikes, as we shall see later) represents the occurrence of a specific *event*, rather than a real-valued sample of some continuous signal. Conceptually, this is much closer to an *interrupt-* or *event-driven* rather than a sampling-based mode of communication, which is also used in digital electronics to convey sparse signals with little latency.

A biological example of such an extremely parsimonious event-based code can be found in the fast sensory pathway of the weakly electric fish [191], where spherical neurons produce only a single individual spike in response to a prolonged stimulation. That neurons would use such an event-based style of communication also seems reasonable from an evolutionary perspective, since many of the biological mechanism used by spiking neurons predate the sort of nervous system required to even generate or interpret a rate-, ISI- or phase-coded signal. Consider for example bacteria that can form biofilm and coordinate through chemically communicated electric action potentials [192], or the rudimentary Ca²⁺ signalling mechanism already present in choanoflagellates [193] that predate animal life. Here, an event (e.g. high concentration of a chemical) triggers a specific response (e.g. release of chemicals, formation of a biofilm) — a simple mechanism that might be a precursor to spiking neurons and is best understood from this event-driven perspective. In another evolutionary stage, nerve-nets [91], action potentials often induce some synchronized behavior throughout the body of an animal. For example, pacemaker neurons of the jellyfish generate periodic action potentials that trigger a nerve-net of motor neurons to drive synchronized contraction of swimming muscles [91]. The output of these pacemaker neurons can be best understood as a form of event-coding. The spike-based transmission of information might therefore originate in some form of event-coded sensory or motor signals. By triggering these event detectors at a stimulus-dependent rate, it is conceivable that rate-based codes could also have emerged from such a simpler event-based code.

7.3 Detecting events in spike-trains

But what exactly constitutes an *event*, and how can such an event be detected? While this may be clear for a sensory neuron, we need to specify what *event* means in the context of

cortical neurons that only receive spiking input from other neurons. I will give two different definitions, fixed spatio-temporal patterns, and ordered but variable sequences of such patterns. Most of these ideas apply to continuous signals as well, but in both cases I will focus only on spike events.



Fixed spatio-temporal patterns

The most obvious definition of an *event* would be a stereotypical signal that last for a brief time-interval and always follows the same time-course. For spiking signals, that would be a fixed pattern of spikes over time, distributed across one or more neurons. We can define this as follows:

A spike pattern event $P = \{\tau_{i,j} : i \in \{1, ..., n\}, j \in J_i\}$ that occurs at time τ produces the *n*-dimensional signal $P_i(t) = \sum_{j \in J_i} \delta(t - \tau - \tau_{i,j})$, where $\tau_{i,j}$ are called the *spike-times* of the pattern. J_i is the index set of spikes belonging to neuron i in this pattern.

To detect such a pattern, a neuron could make use of a dendritic filter that implements the matched filter of the pattern, i.e. a kernel $\kappa(t) = P(T-t)$ for some T (see chapter 4). To allow for small jitter in the timing of the spikes, we can additionally smoothen the dendritic filter by convolution with some other kernel g.4 Additional spikes not belonging to the pattern at all can also affect the filter response, so an appropriately high threshold needs to be chosen to allow a reliable distinction between pattern and noise. The detection of such fixed patterns therefore reduces to dendritic filtering and thresholding, which e.g. the Gamma-neuron from chapter 4 with appropriate number of filter taps can approximate very well.

While the ability to detect such stereotypical patterns is certainly useful, this kind of event-detector suffers from two draw-backs. First, biological parameters determine the timescale of dendritic integration, which limits the length of patterns that can be detected by this mechanism and may prove to be too short for many interesting patterns. Second and more importantly, this definition of a spike pattern is extremely rigid, as it prescribes the exact time of each spike in the pattern with little room for variability. This is fine for detecting relatively short patterns, such as volleys of (nearly) synchronous spikes or rapid successions of spikes produced by a "hard-wired" cell assembly or motive. The high timing precision of some cortical neurons [196, 197] shows that such well-timed spike patterns are certainly possible to generate. But for longer lasting patterns, in particular if they are driven by external inputs that can vary in length, we'd expect some variability in the timing of the individual spikes.

Ordered (but variable) sequences of spikes and spike patterns

Instead of prescribing the actual spike times as above, we might only be interested in the order in which certain spikes arrive. For example, a spike from neuron A followed by a spike from neuron B would constitute a noteworthy event regardless of the precise timing

Figure 7.1. Left: A pattern-detector for one spike each from two incoming spike-trains. The kernel for each input filter is shown in red and blue, respectively. Only the spike pair $(a) \rightarrow (d)$ is an accepted pattern. Middle: A detector for a sequence of one spike in input 1 that precedes another spike in input 2 by some bounded timeinterval. Both patterns $(a) \rightarrow (d)$ and $(b) \rightarrow (d)$ are accepted. **Right:** The same detector for a sequence of spikevolleys, rather than individual spikes.

⁴ This corresponds to a Janossy distance metric over spike-trains [194] and could be similarly derived from optimal transport theory [195].

of either spike (as long as *B* fires within some time interval after *A*). This would be a very parsimonious code, as well, but it relies on the ability of an individual spike to reliably encode the occurrence of some event. Conversely, a single erroneously generated spike could trigger such a neuron and lead to a false detection.

To improve the reliability of such an event-based code, we can extend this concept to ordered sequences of spike patterns, e.g. sufficiently large volleys of spikes from some assembly of neurons rather than individual spikes. In the notation from above, such a spike volley corresponds to a pattern with a single spike per neuron (i.e. $J_i = \{1\}$), all of which are set to occur at roughly the same time (i.e. $t_{i,j} = 0$). This can be easily detected by a dendritic filter, e.g. the fast exponential filter of the leaky-integrate-and-fire neuron model or a brief rectangular filter would work. Since multiple synchronous spikes are required to elicit a response, such an encoding would be extremely robust to noise, while having very low latency and requiring only relatively few spikes to signal a noteworthy event.

7.4 Active dendritic sequence processing

The ability to detect sequences of spike-volleys as discussed above would be a very useful property for spiking neurons to have, but it requires more sophisticated biological mechanisms than just passive dendritic integration. Over the last century, a lot of research has gone into studying the electrical properties of cortical neurons, but only in the last two decades has the vastly improved technology in neuroimaging and electrophysiology allowed a deeper investigation of one rather fundamental property of cortical neurons: Neural dendrites are not just the passive cables we considered in chapter 4, but they can produce localized long-lasting depolarization, i.e. dendritic NMDA or calcium spikes or, as I will collectively call them, plateau potentials [198]. These actively generated effects have been shown to play an important role in cortical UP-states [199], the generation of spikes and bursts, synaptic plasticity and learning, non-linear dendritic computation, and more [198]. In an apparent case of convergent evolution, physiologically different but functionally similar mechanisms exist not just in cortical pyramidal neurons, but also in other cell types such as Purkinje cells [200].

London and Häusser [201] suggested that such localized processes would endow a single dendrite with countless *functional subunits*, which might be the key to understanding a neuron's computational capabilities. Given the importance and ubiquity of this phenomenon, it is surprising how few models in theoretical neuroscience and machine learning currently incorporate active dendritic processes or offer an explanation of their contribution. This may in part be due to inconclusive and sometimes even contradictory biological evidence⁵, which makes it difficult for theoreticians to decide, which phenomena are fundamental, and which are merely "quirks of nature". Two very interesting models by Hawkins and Ahmad [204] and Brea, Gaál, Urbanczik, and Senn [205] include such active dendritic processes to explain the emergence of a predictive UP-state in the neuron's somatic membrane potential, which allows the individual neuron to predict (and learn) "state-transitions" and a network of such neurons to (learn to) detect long-lasting sequences of input! We build on these ideas and derive a much more general model of this process, which we call *active dendritic sequences processing*.

The proposed model and its derivation from basic biological principles and observations is detailed in contribution 7. It uses passive dendritic filtering (i.e. the integration of EPSPs) in individual, electrically isolated dendritic compartments to detect volleys of coincident spikes originating from some populations of neurons. Upon detecting such an event, an *active* process generates a localized, long-lasting depolarization (a *plateau potential*), which enables a nearby dendrite segment to detect the next volley event in the sequence. If the second

⁵ For example, it has been reported that individual spike-inputs at apical dendrites might have no measurable impact on somatic membrane potentials due to strong signal attenuation [202] just as it has been reported that this effect might be completely compensated for by synaptic scaling [141]. Strongly nonlinear interactions between localized dendritic membrane potentials have been demonstrated [142], but other results show a nearly linear integration through the entire dendrite [203]. The list goes on.

Note: Stochastic population codes with event-based spiking neurons

Just like a single spike, a spike volley is a well-timed event that can be used for an event-based code. But unlike the single spike, it conveys an additional piece of information besides the timing of the event: the magnitude of the volley, i.e. the number of participating spikes. This could carry either of two different interpretations: it could signify the magnitude of the corresponding event, just like an earth-quake event has a time and magnitude, or it could signify the probability with which a binary event has occurred, encoding the detector's uncertainty. The latter interpretation is very useful in the context of detecting sequences, as it allows a population of sequence detectors to encode their uncertainty. It also offers a simple interpretation for the unreliable transmission of spikes by stochastic synapses: With deterministic synapses, a spike volley of a given magnitude will either always or never suffice to trigger a plateau potential. But with stochastic synapses, only a random subset of those spikes will be transmitted. Therefore, the probability, that the effective size of the spike volley is sufficient to trigger detection depends on the magnitude of the volley. This turns the individual ADSP neuron into a probabilistic detector, which responds to a sequence of input patterns with a probability that reflects the uncertainty encoded in the input signals. For example, a sequence where each required event has occurred with high certainty will lead to a sequence of large spike volleys, which will be detected with high probability. However, if any of the events only occurred with reduced probability, the corresponding spike volleys will be smaller, and hence the detector is more likely to not respond. By combining multiple such detectors into an assembly, we again produce a code of spike-volleys, where the magnitude of the volley is the number of triggered detectors, and thus encodes the probability of the sequence having occurred. This is highly beneficial if we want the population to quickly produce a graded response to a spike-based input [206]! Therefore, ADSP neurons with stochastic synapses can be combined into assemblies or populations that communicate via a stochastic, event-based population code (see also chapter 6 of [115])!

pattern actually occurs during this plateau, a new plateau is generated in that segment, which in turn activates another dendrite segment, and so on. This procession can make its way to the soma, where it then triggers a spike, if and only if the entire sequence of events has occurred in the correct order. Importantly, the precise timing of the individual volleys doesn't matter here, as long as they happen in the correct order (and within the specified time-intervals). This allows individual neurons to detect ordered sequences of incoming spike volleys that can last hundreds of milliseconds! Not only does this mechanism allow an individual neuron to detect sequential inputs, but it also provides a simple yet reliable mechanism to do non-linear computations with spike-volleys in continuous time. This also resolves the important question, how the fast (passive) neural membrane potential dynamics can contribute to the detection of patterns on a much slower, behaviorally relevant timescale: Our model only uses the fast dendritic filter to detect brief volleys of coincident spikes, rather than complex temporally extended patterns as in chapter 4. The further integration of that information on a slower time-scale is then due to long-lasting plateau potentials. This is a more realistic interpretation of biological evidence [207]. Since this mechanism is invariant to changes in the precise timing of the spikes (or spike volleys), it would allow the detection of such sequences across multiple time-scales, which might be relevant e.g. for reactivation, replay or preplay of hippocampal place-cell activity [208-210]. If we once

again draw a comparison to concepts from computer science and electronics, this behavior is better described by a *state machine* or *timed automaton* [211, 212], rather than the logic gate we saw in chapter 3.

Contribution 7: Event-based pattern detection in active dendrites

In this manuscript, we derive a simple yet powerful mechanisms of dendritic computation in single neurons from first biological principles. Our model makes use of actively generated dendritic plateau potentials, which provide the neuron with distributed processing elements and memory traces that collectively allow a single neuron's dendritic tree to process information in nonlinear ways and on timescales that exceed the typical timescales of membrane potentials by orders of magnitude. We show how this event-based mechanism can be used to reproduce well known nonlinear computations when viewed from a rate-coding perspective, but also how it goes much further than that by detecting specific long-lasting sequences of spike volleys and integrating information from a vast number of inputs over comparatively long time-scales. A pre-print of this paper is publicly available, and a revised version of the same manuscript is currently still under review.

Reference:

J. Leugering, P. Nieters, and G. Pipa, "Event-based pattern detection in active dendrites," *bioRxiv*, 17, 2020. DOI: 10.1101/690792v3.

To better understand how such a neuron can process information through the interaction of localized process that are distributed throughout the dendrite, an analogy can be made to decision trees, which rely on a similar hierarchical structure to classify high-dimensional inputs. A modified learning rule for decision trees can therefore even be used to train this biologically motivated neuron model! I investigated this perspective in contribution 8.

The event-based detection of long sequences results in a highly parsimonious code, which offers potentially large savings in energy consumption for biological neurons and neuromorphic hardware alike, which is why we also filed a patent for a digital neuromorphic circuit model of an ADSP neuron that can be implemented in a fully digital electronic circuit (see contribution 9).

7.5 Rate-, phase-, ISI-, or event-coding?

So which code do spiking neurons *actually* use: rate-, phase-, ISI-, or event-coding? This question goes back almost 70 years to [166], who used information theory to analyze the maximum capacity of a synapse under various assumed codes. But the answer is context dependent and differentiating between these paradigms can be surprisingly difficult.

To illustrate how much the answer to this question depends on context, consider for the sake of argument a hypothetical neuron that can detect the presence of a specific odor of a predator and fires a single spike (or a burst of spikes) whenever it detects a few of its molecules, which happens once every couple of milliseconds.

Looking at the exact same spike-train, a proponent of rate-coding could rightly argue: "The more molecules there are, the higher the firing rate of the neuron, hence the neuron uses a rate-coding approach to encode the concentration of the molecules.".

Contribution 8: Making spiking neurons more succinct with multicompartment models

In this conference paper, which accompanies a full-length presentation (which was postponed due to the ongoing SARS-Cov-2 pandemic and is now to be held in March, 2021), I analyze the computational properties of the biologically motivated multicompartment neuron model of contribution 7 from a machine-learning perspective. By transferring and adapting concepts and learning rules developed for decision trees to this neuron model, I give an intuition for how such a hierarchical structure like a neural dendrite can be useful for computation, and how simple, local learning rules might be enough to optimize such models.

Reference:

J. Leugering, "Making spiking neurons more succinct with multi-compartment models," in Proceedings of the Neuro-Inspired Computational Elements Workshop, 17, 2020, ISBN: 978-1-4503-7718-8. DOI: 10.1145/3381755.3381763.

A proponent of ISI-coding could argue with equal justification: "Since the incoming events are essentially Poisson-distributed with time-varying rate, two detections in short succession are a good indicator of a high concentration, so the neuron uses an ISI-coding approach.".

A proponent of Phase-coding could say for much the same reason: "The waiting time between phase zero of some reference oscillation and the first detection of a molecule gives an estimate of the concentration, so the neuron uses a phase code.".

From an event-coding perspective, I would argue: "The neuron merely signals each event, i.e. the detection of a molecule, with a spike.".

Neither of these explanations is wrong, but the way this thought experiment was set up, they are not equally useful: In order for some downstream neuron to make the decision whether the animal should stay or run away, the rate-coding perspective would require passing the spike train through a low-pass filter with a long enough time-constant to combine the effects of multiple spikes (see chapter 6), which introduces an inevitable and irreversible delay into the signal (see chapter 4). To reach some specific accuracy in the decoded signal, the filter must be longer and hence the response must slower the fewer spikes there are. This is obvious a problem in our thought-experiment, since the animal wouldn't have the luxury of waiting that long! The phase coding approach would require some reference signal, the rate of which limits the response time of the animal and the phase of which introduces an independent random variable. ISI coding would require multiple spikes in order to assess their relative timing. In the extreme case, where a decision must be made based on a single spike, event-coding thus seems to be the only viable explanation of the neuron's code.

But in a slightly different situation, e.g. if a decision to stay or run away is not based on the detection of individual molecules, but rather on whether the average concentration exceeds some higher threshold for some period of time, a rate-coding view might very well offer the better explanation! In fact, two different "decoders" downstream from the neuron might even simultaneously decode the same spike-train for different purposes by ways that can be explained by different paradigms. A crucial take-away of this thought experiment is therefore, that the coding paradigm we use to explain neural firing depends as much on the receiver as it does on the transmitter!

Contribution 9: "Neuromorpher Musterdetektor und neuromorphe Schaltkreisanordnung hiermit" (German patent filing)

Based on the insights derived from the neuron model of contribution 7, we designed a digital neuromorphic circuit that can efficiently realize the computation required for active dendritic sequence processing without the need for any general purpose processing elements like arithmetic-logic-units or micro-processors. It implements a processor for temporal patterns and sequences in each hierarchically structured neuron through a combination of pulses of different lengths, just like its biological counterpart. Homogeneous assemblies of multiple such neurons then communicate with each other through a code that serializes and transmits multiple spike-trains over a single binary connection.

Reference:

J. Leugering, P. Nieters, and G. Pipa, "Neuromorpher Musterdetektor und neuromorphe Schaltkreisanordnung hiermit," patent application DE 10 2019 134 044 A1

But one obvious benefit of event-based communication is that it allows for a maximally sparse code, where each event of interest is represented by just a single spike. A neural network thus becomes a network of interconnected pattern detectors, where neurons close to the periphery detect patterns in the input stimuli, while neurons deeper within the nervous system can be thought of as detecting "patterns of patterns". This argument is very appealing for both computational neuroscience and for implementations of spiking neural networks in the context of machine learning and neuromorphic hardware. However, this requires individual neurons to be able to detect relevant patterns in the first place. This requires an extension of our spiking neuron models, which incorporates well-known but often neglected active processes that occur within the dendrites. Due to the solid foundation of this mechanism on biological evidence and the powerful computation it enables, I believe event-coding to be a fundamental, if not the primary, mode of spike-based communication. Consequently, I have come to view rate- and phase-coding as modifications or refinements thereof, which become relevant when some type of event occurs often enough to admit a notion of *rate*, or when its timing is only relevant in relation to some reference signal.

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Perfection is finally attained not when there is no longer anything to add, but when there is no longer anything to take away.

- Terre des hommes, Antoine de Saint-Exupéry

I feel like I'm just gluing oranges to hair-dryers!

- Anonymous

It's the most powerful words in the world. No argument or eloquence can stand a chance against it. [...] It's: "So what?"

— Dusty Attenborough, Ginga Eiyū Densetsu

Conclusion

As we have seen in chapter 1, the study of artificial neural networks started with an attempt to capture the basic mechanism by which biological neurons process information, and to distill it into an abstract mathematical model. And still today, such artificial neural networks are used as the dominant metaphor to explain "how the brain works", i.e. how it is, that sensory information is processed, decisions are made and actions are taken. The success of deep learning has left many with the illusion that we have finally "cracked the code" of neural information processing, and in a (surprisingly unsurprising) twist, the answer appears to be the same function approximation framework that was already proposed by cyberneticists in the 1960s, and then again by Connectionists in the 1980s. Of course, the capability of recent (deep and/or recurrent) artificial neural networks to solve all sorts of machine learning problems has improved to an impressive degree, and demonstrates the enormous potential of neural networks much more effectively than either cyberneticist or theoretical neuroscientists could. However, these models, which we briefly looked at in chapters 2 and 3, are primarily designed with machine learning applications in mind and hence provide an extremely simplistic, sometimes even misleading, perspective on information processing in the brain. This does not discredit deep neural networks in the least, but it shows that despite their common origin, neuroscience and deep learning have fundamentally different objectives, and caution is required when transferring intuitions from one to the other. Goodfellow, Bengio, and Courville [39] summarized this clearly:

[O]ne should not view deep learning as an attempt to simulate the brain. [...] It is worth noting that the effort to understand how the brain works on an algorithmic level is alive and well. This endeavor is primarily known as "computational neuroscience" and is a separate field of study from deep learning. It is common for researchers to move back and forth between both fields. The field of deep learning is primarily concerned with how to build computer systems that are able to successfully solve tasks requiring intelligence, while the field of computational neuroscience is primarily concerned with building more accurate models of how the brain actually works.

But this is not to say that machine learning and theoretical neuroscience couldn't benefit from each other. Quite to the contrary, I believe that neuroscientists could benefit greatly from the analysis tools developed in machine learning, electrical engineering and computer science, whereas computer scientists and engineers interested in machine learning would do well to take more inspiration from the biological mechanisms analyzed in neuroscience!

Throughout this entire thesis, I have therefore attempted to discuss several inherently biological phenomena that defy this framework in the language of engineering, i.e. dendritic filtering in chapter 4, homeostatic plasticity mechanisms in chapter 5, spike-based

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communication in chapter 6 and finally event-based mechanisms of neural computation in chapter 7.

To decide whether these additions are *actually* instrumental for information processing or merely abstract descriptions of a needlessly complicated biological mechanism, we will have to put them to the test. I have come to believe that the best way to do that is by looking across the domain boundaries between neuroscience and adjacent disciplines, in order to find inspiration and to test ideas in a less forgiving environment outside one's own control. Particularly the *embodiment* of concepts from theoretical neuroscience in neuromorphic hardware appeals to me as a tough, but honest benchmark that makes it possible to evaluate the merit of many theoretical models of neural computation "in the real world". I therefore believe that many future innovations in theoretical neuroscience will originate in or be driven by such application oriented fields, which also motivated my own transition towards neuromorphic hardware.

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A Appendix for chapter 4

Some derivations in this appendix can also be found in similar form in the original work on the Gamma neuron [125, 126], others come from (unpublished) work in our lab. I have decided to re-derive and compile them here for conciseness and completeness. At some points, we have also slightly extended the Gamma neuron beyond its original definition or taken it out of its original context, but decided to keep the established name for our more general extension.

Equivalence between filtering and continuous delays

Filter and (continuous) delay operators are very closely related.

To show the first direction of this relationship, consider a continuous signal s(t). Any operator D that delays s by some fixed delay $\Delta t \ge 0$ can be expressed as a causal filter with a shifted Dirac-distribution as its kernel:

$$Ds(t) = s(t - \Delta t) = \int_{-\infty}^{t} s(\tau)\kappa(t - \tau)d\tau$$
$$\kappa(t) := \delta(t - \Delta t)$$

Conversely, for a continuous signal s(t) and a causal filter with (piece-wise) continuous kernel $\kappa(t)$, we can use the Riemann integral to approximate the effect of the kernel κ by a linear combination $\bar{\kappa}$ of shifted and scaled δ -pulses:

$$(s * \kappa)(t) = \int_{-\infty}^{t} s(\tau)\kappa(t - \tau)d\tau$$

$$= \lim_{\Delta t \to 0} \Delta t \sum_{k=0}^{\infty} s(t - k \cdot \Delta t)\kappa(k \cdot \Delta t)$$

$$= \lim_{\Delta t \to 0} \Delta t \sum_{k=0}^{\infty} \left(\int_{-\infty}^{t} s(\theta)\delta(t - \theta - k \cdot \Delta t)d\theta \right)\kappa(k \cdot \Delta t)$$

$$= \int_{-\infty}^{t} s(\theta) \lim_{\Delta t \to 0} \Delta t \sum_{k=0}^{\infty} \kappa(k \cdot \Delta t)\delta(t - \theta - k \cdot \Delta t)d\theta$$

$$= (s * \bar{\kappa})(t) \quad \text{where } \bar{\kappa}(t) = \lim_{\Delta t \to 0} \Delta t \sum_{k=0}^{\infty} \kappa(k \cdot \Delta t)\delta(t - k \cdot \Delta t)$$

A.2 Transfer function of the Gamma neuron

Consider the Gamma neuron as described in section 4.4. We can analyze the linear system realized by its dendritic filter in the Laplace domain. Let us denote the Laplace transform of the j^{th} input signal s(t) with $S_j(s)$, and the Laplace transform of the output signal of the i^{th} tap of the dendritic filter with $X_i(s)$, where $X_1(s)$ is the output of the filter tap closes to the soma. To simplify the analysis, we separate the feed-forward and the feed-back paths of the model by defining the *open-loop impulse-response* to the j^{th} external input signal (denoted κ_j^{fwd}) as well as the open-loop impulse-response of the neuron to its *own output* y (denoted κ_j^{fwd}).

In the absence of feedback, an impulse arriving at the i^{th} tap from the soma has to traverse i filters on its way to the soma, and hence produces (in the Laplace domain) a filter response $\left(\frac{\alpha}{s+\alpha}\right)^i$ — which corresponds to the probability density function of a Gamma distribution with coefficients i and α .

For a neuron with n taps and m input signals and time-constant α , these individual impulse responses of the taps are scaled by the forward and feedback weights $w_{i,j}$ and v_i , respectively, and linearly combined to yield the effective dendritic filter kernels:

$$\kappa_j^{\text{fwd}} = \sum_{i=1}^n \left(\frac{\alpha}{s+\alpha}\right)^i w_{i,j}$$

$$\kappa^{\text{fb}} = \sum_{i=1}^n \left(\frac{\alpha}{s+\alpha}\right)^i v_i$$

We can then derive the transfer function:

$$\begin{split} X_{1}(s) &= \kappa^{\text{fb}} X_{1}(s) + \sum_{j=1}^{m} \kappa_{j}^{\text{fwd}} S_{j}(s) \\ \Rightarrow X_{1}(s) &= \sum_{j=1}^{m} \frac{\kappa_{j}^{\text{fwd}}}{1 - \kappa^{\text{fb}}} S_{j}(s) \\ &= \sum_{j=1}^{m} \frac{\sum_{i=1}^{n} \left(\frac{\alpha}{s + \alpha}\right)^{i} w_{i,j}}{1 - \sum_{i=1}^{n} \left(\frac{\alpha}{s + \alpha}\right)^{i} v_{i}} S_{j}(s) \\ &= -\sum_{j=1}^{m} \frac{\sum_{i=1}^{n} \alpha^{i} (s + \alpha)^{n-i} w_{i,j}}{\sum_{i=0}^{n} \alpha^{i} (s + \alpha)^{n-i} v_{i}} S_{j}(s) \\ &= -\sum_{j=1}^{m} \frac{\sum_{i=1}^{n} \sum_{k=0}^{n-i} \binom{n-i}{k} s^{k} \alpha^{n-k} w_{i,j}}{\sum_{i=0}^{n} \sum_{k=0}^{n-i} \binom{n-i}{k} s^{k} \alpha^{n-k} v_{i}} S_{j}(s) \end{split}$$

The last expression can be simplified using matrix-vector multiplications:

$$(w^{j})_{i} := w_{i,j}, \quad w^{j} \in \mathbb{R}^{n}$$

$$(M)_{k,i} := \begin{cases} \alpha^{n-k} \binom{n-i}{k} & \text{if } k \leq i \\ 0 & \text{otherwise} \end{cases}, \quad M \in \mathbb{R}^{n \times n}$$

The matrix M above is an invertible matrix, therefore the coefficient vectors Mw^j and Mv of this rational transfer function can be freely determined by an appropriate choice of the feed-forward and feedback weights $w_{k,j}$ and v_k , respectively. The resulting transfer function of the dendritic filter simplifies to:

$$X_1(s) = \sum_{i=1}^m \frac{\sum_{k=1}^n s^{k-1} (Mw^i)_k}{1 - \sum_{k=1}^n s^k (Mv)_k} S_j(s).$$

The special case of a Gamma neuron without feedback occurs when setting the feedback weights v to zero, which results in the simpler form:

$$X_1(s) = \sum_{j=1}^m \left(\sum_{k=1}^n s^{k-1} (Mw^j)_k \right) S_j(s)$$

The Gamma neuron with n filter taps and linear feedback can thus be used to implement a dendritic filter with arbitrary *proper rational* transfer function (for a single input signal) with degrees up to n-1 in the numerator and n in the denominator. For multiple inputs, the numerator of this transfer function can be individually chosen for each input, whereas the denominator (determined by the feedback coefficients) is shared among all inputs.

The ability to freely place zeros and poles (i.e. zeros of the denominator) of the transfer function makes this type of filter bank with feedback extremely versatile. It is capable of implementing a wide range of practically relevant filters, such as higher-order Butterworth, Chebyshev and Elliptic filters in low- and band-pass form. High-pass and band-stop filter can similarly be implemented, but some limitations apply. For an in-depth look at continuous filter design, see e.g. chapter 7 of [124].

Derivatives of Gamma filters are Gamma filters

The time-derivative of the open-loop impulse-response of tap $k \ge 2$ in the Gamma neuron's filter can be expressed simply in terms of just two neighboring taps (see also [125]):

$$\kappa_k(t) = \frac{\alpha^k}{\Gamma(k)} t^{k-1} \exp(-\alpha t)$$

$$\kappa_k'(t) = \frac{\alpha^k}{\Gamma(k)} \left(t^{k-2} \exp(-\alpha t) - \alpha t^{k-1} \exp(-\alpha t) \right)$$

$$= \frac{\alpha}{k-1} \kappa_{k-1}(t) - \alpha \kappa_k(t)$$

Therefore, the time derivative of *any* filter constructed without feedback by linear combination of the taps $k \geq 2$ can be implemented by the same filter bank, as well. The same argument applies to the feedback path, and hence the time-derivative can be implemented for (most) filters constructed with feedback, as well.

In short, this means that the Gamma neuron model can combine filtering and differentiation in the linear operator implemented by its dendrite. The Gamma neuron therefore posses all the capabilities required of a *PID-controller*: proportional input (i.e. a δ -impulse response, e.g. approximated by a quickly decaying exponential filter), integration of input

(e.g. approximated by a slowly decaying exponential filter) and differentiation of input. See also [129] for more information about the design of PID-controllers.

A.3 The ring of Gamma filters

As shown in appendix A.2, the Gamma neuron's dendritic filter has a proper rational transfer function with order $\leq n$ in the denominator. While rational functions form a *field*, *proper* rational functions only form a *ring*, since the multiplicative inverse of a proper rational function would not necessarily be proper. This has an analog physical interpretation: The sum of two causal filters or the concatenation of two causal filters (with rational transfer functions) is again a causal filter (with rational transfer function), but a causal filter cannot be inverted by application of a causal filter (consider as a simple counter-example, that inverting a delay would require an acausal advance of the signal). This ring is commutative, but it doesn't contain a multiplicative identity (this would correspond to a Dirac- δ in the time-domain), and could therefore be called a *pseudo-ring*.

While multiplicative inverses don't exist in this ring, we can approximate a solution of $\kappa_2 = \kappa^{\dagger} \kappa_1$ for κ^{\dagger} by Euclidean division with remainder r.

$$\exists! \kappa^{\dagger}, r : \kappa_2 = \kappa^{\dagger} \kappa_1 + r.$$

Now consider a signal *s* and its filtered version $\tilde{s} := \kappa_1 * s$. We then have

$$\kappa^{\dagger} * \tilde{s} = (\kappa_2 - r) * s$$

Therefore, while we cannot implement *de*convolution, we can use filtering to "replace" the effect of the filter κ_1 with the effect of the filter $\kappa_2 - r$, an approximation of κ_2 . If $\kappa_2(s)$ is now chosen e.g. to approximate the delay $\delta(t-T)$ for some large T, then this *re*convolution with the filter κ^{\dagger} approximates a delayed *de*convolution, i.e. it approximately recovers the delayed original signal s(t-T)!

Appendix for chapter 6

Note:

Many of the derivations in this appendix are not new results, and can hence be found across standard literature like [105, 174, 213]. I have nevertheless decided to re-derive them here from scratch and compile them in order to provide a more concise summary and to allow for a direct comparison between the different models using a common language.

B.1 Rate-coding with (L)IF neurons

Activation function of integrate-and-fire neurons

The pure integrate-and-fire neuron integrates its input up until it hits a threshold θ , at which point it resets and the process begins anew. For an incoming signal s(t) with integral $S(t) = \int_0^t s(t)$, the integrate-and-fire neuron thus produces spikes at the times $t_k = S^{-1}(k\theta)$, $k \in \mathbb{N}$. If we filter the resulting spike train $\chi(t) = \sum_{t_k} \delta(t - t_k)$ by a filter $\kappa(t)$, we get the decoded signal z_{IF} :

$$z_{\rm IF}(t) = \theta \cdot (\kappa * \gamma)(t) \tag{B.1}$$

$$= \theta \sum_{t_k \le t} \int_{-\infty}^{t} \kappa(t - \tau) \delta(\tau - t_k) d\tau$$
 (B.2)

$$=\theta \sum_{t_k < t} \kappa(t - t_k) \tag{B.3}$$

For a constant signal $s(t) = c \ge 0$, we get S(t) = ct and hence $t_k = k\theta/c$. To simplify notation, we can introduce $K_t := \max\{k : t_k \le t\} = \lfloor ct/\theta \rfloor$, which is the index of the last spike before time t. If we choose the exponential kernel $\kappa(t) = H(t)\alpha \exp(-t\alpha)$, where H is the Heaviside

step-function and $\alpha > 0$ sets the time-scale of the filter, we can compute $z_{\rm IF}$:

$$z_{\text{IF}}(t) = \sum_{k=-\infty}^{K_t} \alpha \theta \exp(-\alpha (t - k\theta/c))$$

$$= \frac{\alpha \theta}{1 - \exp(-\alpha \theta/c)} \exp(\alpha (K_t \theta/c - t))$$
(B.4)
(B.5)

$$= \frac{\alpha \theta}{1 - \exp(-\alpha \theta/c)} \exp(\alpha (K_t \theta/c - t))$$
(B.5)

$$= \frac{\alpha \theta}{1 - \exp(-\alpha \theta/c)} \exp(-\alpha \Delta t) \quad \text{where } \Delta t := t - t_{K_t}$$
 (B.6)

This expression depends only on the relative time Δt since the previous spike, and repeats after every spike. Therefore, the "decoded" spike-train is a periodic signal of discontinuous jumps after every spike with period $t_{k+1} - t_k = \theta/c$, followed by exponential decay. We can thus compute the mean signal by averaging it between two successive spikes:

$$\bar{z}_{\rm IF}(c) = \frac{1}{\theta/c} \int_0^{\theta/c} z_{\rm IF}(t) \mathrm{d}\Delta t \tag{B.7}$$

$$= \frac{\alpha \theta}{\theta/c} \cdot \frac{1}{1 - \exp(-\alpha \theta/c)} \int_0^{\theta/c} \exp(-\alpha(\Delta t)) d\Delta t$$
 (B.8)

$$= c \cdot \frac{1 - \exp(-\alpha\theta/c)}{1 - \exp(-\alpha\theta/c)} \tag{B.9}$$

$$=c (B.10)$$

Therefore, the (linearly decoded) output of the integrate-and-fire neuron is given by a rectified linear function $\bar{z}_{IF}(c) = \max(0, c)$ of the constant input c.

Decoding error of integrate-and-fire spike-trains

Given $z_{\rm IF}(t)$ and $\bar{z}_{\rm IF}(c)$, we can similarly calculate the expected root-mean-squared error (RMSE) of the IF neuron.

$$MSE_{IF}(c) = \frac{1}{\theta/c} \int_0^{\theta/c} (z_{IF}(\Delta t) - \bar{z}_{IF}(c))^2 d\Delta t$$
 (B.11)

$$= \frac{c}{\theta} \int_0^{\theta/c} z_{\text{IF}} (\Delta t)^2 d\Delta t - (\bar{z}_{\text{IF}}(c))^2$$
(B.12)

$$= \frac{c\alpha^2\theta}{(1 - \exp(-\frac{\alpha\theta}{c}))^2} \int_0^{\theta/c} \exp(-2\alpha\Delta t) d\Delta t - c^2$$
 (B.13)

$$= \frac{c\alpha\theta(1 + \exp(-\frac{\alpha\theta}{c}))}{2(1 - \exp(-\frac{\alpha\theta}{c}))} - c^2$$
(B.14)

$$= \frac{c\alpha\theta}{2} \coth(\frac{\alpha\theta}{2c}) - c^2 \tag{B.15}$$

$$RMSE_{IF}(c) = \sqrt{\frac{c\alpha\theta}{2}\coth(\frac{\alpha\theta}{2c}) - c^2}$$
(B.16)

$$\lim_{c \to \infty} \text{MSE}_{\text{IF}}(c) = \frac{\alpha^2 \theta^2}{12} \quad \Rightarrow \quad \lim_{c \to \infty} \text{RMSE}_{\text{IF}}(c) \approx \frac{\alpha \theta}{\sqrt{12}}$$
 (B.17)

The last equation shows, that in the limit of relatively high firing rates (which we typically assume when talking about rate coding) the RMSE only depends on the product $\alpha\theta$ and becomes independent of c. To reduce the error, we have to either use a slower kernel with smaller α or lower the threshold θ .

A different way to interpret this result is to consider that in the steady-state, the filtered spike-train makes a jump of fixed magnitude $\alpha\theta$ after each spike and then relaxes back to the same initial value z_0 before spiking again. The higher the firing rate is, the more this exponential relaxation looks like a linear decrease, and the more the filtered spike-train looks like a saw-tooth wave. If we were to estimate the stationary distribution of the error that results from sampling the filtered signal $z_{\rm IF}$ at a random point in time, the residual r would then be uniformly distributed in the range $[c - \alpha\theta/2, c + \alpha\theta/2]$ with mean value c and standard deviation $\alpha\theta/\sqrt{12}$. We will make use of this probabilistic perspective for comparing the information content of various encodings in appendix B.3.

Activation function of leaky integrate-and-fire neurons

Leaky integrate-and-fire (LIF) neurons are very similar to integrate-and-fire neurons, but the integrator is replaced by a first-order low-pass filter with leak-rate α . We will choose the same α for both the LIF neuron and the decoder (which would be another LIF neuron in a spiking neural network, anyway). LIF neurons also fire periodically in response to constant inputs, albeit with a lower firing rate that depends non-linearly on the input and the leak-rate α . We can therefore model the LIF neuron as an IF neuron, whose input signal c' = v(c) has been nonlinearly transformed. This nonlinear distortion ν can be characterized as follows:

For constant input c and following a reset at time 0, the LIF neuron's membrane potential follows the trajectory $c/\alpha(1-\exp(-\alpha t))$, i.e. it exponentially approaches c/α rather than growing at constant rate c like in the IF neuron. The threshold θ is reached at time t= $-1/\alpha \log(1-\alpha\theta/c)$. An IF neuron would produce the same periodic firing for a different constant input c':

$$-1/\alpha \log(1 - \alpha \theta/c) = \theta/c'$$
(B.18)

$$\Leftrightarrow \nu(c) := c' = -\frac{\alpha\theta}{\log(1 - \alpha\theta/c)}$$
 (B.19)

For $c \gg \theta$, this has the asymptote $c'(c) \approx c + \alpha \theta/2$.

This implies, that the LIF neuron has the transfer function is

$$\bar{z}_{\text{LIF}}(c) = \bar{z}_{\text{IF}}(\nu(c)) = \max(0, -\frac{\alpha\theta}{\log(1 - \alpha\theta/c)}).$$

If we use the asymptotic approximation, this reduces further to simply

$$\bar{z}_{\text{LIF}}(c) \approx \max(0, c + \alpha \theta/2).$$

Decoding error of leaky integrate-and-fire spike-trains

Using the same trick of substituting in c' = v(c) for the input of an IF neuron, we can also calculate the RMSE of the LIF neuron. But since we are mostly interested in the high firing-rate regime, where we saw that the RMSE of the IF neuron does not depend on c at all, we get the same approximation for the LIF neuron as well:

$$RMSE_{LIF}(c) \approx \frac{\alpha \theta}{\sqrt{12}}.$$

Rate-coding with linear-nonlinear-Poisson neurons

To allow for a direct comparison between the LNP neuron and the (L)IF neuron from appendix B.1, let's now imagine that we filter the spiking output of an LNP neuron in response to a (piece-wise) constant input. The linear-nonlinear-Poisson spiking neuron produces stochastic spikes at the (time-varying) rate s(t) by an inhomogeneous Poisson process, i.e. the number N of spikes in the time-interval [0,t] is a Poisson random variable with expected value $\mathbb{E}[N] = \int_0^t \lambda s(\tau) d\tau = \lambda(S(t) - S(0))$. Here, λ is a firing-rate gain parameter that plays the same role as $1/\theta$ does in the LIF neuron. When decoding the spike-train, we therefore weigh each spike by $1/\lambda$ instead of θ to compensate for the gain. In contrast to the (L)IF neuron, the spike-times are *not* periodically spaced for the LNP neuron, so we have to follow a slightly different approach to derive the mean and RMSE of the decoded spike-train.

For each of these N spikes, the spike times $t_k \sim \text{Uniform}(0,t)$ are independent and identically distributed random variables. If we use the same constant signal s(t) = c and exponential filter κ as in appendix B.1, we can compute the effect γ_k of the individual spikes on the decoded signal, the combined effect Γ of all spikes since time t = 0, as well as a couple of expectations that will be useful later:

$$\gamma_k(t) = \frac{\alpha}{\lambda} \exp(-\alpha(t - t_k))$$
 (B.20)

$$\Gamma(t) = \sum_{0 \le t_k \le t} \gamma_k(t) \tag{B.21}$$

$$\mathbb{E}[N] = \lambda \int_0^t c d\tau = \lambda t c \tag{B.22}$$

$$\mathbb{E}[\gamma_k(t)] = \frac{1}{t} \int_0^t \frac{\alpha}{\lambda} \exp(-\alpha(t-\tau)) d\tau = \frac{1}{\lambda t} (1 - \exp(-\alpha t))$$
 (B.23)

$$\mathbb{E}[\gamma_k(t)^2] = \frac{\alpha^2}{t\lambda^2} \int_0^t \exp(-\alpha(t-\tau))^2 d\tau = \frac{\alpha}{2t\lambda^2} (1 - \exp(-2\alpha t))$$
 (B.24)

$$\mathbb{E}[\Gamma(t)|N] = N\mathbb{E}[\gamma_i(t)] = \frac{N}{\lambda t}(1 - \exp(-\alpha t))$$
(B.25)

$$\mathbb{E}[\Gamma(t)] = \mathbb{E}[N]\mathbb{E}[\gamma_i(t)] = c(1 - \exp(-\alpha t))$$
(B.26)

With these results, we can finally derive the decoded signal and its expected value:

$$z_{\text{LNP}}(t) = 1/\lambda(\kappa * \chi)(t) \tag{B.27}$$

$$=\sum_{-\infty < t_k < t} \gamma_k(t) \tag{B.28}$$

$$= z_{\text{LNP}}(0) \exp(-\alpha t) + \Gamma(t)$$
(B.29)

$$\mathbb{E}[z_{\text{LNP}}(t)|N] = \mathbb{E}[z_{\text{LNP}}(0)] \exp(-\alpha t) + \mathbb{E}[\Gamma(t)|N]$$
(B.30)

$$\mathbb{E}[z_{\text{LNP}}(t)] = \mathbb{E}[z_{\text{LNP}}(0)] \exp(-\alpha t) + \mathbb{E}[\Gamma(t)]$$
(B.31)

$$= \mathbb{E}[z_{\text{LNP}}(0)] \exp(-\alpha t) + c(1 - \exp(-\alpha t)) \tag{B.32}$$

We can solve the last equation easily by using the fact that the filtered signal (and hence its expectations) must be time-shift-invariant, i.e. $\mathbb{E}[z_{\text{LNP}}(t)] = \mathbb{E}[z_{\text{LNP}}(0)]$. For the mean value $\bar{z}_{\text{LNP}}(c)$ of the decoded signal in response to constant input s(t) = c, we thus get:

$$\bar{z}_{\text{LNP}}(c) = \mathbb{E}[z_{\text{LNP}}(t)] = c \tag{B.33}$$

Filtering an LNP neuron's spike-train thus also produces an unbiased estimate of its (constant) input.

Calculating the RMSE for the LNP neuron is more difficult, because it involves a random number of spikes *N* as well as the random times of each individual spike. But we can use the *law of total variance* and apply the same trick as above, i.e. enforcing that the MSE must be

time-shift-invariant:

$$\operatorname{Var}[z_{\operatorname{LNP}}(t)|N] = \operatorname{Var}[z_{\operatorname{LNP}}(0) \exp(-\alpha t) + \Gamma(t)] \tag{B.34}$$

$$= \operatorname{Var}[z_{\operatorname{LNP}}(0) \exp(-\alpha t)] + \operatorname{Var}[\Gamma(t)] \tag{B.35}$$

$$= \operatorname{Var}[z_{\operatorname{LNP}}(0)] \exp(-2\alpha t) + N\operatorname{Var}[\gamma_i(t)] \tag{B.36}$$

$$\operatorname{MSE}_{\operatorname{LNP}}(c) = \operatorname{Var}[z_{\operatorname{LNP}}(t)] \tag{B.37}$$

$$= \operatorname{\mathbb{E}}(\operatorname{Var}[z_{\operatorname{LNP}}(t)|N]) + \operatorname{Var}(\operatorname{\mathbb{E}}[z_{\operatorname{LNP}}(t)|N]) \tag{B.38}$$

$$= \operatorname{\mathbb{E}}(\operatorname{Var}[z_{\operatorname{LNP}}(0)] \exp(-2\alpha t) + N\operatorname{Var}[\gamma_i(t)]) + \operatorname{Var}(\operatorname{\mathbb{E}}[z_{\operatorname{LNP}}(0)] \exp(-\alpha t) + N\operatorname{\mathbb{E}}[\gamma_i(t)])$$

$$(B.39)$$

$$= \operatorname{Var}[z_{\operatorname{LNP}}(0)] \exp(-2\alpha t) + \operatorname{\mathbb{E}}[N]\operatorname{\mathbb{E}}[\gamma_i(t)^2] + (\operatorname{Var}(N) - \operatorname{\mathbb{E}}[N])\operatorname{\mathbb{E}}[\gamma_i(t)]^2$$

$$(B.40)$$

$$= \operatorname{Var}[z_{\operatorname{LNP}}(0)] \exp(-2\alpha t) + \operatorname{\mathbb{E}}[N]\operatorname{\mathbb{E}}[\gamma_i(t)^2] \tag{B.41}$$

$$\operatorname{Var}[z_{\operatorname{LNP}}(0)] = \operatorname{Var}[z_{\operatorname{LNP}}(t)] \Leftrightarrow \operatorname{MSE}_{\operatorname{LNP}}(c) = \frac{\alpha c}{2\lambda} \tag{B.42}$$

$$\operatorname{RMSE}_{\operatorname{LNP}}(c) = \sqrt{\frac{\alpha c}{2\lambda}} \tag{B.43}$$

Since the filtered spike-train of the LNP neuron can be viewed as a sum of independent and identically distributed random variables, we can assume that the distribution of the membrane potential (for a sufficiently high firing rate) approaches a normal distribution due to the central limit theorem. In contrast to the (L)IF neuron, the RMSE hence increases proportionally to \sqrt{c} ! We will make use of this in appendix B.3.

B.3 The entropy of LIF and LNP encoding

Let's assume as in appendices B.1 and B.2 that $c \gg 0$, so that we may assume a Gaussian distribution of errors when decoding the LNP neuron's output and a uniform distribution for the LIF neuron. The entropy of a Gaussian distribution with standard-deviation $RMSE_{LNP}(c) = \sqrt{\frac{\alpha c}{2\lambda}}$ is $E_{Gauss}^{\sigma} = 1/2 \log(\pi e \frac{\alpha c}{\lambda})$ and the entropy of a uniformly distributed random variable on the interval $[0, \alpha\theta]$ is $E_{\text{Uniform}}^{\theta} = \log(\alpha\theta)$. Therefore, measurements of the two neurons' firing rates are similarly informative if $E_{\text{Gauss}}^{\sigma} = E_{\text{Uniform}}^{\theta}$, i.e. if $\theta = \sqrt{\frac{\pi e c}{\alpha \lambda}}$ Since for a constant signal, the mean firing rate r_{LNP} of the LNP neuron scales linearly with λ whereas the mean firing rate $r_{\rm LIF}$ of the LIF neuron scales linearly with $1/\theta$, we can see that in order to reach similar performance, we have to have $r_{\rm LIF} \propto \sqrt{\frac{\alpha r_{\rm LNP}}{\pi e c}}$.

Spike-coding under metabolic constraints

Maximizing information-rate under metabolic constraints B.4.1

Following the procedure already outlined in section 5.2, we can find the optimal distribution of firing rates Y that maximizes the information-rate under certain metabolic constraints. For example, we might want to keep the expected (RMS) error below some limit θ_{error} , and the energy cost below the limit θ_{cost} . The optimal firing rate distribution must then be of the form:

$$p(y) = \exp(\lambda_0 \mathbb{1}_R(y) + \lambda_1 \operatorname{cost}(y) + \lambda_2 \operatorname{error}(y)),$$

and the optimal coefficients of the distribution $\lambda^* = \begin{pmatrix} \lambda_0^* & \lambda_1^* & \lambda_2^* \end{pmatrix}^T$ can be found by optimization of:

$$\lambda^* = \arg\max_{\lambda} \left(\lambda^T \theta - \int_R p(y) dy \right)$$
 subject to $\lambda_1, \lambda_2 \ge 0$,

where $R = [0, R_{\text{max}}]$ is the range of admissible firing rates and $\theta = \begin{pmatrix} 1 & -\theta_{\text{cost}} & -\theta_{\text{error}} \end{pmatrix}$ holds the constraints we wish to enforce. ¹ In general, it might be difficult to derive a closed-form solution for this, but if we are for instance willing to assume that (a) the firing rate is bounded, (b) the cost increases linearly with the rate and (c) RMSE is (almost) independent of y (as is the case for the (L)IF neuron), then the equation above reduces to a truncated exponential distribution! ²

B.4.2 Maximizing metabolic efficiency

While optimizing the through-put of a neuron seems reasonable from an information bottle-neck perspective, there is convincing evidence that (some) biological neurons appear to be optimized for metabolic efficiency instead, firing at rates as low as two spikes per second and with an information content as high as 5.6 bits per spike (see e.g. chapter 4 of [7]). This may be much less than the maximum bit-rate that a single neuron could deliver in principle, but it appears to make optimal use of the invested energy. To find the optimal firing rate distribution P_Y^* , we can proceed as follows: If we again use $\cos(y)$ to denote the power required for maintaining a firing rate y, the cost associated with the firing rate distribution P_Y is $\cos t_{P_Y} = \mathbb{E}_{P_Y}[\cos(y)]$. We can then express the metabolic efficiency of the neuron $\varepsilon = \frac{I_{P_Y}}{\cot t_{P_Y}}$ where I_{P_Y} denotes the information-rate of the neuron for firing rate distribution P_Y . If the cost was independent of the firing rate distribution (i.e. if generating spikes required no additional energy), then most the "powerful" encoding from above would also be the most metabolically efficient. But if we make the more realistic assumption, that each spike costs a finite amount of energy $e_{\rm spike}$ in addition to the $t_{\rm static}$ power $t_{\rm static}$ required to keep the neuron operational, the metabolic efficiency is optimized by the distribution

$$P_Y^* = \operatorname*{arg\,max}_{P_Y} \frac{I_{P_Y}}{\mathbb{E}[Y] \cdot e_{\mathrm{spike}} + \mathrm{cost}_{\mathrm{static}}}.$$

If we only consider exponential distributions and assume a one-to-one deterministic encoding by the neuron, then the information rate (which is then proportional to the entropy of Y) is just a function of the expected firing rate $I_{P_Y} \propto \log_2(\mathbb{E}[Y]) + 1$, and hence P_Y^* becomes a function of just the mean firing rate, as well. In this case, we see that the metabolic efficiency would be maximized for

$$\mathbb{E}[Y] = \frac{\gamma}{\gamma W(2 \exp(-1))}$$
, where $\gamma := \frac{\cos t_{\text{static}}}{e_{\text{spike}}}$ and W is Lambert's function.

Note that this gives us a finite, optimal firing rate for the neuron, even if we impose no hard limits on either the energy constraint or the maximum firing rate.

 1 The first term 1 and corresponding coefficient λ_0 originate from the constraint that the distribution of *Y* has to be normalized over *R*, the other two from the metabolic constraints.

² If the cost constraint is redundant, this simplifies further to a uniform distribution.

$\mathbb{C}_{\mathbb{R}}$

Index of included contributions

In the following, I have compiled all the relevant own publications referenced in the thesis. Full-text sources are available through the respective publishers. Unpublished or inaccessible contributions are attached in appendix D.

Description of the contribution

†1 **Title:** Bistable Perception in Conceptor Networks.

Publication type: conference paper

Summary: This conference paper explores, how a hierarchy of conceptor networks, which can act as generative models for time-series signals, can be used to actively suppress noise and minimize prediction errors. The idea of such a hierarchical predictive coding scheme is in line with biological observations and provides an appealing model of perception. When presented with ambiguous superposition of two stimuli, this architecture reproduces the well known psychological phenomenon of bi-stable perception, where either of the two pure stimuli is perceived in isolation for a period of time, before the percept switches to the other. It matches empirical results with surprising fidelity, including the distribution of the time-spans for which either of the stimuli is perceived! This paper extends ideas developed within Felix Meyer zu Driehausen's thesis, which I had the pleasure of supervising. Felix and Rüdiger Busche subsequently turned it into a viable model and a nice conference paper, for which they deserve all the credit.

Full-text available at: https://doi.org/10.1007/978-3-030-30493-5_3 Bibliographic reference:

F. Meyer zu Driehausen, R. Busche, **J. Leugering**, and G. Pipa, "Bistable Perception in Conceptor Networks," in *Artificial Neural Networks and Machine Learning – ICANN 2019: Workshop and Special Sessions*, 2019, ISBN: 978-3-030-30493-5. DOI: 10.1007/978-3-030-30493-5_3.

#2 **Title:** Computational Elements of Circuits.

Publication type: book chapter

Summary: The book "The Neocortex", published by the Ernst Strüngmann Forum, compiles the current state of knowledge about the basic principles of operation of the neocortex. In our contribution to this work, the book chapter entitled "Computational Elements of Circuits", we discuss several fundamental properties of neural computation — from homeostasis to delayed interactions, synchronization, random feature expansion and reservoir computing. Within this book chapter, my own largest contribution can be found in the section "Information Processing in Single Neurons and Populations", which elaborates and generalizes ideas from contribution 6.

Full-text available at: https://mitpress.mit.edu/books/neocortex Bibliographic reference:

J. Leugering, P. Nieters, and G. Pipa, "Computational Elements of Circuits," in *The Neocortex*, W. Singer, T. J. Sejnowski, and P. Rakic, eds., red. by J. Lupp, vol. 27, The MIT Press, 2019, pp. 195–209, ISBN: 978-0-262-04324-3. DOI: 10.7551/mitpress/12593.003.0016.

#3 **Title:** A visit to the neuromorphic zoo.

Publication type: conference/fair paper

Summary: In this paper, which accompanies a public talk held at the Embedded World conference, I provide a brief overview over current concepts and academic as well as commercial developments in the field of AI-hardware acceleration in general, and neuromorphic hardware in particular. This paper was selected by WEKA Fachmedien for a re-publication in the magazine DESIGN&ELEKTRONIK, where it appeared in German translation under the title "Neuromorphe Hardware". Since the Embedded World conference accompanies an industry fair, the proceedings are targeted towards an engineering audience and are intended to offer an accessible high-level perspective. (This paper passed an editorial process, but no scientific peer-review.)

Full-text available at: appendix D, page 106ff

Bibliographic reference:

J. Leugering, "A visit to the neuromorphic zoo," in *Embedded World Conference* 2020 – *Proceedings*, 2020, ISBN: 978-3-645-50186-6.

A German translation also appeared in:

J. Leugering, "Neuromorphe Hardware," *DESIGN&ELEKTRONIK*, no. 7/2020, pp. 41–47, 2020.

#4 **Title:** Neuromorphic Adaptive Filters for event detection, trained with a gradient free online learning rule.

Publication type: conference poster

Summary: We extended the Gamma Neuron to a spike-based temporal pattern detector for a conference poster presented first at the Cognitive Computing 2018 conference in Hannover, Germany, and then again at the Machine Learning Summer-School (MLSS) 2019 held in Cape Town, South Africa. Here we investigated how this type of neuron model could be trained to produce a spike-based classification of temporal patterns through a local, reward-modulated synaptic learning rule.

Full-text available at: appendix D, page 113ff

Bibliographic reference:

P. Nieters, **J. Leugering**, and G. Pipa, "Neuromorphic Adaptive Filters for event detection, trained with a gradient free online learning rule," presented at the Machine Learning Summer School (MLSS-Africa 2019), 1, 2019.

#5 **Title:** Neuromorphic computation in multi-delay coupled models.

Publication type: journal paper

Summary: In this paper, we explored how delayed feedback, in particular the interaction between differently delayed feedback-loops, can be exploited to endow a single neuron, which could be implemented in an electrical or photonic circuit, with memory and the capability to compute complex functions of its input history. For a simple single-node multi-delay-coupled reservoir neuron, we show how the relationship between the delay terms leads to different complexity of behavior, and hence different performance of the trained neuron across different time-series regression tasks. Curiously, we can show that — and why — co-prime delays result in the best performance, and thus give some intuition for the complex behavior of delay-coupled systems.

Full-text available at: http://ieeexplore.ieee.org/document/7921635/Bibliographic reference:

P. Nieters, **J. Leugering**, and G. Pipa, "Neuromorphic computation in multidelay coupled models," *IBM Journal of Research and Development*, vol. 61, no. 2/3, 8:7–8:9, 1, 2017, ISSN: 0018-8646, 0018-8646. DOI: 10.1147/JRD.2017.2664698.

#6 **Title:** A Unifying Framework of Synaptic and Intrinsic Plasticity in Neural Populations.

Publication type: journal paper

Summary: In this rather long paper, I explore the relationships and interaction of intrinsic and synaptic plasticity for computation. The entire chapter 5 of my thesis is largely based on ideas contained within this publication. I try to motivate the same main results here using a slightly different approach that introduces concepts like optimal transport theory. But for most content of the current chapter, a more in-depth discussion can be found within this original publication.

Full-text available at: https://www.mitpressjournals.org/doi/10.1162/neco_a_01057

Bibliographic reference:

J. Leugering and G. Pipa, "A Unifying Framework of Synaptic and Intrinsic Plasticity in Neural Populations," *Neural Computation*, vol. 30, no. 4, pp. 945–986, 17, 2018, ISSN: 0899-7667. DOI: 10.1162/neco a 01057.

#7 **Title:** Event-based pattern detection in active dendrites.

Publication type: journal paper (pre-print)

Summary: In this manuscript, we derive a simple yet powerful mechanisms of dendritic computation in single neurons from first biological principles. Our model makes use of actively generated dendritic plateau potentials, which provide the neuron with distributed processing elements and memory traces that collectively allow a single neuron's dendritic tree to process information in nonlinear ways and on timescales that exceed the typical timescales of membrane potentials by orders of magnitude. We show how this event-based mechanism can be used to reproduce well known nonlinear computations when viewed from a rate-coding perspective, but also how it goes much further than that by detecting specific long-lasting sequences of spike volleys and integrating information from a vast number of inputs over comparatively long time-scales. A pre-print of this paper is publicly available, and a revised version of the same manuscript is currently still under review.

Full-text available at: appendix D, page 114ff

Bibliographic reference:

J. Leugering, P. Nieters, and G. Pipa, "Event-based pattern detection in active dendrites," *bioRxiv*, 17, 2020. DOI: 10.1101/690792v3.

#8 **Title:** *Making spiking neurons more succinct with multi-compartment models.*

Publication type: conference paper

Summary: In this conference paper, which accompanies a full-length presentation, I analyze the computational properties of the biologically motivated multi-compartment neuron model of contribution 7 from a machine-learning perspective. By transferring and adapting concepts and learning rules developed for decision trees to this neuron model, I give an intuition for how such a hierarchical structure like a neural dendrite can be useful for computation, and how simple, local learning rules might be enough to optimize such models.

Full-text available at: https://doi.org/10.1145/3381755.3381763 Bibliographic reference:

J. Leugering, "Making spiking neurons more succinct with multicompartment models," in *Proceedings of the Neuro-Inspired Computational Elements Workshop*, 17, 2020, ISBN: 978-1-4503-7718-8. DOI: 10.1145/3381755.3381763.

#9 **Title:** Neuromorpher Musterdetektor und neuromorphe Schaltkreisanordnung hiermit.

Publication type: patent (filed & pending)

Summary: Based on the insights derived from the neuron model of contribution 7, we designed a digital neuromorphic circuit that can efficiently realize the computation required for active dendritic sequence processing without the need for any general purpose processing elements like arithmetic-logic-units or micro-processors. It implements a processor for temporal patterns and sequences in each hierarchically structured neuron through a combination of pulses of different lengths, just like its biological counterpart. Homogeneous assemblies of multiple such neurons then communicate with each other through a code that serializes and transmits multiple spike-trains over a single binary connection.

Full-text available at: appendix D, page 131ff

Bibliographic reference:

J. Leugering, P. Nieters, and G. Pipa, "Neuromorpher Musterdetektor und neuromorphe Schaltkreisanordnung hiermit," patent application DE 10 2019 134 044 A1.

#10 **Title:** A Bayesian Monte Carlo approach for predicting the spread of infectious diseases.

Publication type: journal paper

Summary: In this publication, we investigate a simple class of probabilistic models of the spread of infectious diseases using Hamiltonian Markov Chain Monte Carlo sampling techniques and spatio-temporal kernel functions. Since this work is thematically disconnected from the central focus of this dissertation, it is referenced here for completeness but not discussed further.

Full-text available at: https://journals.plos.org/plosone/article?id =10.1371/journal.pone.0225838

Bibliographic reference:

O. Stojanović, **J. Leugering**, G. Pipa, S. Ghozzi, and A. Ullrich, "A Bayesian Monte Carlo approach for predicting the spread of infectious diseases," *PLOS ONE*, vol. 14, no. 12, e0225838, 18, 2019, ISSN: 1932-6203. DOI: 10.1371/journal.pone.0225838.

D

Full-text sources of further contributions

In the following, I have compiled unpublished or inaccessible contributions referenced in the thesis. The documents are included "as-is" in the exact form they were submitted or presented (except, of course, for the added page numbers and legal notices).



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A visit to the neuromorphic zoo

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Abstract— Over the course of the last decade, neural networks have finally found their way from mostly academic research into commercial applications. So far, this transition has taken place primarily behind closed doors at high-performance computing centers — but with ever more powerful mobile devices and a growing interest in the Internet of Things, a similar revolution is ahead of us in the embedded device market. One technology that takes center stage in these developments is specialized neuromorphic hardware, custom designed for executing neural network applications. In this paper, we would like to provide some background information on this fascinating branch of hardware development for interested readers from other disciplines, compare different approaches and provide an overview of the current state of the field.

Keywords—neuromorphic hardware; embedded AI; neural networks; deep learning; accelerators

I. THE DEEP LEARNING REVOLUTION

The last decade has brought with it a remarkable transformation of the field of artificial intelligence. Starting in 2012, when a Deep Neural Network now known as AlexNet [1] beat all competing approaches in a highly competitive computer vision challenge for the first time, Deep Learning has enjoyed a stellar rise in popularity - both in academia and industry alike. Today, it has become a ubiquitous and indispensable tool for a broad range of applications: Unlocking a mobile phone through facial recognition [2], controlling it through voice [3, 4], or having it translate a website [5] likely relies on a deep neural network in the background to bear the brunt of the work. On the other end of the spectrum, powerful servers and clusters employ deep neural networks for the automated analysis of big medical data sets [6], economic forecasts [7] or epidemiological predictions [8] etc., and ever-more complex neural networks are being developed to tackle increasingly harder problems.

In the wake of this "Deep Learning revolution" and as we mourn the gradual decline of Moore's law [9], there is a growing demand for innovative hardware solutions to sustain this development – even as we approach the limits of established technologies. In particular *neuromorphic hardware*, which is often subsumed with other approaches under the generic umbrella terms *next generation computing* (NGC) or *non-von-Neumann computing*, promises to deliver critical performance benefits that pave the way for further market adoption of machine learning and artificial intelligence. But what exactly *is*

neuromorphic hardware, and how does it work? Why is it becoming increasingly relevant today, and where might it lead us in the future? We'll discuss these questions in the following, starting with a little bit of background.

II. WHAT EXACTLY IS NEUROMORPHIC HARDWARE?

The term "neuromorphic" is obviously a portmanteau of "neuro-" and "-morphic", and it describes hardware that is in some way inspired by the *morphology* of biological *neural* systems. Since biological inspiration can take many forms and can be taken to different levels, the label "neuromorphic hardware" is applied rather loosely to an entire research field that shares the common objective to *implement (some) neural network models efficiently in hardware*. The best way to achieve this varies, depending on the type of neural network to accelerate, design constraints and the performance criteria that have to be optimized, e.g. power, latency, noise robustness etc. There are many degrees of freedom in the design process, among them:

- What type of networks should be used, e.g. neurons with continuously valued states that change continuously with time, or with discrete states that are updated at discrete time-steps, or spiking neurons that communicate asynchronously?
- Should these computations be realized by analog or digital circuitry?
- Should the *topology of the networks* be restricted to some specific structure, e.g. to feed-forward, recurrent or convolutional networks?
- Which components of the network should be directly implemented by dedicated hardware components, and which, if any, should be emulated with a more conventional processor design?
- Can the design be decomposed into functional modules, and if so, how should these modules communicate with each other?

Any combination of these (and many more) design choices leads to a different species in zoo of neuromorphic hardware designs. In practice, this results in a broad continuum from specialized multi-processor designs on the one end, all the way to full analog instantiations of biological neural network models

on the other end, where each neuron and synapse of the neural network model has a corresponding dedicated electrical counterpart. Neuromorphic hardware therefore encompasses several diverse technologies, rather than any one in particular, so an exact technical definition of the term is difficult. Instead, we can identify some characteristic features, or *design principles*, that distinguish (most) neuromorphic hardware designs from more conventional approaches:

- *highly parallel* computing instead of to the sequential operation of a single central processing unit (CPU)
- the use of distributed and decentralized memory instead of a central dedicated storage
- a system design specifically *optimized to implement neural networks* of some form.

III. HOW CAN/DOES NEUROMORPHIC HARDWARE WORK?

Since neuromorphic hardware is designed to accelerate neural networks, its merits can only be explained in the context of neural networks. Luckily, the mathematical model underlying most machine-learning applications of neural networks is actually very simple. To get everyone on the same page, what follows is a (hopelessly incomplete) high-level glance at the theory of neural networks, focusing only on those aspects that are relevant for neuromorphic hardware designers. For a more complete introduction, we refer to [10] and references therein. Everyone who is familiar with the theory already can safely skip right ahead to the next section, where we then discuss different approaches to implementing such network models in hardware.

A. Excursion: How do (deep) neural networks work?

In the machine-learning context¹, a neural network is a graph structure composed of *neurons* that are connected by *synapses*. Each synapse can multiplicatively scale its input, which may be the output of some neuron or an external source, by some number called the synapse's *weight*, and transmit the result to another neuron. Each neuron linearly combines all of the signals from its incoming synaptic connections (scaled by their respective weights) into a single signal. This signal is then nonlinearly transformed to produce the neuron's output.

Mathematically [10], the neuron j's output $y_j(t)$ at time t is therefore a function of the input signals $x_i(t)$, and has the simple form $y_j(t) = f(\sum_i w_{j,i} x_i(t) + b_j)$, where $w_{j,i}$ represents the scalar weight of a synaptic connection from input x_i to neuron j, and b_j is a neuron specific offset or bias term. We can group all the weights and bias terms in a single weight matrix W and the bias vector \vec{b} , which results in the simple matrix equation $\vec{y}(t) = f(W\vec{x}(t) + \vec{b})$. From this equation, it should become clear, that the multiply-accumulate operations (MACs) required for matrix-vector multiplication constitute the main computational cost for simulating neural networks.

One major exception to this story are *spiking neural* networks (SNNs, [11]), which encode a neuron's output instead

by a sequence of brief, stereotypical pulses (*spikes*). In these networks, much like pulse code modulate (PCM, [12]) in digital signal processing, it is the *number of spikes per unit time* that conveys the magnitude of a signal, not the amplitude. The corresponding mathematical model requires an explicit representation of time and is hence best suited for the processing of time-series.

Despite the simple mathematical formalism, many types of neural networks can be realized by different classes of weight matrices. To give a few examples, a (block)-triangular weight matrix represents a (layered) feed-forward network. Diagonal blocks represent recurrently connected layers (i.e. groups of mutually connected neurons), and all blocks on the second or higher off-diagonal represent so-called skip connections. Offdiagonal blocks in Toeplitz-form resemble convolutional layers a structure that has proved to be invaluable in image processing tasks, and so on. Besides the structure of the weight matrix, we can also choose what numeric type of entries it should contain. For example, instead of real valued weights, we can use integer valued weights, pick from an arbitrary set of discrete weights, e.g. binary ($w \in \{-1,1\}$) or ternary ($w \in \{-1,0,1\}$), or even use a compressed encoding of the weights. To summarize, there are countless interesting classes of weight matrices, and each of them has specific implications for the corresponding class of neural networks and offers specific opportunities for hardware acceleration.

In order for a neural network to do anything useful at all, the free parameters, i.e. weights and bias terms, must be set to specific task-dependent values. Earlier engineering approaches like the famous *Neocognitron* [13] anticipated many of features of deep neural networks, but relied heavily on domain-knowledge and inspiration from the network structures observed in nature². The real breakthrough happened decades later, when improvements in computer technology suddenly made it feasible to directly optimize (or *train*) the weight matrices (and bias terms) of highly structured networks (e.g. deep convolutional feed-forward networks [1]) to minimize errors (or *loss*) on extremely large data-sets.

Since global optimization of such large non-linear systems is near impossible, the work-horse for the optimization of deep neural networks are simple, greedy, gradient-based algorithms, that differentiate the loss function on a training data-set with respect to the network's parameters, and use this information to iteratively improve the parameters [10]. While this is significantly more difficult for spiking neural networks (the discrete-time nature of their event-based communication makes the calculation of gradients difficult), some remedies exist [14, 15] that allow us to use similar tools even for training spiking neural networks. As a result, gradient-based methods have *de facto* become such a central part of deep learning, that *differentiable programming* has even been suggested as a more accurate label for the entire field [16].

¹ We only discuss neural network models for machinelearning, not bio-physically accurate models of nervous systems.

² The study of these biological and artificial *connectomes* under the name *connectionism* was the intellectual precursor of modern deep learning.

B. The Neuromorphic Zoo

As this brief overview/recap of neural networks hopefully shows, there are many knobs to turn in the construction of neural networks, and the possible hardware implementations are similarly plentiful. In the following, let's have a brief look at several different approaches to neuromorphic hardware design. We stay on a rather high conceptual level here, and discuss five clusters of approaches, grouped by the degree to which the network model is directly reflected in the hardware. A more complete overview and a more in-depth discussion of the various underlying hardware design concepts can be found e.g. in [17, 18]. We begin with conventional computing devices, and end with truly neuromorphic, fully analog designs that replicate each individual synapse in hardware.

1) Generic co-processors and graphics cards

Since a major fraction of the simulation and training time for neural networks is spent on MAC operations, the key innovation in most accelerator designs is an efficient hardware implementation of matrix multiplication for some class of weight matrices. Arguably the most flexible and generic form of hardware accelerators for neural networks are therefore conventional many-processor designs like graphics processing units (GPUs, [19]) or other "number crunching" co-processors like tensor processing/streaming units (TPUs, [20, 21]), which have been thoroughly optimized for large and fast matrixconsidered multiplications. They are generally not neuromorphic hardware, but the high demand of such devices for deep learning applications, among others, has driven the development of a new generation of GPUs and TPUs optimized entirely for generic parallel compute tasks, and software libraries [22, 23] have accordingly begun to delegate more and more parallel operations away from the CPU to such co-processors. However, their versatility comes at a high price: the ability to execute arbitrary programs requires an extensive control-logic, powerful arithmetic logic units and a cache-, memory- and bussystem optimized for arbitrary memory access and fast data transfer. This overhead is unnecessary for many neural network architectures and can lead to power, performance or latency penalties. In addition, since the main speed-up offered by such co-processors is through the acceleration of matrix multiplication, they yield hardly any benefit at all for certain types of networks like SNNs, extremely sparsely connected networks or networks with non-linear synaptic effects.

2) Custom many-processor designs

Similar in spirit, albeit more closely focused on neural network applications, are specialized many-core designs (e.g. [24, 25] among many others), that distribute the task of simulating or training a large neural network across many independent processor cores. They typically support a (reduced) instruction set tailored and optimized towards neural network applications. Rather than by arbitrary access to shared memory, these designs typically implement an efficient routing or message passing system for the exchange of information between the nodes. Despite the focus on neural network applications, the neurons are here emulated algorithmically in software, and data flows through a shared bus-system, rather

than dedicated synapses. Since these devices do not directly implement any of the components of a neural network in hardware, they are not genuine neuromorphic hardware in the narrowest sense³, but they are generally discussed alongside neuromorphic hardware due to their near-identical application areas and user interfaces.

3) Digital deep learning accelerators

There is another class of digital accelerators (e.g. [26, 27, 28]), that is designed and optimized on a low level entirely for the implementation of (some) deep neural networks. Here, the operation of individual neurons is approximated by a dedicated digital logic circuit, that realizes the specific MAC operations required for the network class of interest (respecting the structure as well as the bit-precision the neurons' input weights and activation functions). These devices therefore don't allow arbitrary code execution, but instead require provisioning with the precise topology and coefficients of the network. Once configured, they act as a black box that efficiently executes the provisioned network, mapping digital input signals onto the network's digital outputs.

Rather than on their own, such deep learning accelerators can be used as small cores embedded within a larger many-core system similar to the ones discussed above. Such a modular design can be more effective, flexible and easier to scale, in particular when a specific network architecture to be accelerated allows for specific optimizations. For example, convolutional neural networks re-use the same structure of synaptic weights repeatedly for different neurons (also called weight-sharing), which can be implemented very efficiently by re-using the same hardware substrate in a time-multiplexed design that updates the network one neuron at a time. Generally in feed-forward networks, the neurons within one layer are conditionally independent given their input and can thus be processed in parallel. Here, an efficient hardware solution could update one entire layer at a time. For sparse weight matrices, an optimized handling of zeros can further improve performance, while networks with low-precision (e.g. ternary) weights can be implemented by much more compact circuits.

4) Analog deep learning accelerators

As we saw before, the mathematical models of neural networks are typically given in terms of real numbers. So rather than approximating them via discrete digital circuitry, another natural approach is to instead represent the real-valued quantities of the model by real-valued physical quantities like analog voltages, currents or charges. Such a use of analog circuit design goes back to the earliest attempts of neuromorphic hardware design in the 1950s [29], but fell out of favor during the digital revolution in electronics. While the susceptibility to noise is still a major challenge for most applications of analog circuitry, some neural networks have, quite surprisingly, proven to be remarkably robust to the effects of noise [30] – in fact, some forms of noise may even *help to improve* the robustness of the system [31]! Other pit-falls of analog circuit design, e.g. the difficulty of precisely controlling (non-)linearities in the system,

³ In fact, they are not limited to neural network applications at all, and can be used for other tasks with similar demands.

are much less critical for neural networks than other applications, because the networks have sufficient degrees of freedom to counteract such defects (provided, of course, that the defects are known). Despite these challenges and limitations, which ultimately caused the transition to digital circuit design, there are of course also major benefits for the implementation of neural networks in the analog domain. First, our continuous model of a neuron is remarkably similar to that of a logic gate⁴ - one might even view it as a continuous, weighted extension of logic gates - with one crucial difference: neurons can be differentiated with respect to their parameters, which, as we have seen above, is critical for deep learning [10]. Deep learning therefore provides a framework to optimize analog circuits in a way that cannot be directly applied to digital circuits. While it is of course possible to approximate the behavior of continuous neuron models by digital circuitry (see above), this can result in a high component count of transistor and logic gates, each of which has itself a complexity rivaling that of an analog implementation of the neuron model [17]. In the analog approach, multiplication and addition is instead realized by direct application of Ohm's and Kirchhoff's laws, i.e. by choosing appropriate values of resistive elements to represent individual synaptic weigths and accumulating the resulting currents. This also allows for ultra-low power applications, possibly at the expense of an increased noise-floor, and alleviates the need to wait for signals to settle, which makes lowlatency asynchronous designs possible.

Due to strong barriers to entry, e.g. high manufacturing costs and long development cycles, only a comparatively small number of fully analog deep learning accelerators have actually seen the light of day. However, a vast amount of literature has been written about this already (see e.g. [17, 18] and references therein) and if the number of recent start-ups and research projects in that field is any indication, there is a substantial and growing commercial and academic interest, as well.

5) Spiking neural network accelerators

Last but not least, neuromorphic hardware for spiking neural networks, currently an outlier in the machine learning world, is set to become another major branch of hardware accelerated embedded AI. Contrary to conventional neural networks, the purely event-driven operation of spiking neural networks defies the simple mathematical frameworks of continuous function approximation and periodic sampling. An efficient implementation of such networks is therefore difficult for both clocked digital logic as well as in conventional imperative programming paradigms. Combined with the increased complexity of training algorithms for SNNs [14, 15, 32], this may explain the relative lack of attention these networks have received within the machine learning community – despite advocacy by some leading theoreticians in the field [11]. However, the very same properties of spiking neurons that appear as major obstacles for efficient software implementations (e.g. integration and low-pass filtering of signals over time and rising-edge triggered generation of pulses) are commonplace [12] in signal processing and can be implemented by simple analog circuits [33]. As a result, there is little overhead in complexity when switching from an analog to a spike-based network design. To the contrary, since each neuron's spiking output is a binary signal that can be converted into an analog signal merely by low-pass filtering (one of the axioms of the neural engineering framework [34]), spike-based neuromorphic hardware can combine the best of both worlds: the highly energy-efficient computation of analog circuitry and the binary transmission of signals via spikes, which decreases susceptibility to noise and simplifies routing and buffering.

Just as for digital hardware accelerators, the communication between individual neurons of a spiking network can therefore be implemented either through dedicated electrical lines or through a (digital) package routing system, the most popular of which is *address event representation* [35] encoding, where each spike is conveyed as a package containing the "address" of the neuron from which it originated. While such a routing system greatly improves the scalability of the system by time-multiplexing the usage of the same communication channels, it requires sophisticated scheduling and low latencies that can become prohibitive as the number of interconnected neurons increases. A hybrid approach that uses many cores with full *internal* connectivity through dedicated lines, connected *to each other* via a common bus system, is therefore a popular compromise (e.g. in [36, 37]).

IV. WHY IS NEUROMORPHIC HARDWARE RELEVANT TODAY?

With this brief overview in mind, one might wonder, why neuromorphic hardware has so suddenly become a hot topic among AI researchers and ASIC developers. None of these ideas seem novel enough to justify this rise in popularity — in fact, similar ideas have been continuously suggested since the very beginning of artificial intelligence research and computer science in the 1950s [29]. Even the term "neuromorphic hardware" was popularized already in the 1990s by Carver Mead [38], who has been a pioneer in this research area since its early days. So why should we invest in neuromorphic hardware today, and why didn't this happen before?

The most obvious argument is purely opportunistic: never before have neural networks had sufficient size to be practically useful for complex, data-driven applications. Now, with the breakthroughs in image classification competitions during the recent years, neural networks have finally proven their worth for commercial applications, and have received massive exposure to the public and industry ever since. This growing popularity has correspondingly lead to an increased demand of efficient hardware on which to run neural networks.

The applications in fields like image processing [1], gaming [39], text analysis [40], audio processing [41, 4] and datascience, e.g. in medical image analysis [6], have diversified and become more complex, with weight coefficients numbering anywhere from hundreds of thousands up to a staggering billion

⁴ The earliest theories of neural networks by McCulloch & Pitts [67] already established this connection.

for extreme cases [42]⁵. The range of applications is likely only going to increase, as a growing number of mobile devices from smartphones [43] all the way to autonomous vehicles do already (or will soon) use neural networks for demanding image recognitions tasks, and require the corresponding computing power.

So far, we have been riding the wave of ever-improving CPUs and GPUs, and technological progress alone could sustain the growing demand, but as we begin to witness the end of Moore's [9] law, we need fundamentally new ideas. Current state-of-the-art 7nm CMOS technology approaches physical limits and it seems unlikely that we can continue scaling down size, power-consumption or latency much further. At the same time, the total training time, power-consumption and initial cost of systems capable of simulating larger state-of-the-art neural networks has sky-rocketed to an unsustainable level [44], while the limited power budget of mobile devices has been a limiting factor for many potentially interesting applications.

Of course, neuromorphic hardware, too, has benefitted from the technological advances in electronics manufacturing during the last few decades. Besides a new market, the availability of new technologies is thus another reason for a renewed interest in neuromorphic hardware development. For example, new transistor design principles like fin field-effect transistors (FinFETs, [45]), fully depleted silicon on insulator (FD-SOI, [46]) and floating multi-gate MOSFET transistors [47, 48] as well as special neuro-transistors (vMOS, [49]) have enabled extremely low-power applications and novel neuromorphic hardware designs. Since neural networks require a sizeable amount of memory for storing the network topology and synaptic weights, neuromorphic hardware also stands to gain a lot from new trends in memory technology. With small feature sizes of 28nm and below and advances in dynamic RAM (DRAM) and static RAM (SRAM), it has become possible to store reasonably sized networks directly in silico, and process data right where it is stored. While this new paradigm of inmemory computing [50, 51] is by no means limited to neuromorphic hardware, the highly distributed structure of neural networks can leverage this advantage particularly well, and thus overcome the memory-bottleneck that conventional von-Neumann architectures suffer from.

Since the network coefficients (typically) do not change at all during inference, emerging non-volatile memory technologies (eNVM, [52]) are particularly interesting for neuromorphic hardware. One major development is the emergence of several forms of *memristive* devices, which can act as the programmable resistive components required for reconfigurable analog hardware accelerators [53]. Competing technologies like charge trap flash memory (CTF), ferroelectric field-effect transistors (FeFETs), resistive RAM (ReRAM), conductive bridge RAM (CBRAM) and phase change memory (PCM) [52, 54] all exploit different physical phenomena to allow non-volatile storage on chip, many of them supporting the storage of analog values at a multi-bit resolution

[51], which is critical for analog hardware accelerators and reduces the die-space required for memory.

Lastly, we have learned a lot about neural networks in the meantime: we have demonstrated their capability, know now that there are indeed use-cases for large networks, and we have found better ways and tools [22, 23] to train even large networks. As we continue to learn more in-depth about which network topologies are effective for which specific tasks, and why, we also develop a better understanding of what sort of networks are worth accelerating in hardware – and which are not.

V. WHERE IS THE DEVELOPMENT OF NEUROMORPHIC HARDWARE HEADED IN THE NEAR FUTURE?

Of course, predicting the future is hard, particularly for a field that has been through all the season (including the dreaded, recurring "AI winter") several times already. But with new exciting technologies on the horizon, such as 3D [55] and wafer-scale [56] integration, nano-wire transistors [57], carbon nano-tubes [58] on-chip, silicon photonics [59], spintronics [60], ever-smaller micro- and nano-electro-mechanical systems (MEMS, NEMS [61]), integrated sensor-processor systems and more, it is hard to not feel optimistic about the future of neuromorphic hardware.

The technological progress is likely to bring neuromorphic hardware into new application areas, where it can reduce energy consumption, latency or the cost of existing solutions. For example, co-processors for AI are already being included in modern smartphones [43] to reduce CPU load during AI applications and therefore prolong battery life. On the other end of the spectrum, the increasing demand for high-performance computing clusters and cloud-services that provide "deeplearning-as-a-service" [62] shows a market for server-side energy efficient, dedicated neuromorphic hardware co-processors.

Neuromorphic hardware can also enable applications that are in principle possible right now, but not economically viable yet, such as natural language [63] or gesture based [64] user-interfaces for controlling a wide range of electrical devices, e.g. in the context of the internet-of-things or home-automation and appliances. Industry could use neuromorphic hardware to make even low-level processes in manufacturing more adaptive or responsive, or improve the interaction between humans and machines.

Finally, the adoption of neuromorphic hardware might even enable machine learning solutions that are flat-out impossible right now, such as many sophisticated real-time applications like the processing and fusion of complex, high-dimensional sensor data and the intelligent real-time control of sophisticated robots or production plants. Power savings might enable novel mobile applications like large-scale distributed sensor networks [65], or autonomous systems that are smart enough to act on their own and resilient enough to survive in difficult environments. By optimizing die-space, neuromorphic hardware could also find its way into miniaturized sensors, e.g. ingestible medical sensors

⁵ It should be recognized that despite the general trend of increasing network sizes, there are also contrary efforts to reduce the number of parameters, e.g. [68].

[66] and much more. Of course, the most important applications might turn out to be entirely different from the ones listed here, but we are confident that industry and academia will find countless ways to capitalize on neuromorphic hardware in the future!

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Neuromorphic Adaptive Filters for event detection, trained with a gradient free online learning rule

Machine learning problems are typically framed in a regression, classification or prediction setting, where a set of distinct data points is to be identified with corresponding labels. Artificial neural networks excel at such problemse, because their universal function approximation capability and differentiability can be leveraged for powerful gradiente-based optimization algorithms. Neuromorphic hardware however, interacting with its environment in real time, faces challenges that defy this framework. One such example is the detection of specific events in real time, where the mapping from a continuous stream of noisy input signals onto a discrete set of events is to be learned.

The temporal dimension of this task entails a **credit** assignment problem for learning, since the detector must evaluate a history of input signals and needs to be afforded some flexible **processing delays**, which makes defining a differentiable loss function for the event detection task difficult. This is aggravated in a setting where the **target** signals themselves are delayed. The constraints of neuromorphic hardware design further restrict the available learning algorithms to "any-time" computations implementable just by (traces of) locally available information, which precludes many of the established gradient-based optimization procedures.

We propose a neuromorphic event detector, the **Neuromorphic Adaptive Filter** (**NAF**) and ensembles thereof, that utilizes **Gamma Filter** banks [4,5] to learn a parameterized multidimensional signal filter through a **supervised gradient-free online learning rule**.

State & Weight Traces:

$$\dot{x}_0^j = \alpha \cdot (I_j - x_0^j)$$

$$\dot{x}_i^j = \alpha \cdot (x_{i-1}^j - x_i^j)$$

$$\dot{\bar{w}}_i^j = \eta \cdot \left(x_i^j \cdot \left(\frac{y-1}{2}\right)^2 - \bar{w}_i^j\right)$$

Derived Variables:

$$\nu = \sum_{j} \sum_{i} x_{i}^{j} w_{i}^{j}$$

$$I_{\rm fb} = y = \operatorname{sat}(\frac{\nu - \mu}{\sqrt{2\sigma^2}})$$

Update on signal:

$$\begin{aligned} \theta &\leftarrow \theta + \tau^+ \\ r &\leftarrow r + 1 \\ w_i^j &\leftarrow w_i^j - \kappa \cdot \bar{w}_i^j \\ w &\leftarrow \frac{w}{||w||} \end{aligned}$$

Intrinsic Plasticity:

 $\dot{\sigma}^2 = \beta \cdot ((\nu - \mu)^2 - \sigma^2)$ $\dot{\theta} = -\gamma \cdot \theta$

Signal condition:

 $y \ge \theta + r$

 $\dot{\mu} = \beta \cdot (\nu - \mu)$

Update on target:
$$\theta \leftarrow \theta - \tau^-$$

$$w_i^j \leftarrow w_i^j + \kappa \cdot \bar{w}_i^j$$

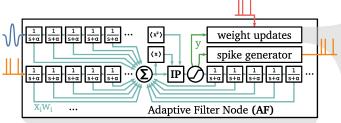
$$w \leftarrow w$$

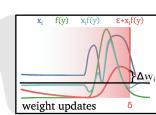
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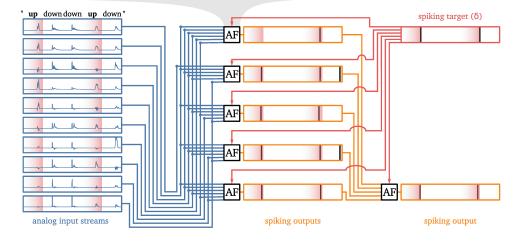
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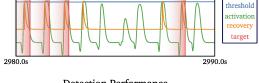
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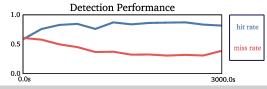






Early dynamics threshold activation recovery target 10.0s Trained dynamics threshold activation activation recovery target





♦Speech command detections with ENAF.

A continuous audio stream is represented by its spectral power in 10 Mel-adjusted frequency bands. 5 NAFs each receive as input a subset of 6 random chosen frequency bands. Their spiking outputs are used as input signals for another NAF, which then provides the ensemble output. Each occurrence of the desired word ("UP") in the audio stream is shortly followed by a spiking target signal to each ensemble member.

⟨=NAF performance during training.

Detection rates increase while false positive rates decrease during training as the ensemble members adjust. Early on, the filter responses within an individual NAF show little specificity for the target word ("UP"), but towards the end of training, the NAF has become selective and the threshold is set to provide a reasonable trade-off between false-positive and missed detections.

Key properties of the model:

- It can be formalized as a **system of ordinary differential equations** with discontinuous updates at distinct time points.
- Event signals are generated as output akin to spiking neuron models, and target signals are similarly provided as a sequence of events.
- The learning rule, similar in spirit to the Tempotron-rule[1], in concert with weight normalization adjusts the shape of the learned filter.
- The learning mechanism can be thought of as enforcing an "energy budget", where parameter updates are performed only at two kinds of distinct events: the emission of a detection signal incurrs a penalty, leading to a reduction in the weights of contributing features, whereas at each target signal, the weights assigned to those features that recently contributed to high membrane potentials are amplified.
- An adaptive threshold paired with intrinsic

- plasticity mechanisms in- or decreased the detector's sensitivity in response to target or emitted signals, respectively, thus adjusting the trade-off between false positives and false negatives.
- An individual NAF can be viewed as a variation of a generalized functional linear model [6] and thus represents a reasonable week learner for machine learning applications.
- Performance and robustness can be enhanced by routing the output of multiple NAFs with diverse inputs through another NAF capable of compensating the different incurred processing delays, thus forming an ensemble (ENAF).
- The training and inference procedure can be realized purely by instantaneous local computation and exponential memory traces.
 (E)NAFs are thus good candidate systems for neuromorphic hardware implementation.

EVENT-BASED PATTERN DETECTION IN ACTIVE DENDRITES

PREPRINT

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ABSTRACT

Many behavioural tasks require an animal to integrate information on a slow timescale that can exceed hundreds of milliseconds. How this is realized by neurons with membrane time constants on the order of tens of milliseconds or less remains an open question. We show, how the interaction of two kinds of events within the dendritic tree, *excitatory postsynaptic potentials* and locally generated *dendritic plateau potentials*, can allow a single neuron to detect specific sequences of spiking input on such slow timescales. Our conceptual model reveals, how the morphology of a neuron's dendritic tree determines its computational function, which can range from a simple logic gate to the gradual integration of evidence to the detection of complex spatio-temporal spike-sequences on long timescales. As an example, we illustrate in a simulated navigation task how this mechanism can even allow individual neurons to reliably detect specific movement trajectories with high tolerance for timing variability. We relate our results to conclusive findings in neurobiology and discuss implications for both experimental and theoretical neuroscience.

Author Summary

The recognition of patterns that span multiple timescales is a critical function of the brain. This is a conceptual challenge for all neuron models that rely on the passive integration of synaptic inputs and are therefore limited to the rigid millisecond timescale of post-synaptic currents. However, detailed biological measurements recently revealed that single neurons actively generate localized plateau potentials within the dendritic tree that can last hundreds of milliseconds. Here, we investigate single-neuron computation in a model that adheres to these findings but is intentionally simple. Our analysis reveals how plateaus act as memory traces, and their interaction as defined by the dendritic morphology of a neuron gives rise to complex non-linear computation. We demonstrate how this mechanism enables individual neurons to solve difficult, behaviorally relevant tasks that are commonly studied on the network-level, such as the detection of variable input sequences or the integration of evidence on long timescales. We also characterize computation in our model using rate-based analysis tools, demonstrate why our proposed mechanism of dendritic computation cannot be detected under this analysis and suggest an alternative based on plateau timings. The interaction of plateau events in dendritic trees is, according to our argument, an elementary principle of neural computation which implies the need for a fundamental change of perspective on the computational function of neurons.

Introduction

- 2 The ability to detect long-lasting sequences of neural activity is crucial for complex behavior, but poses a serious
- 3 challenge for most established neuron models. Consider a rodent navigating through an environment in search for
- 4 food. Receptive fields of place and grid cells tile a spatial map of the environment and encode the current position by
- 5 their respective population activities [1, 2]. But in order to find its way back, the animal needs to know not only its
- 6 present location, but also which path it took to get there. Decoding this path from the sequential activation of place and
- 7 grid cells requires the integration of information on behavioural timescales that can span hundreds of milliseconds or

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- more [3, 4]. Relevant patterns on such long timescales may prove to be a ubiquitous phenomenon, and have already been documented for a wide range of sensory processing tasks, such as olfaction [5, 6] or cortical auditory processing [7].
- This raises the puzzling question, how such long sequences of neural activity can be processed by volatile neurons with membrane time constants on the timescale of tens of milliseconds or less [8]. While this problem is typically addressed on a network level, e.g. by relying on effects of fast-acting synaptic plasticity [9] or slow emergent dynamics due to recurrent connections [10], we argue that it can be solved on the level of individual neurons by active processes within their dendritic trees. These localized processes endow neurons with internal memory traces on the timescale of hundreds of milliseconds, and can be captured in a simple, conceptual model that adheres to recent biological evidence not accounted for in integrate-and-fire neuron models.
- 17 By investigating the computational properties of neurons with active dendrites, we draw three conclusions.
- Firstly, active dendritic processes can implement complex spatio-temporal receptive fields for ordered sequences of synaptic inputs. Secondly, active dendritic processes enable the robust integration of weak signals over timescales much longer than post-synaptic responses. Thirdly, when analyzed from a rate-coding perspective, active dendritic processes implement sophisticated non-linear computations that are characterized by the neuron's dendritic morphology.
- We demonstrate these propositions in a general computational framework for event-based, active dendritic sequence processing (ADSP), which offers an elegant solution to the problem of detecting highly variable, long lasting patterns in a neuron's input.

The functional role of active dendritic processes.

We derive our abstract model of dendritic computation from a few basic biological observations: Most of a cortical pyramidal neuron's excitatory synaptic inputs terminate on dendritic spines [11], where post-synaptic ion channels are activated via the stochastic, pre-synaptic release of glutamate-carrying vesicles [12, 13]. The activated channels, primarily controlled by α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid receptors (AMPARs) [14], become conductive to a mixture of ions, which leads to a brief depolarization in the corresponding spine, referred to as the excitatory post-synaptic potential (EPSP) [15]. These voltage changes in nearby spines induce a modest depolarization in the local dendritic membrane potential [16], which passively propagates along the dendrite as described by neural cable theory (Fig. 1c). For very specific branching patterns, the passive propagation of activity along a neuron's dendrite can be simplified to an equivalent model of a cylinder, in which the contribution of individual synaptic inputs sum (sub-)linearly [17]. Since propagation along the cylinder is very fast, abstract point-neuron models such as leaky integrate-and-fire neurons ignore the spatial dimension of the dendritic tree entirely and model the neuron as if it were a single electric compartment [18]. However, in this purely passive model of dendritic integration, the attenuation of signals along the dendritic cable is so strong, that synaptic input onto thin apical dendrites should have little, if any, measurable effect on the membrane potential at the soma far away [19, 20]. A synaptic plasticity mechanism that proportionally up-scales synaptic efficacies depending on the synapses' distance to the soma may counteract this phenomenon. Aptly termed "dendritic democracy" [21], it has been shown in hippocampal pyramidal neurons [22], where it results in a similar contribution of synaptic inputs onto the somatic membrane potential — regardless of the synapse's position along the dendrite. We instead look at a different mechanism to boost weak synaptic inputs, which relies on localized depolarizations that are actively generated and maintained within the dendritic tree.

Such active dendritic processes are ubiquitous [23, 24] and largely rely on N-methyl-D-asparate receptor (NMDAR) 45 gated ion-channels [14] (see Fig. 1c for a schematic representation of this mechanism). NMDAR gated channels, like their AMPAR gated counterparts, are activated in the presence of glutamate, but do not become conductive unless a 47 channel-blocking Mg⁺ ion is first displaced by a sufficiently strong depolarization [25, 26]. This depolarization can be 48 achieved by the coactivation of multiple AMPAR channels on nearby spines within a short time-window. Experimental 49 as well as simulation studies report that this requires a volley of 4-20 or even up to 50 spikes within 1-4ms, depending on the location along the dendritic tree [16, 27, 28, 29]. The opening of NMDAR channels triggers a massive influx 51 of different ionic currents that lead to a complete depolarization of a small segment of the dendritic arbor. While the 52 isolated NMDAR response itself is reported to last on the order of at least 25ms [30], in vivo recordings reveal that 53 voltage-gated channels in the dendritic membrane [20] prolong this effect, resulting in a depolarization that can last 54 from tens to hundreds of milliseconds [31]. We focus on these longer lasting events, which we collectively refer to as dendritic plateau potentials, and argue, that they provide useful memory traces within the dendritic tree that can last 56 hundreds of milliseconds. 57

The much larger depolarization during a plateau potential propagates further along the dendrite than the weaker effect of individual EPSPs and thus extends the range at which they can contribute to somatic action potential generation. This may even be required for generating or spiking [32] or bursting [33] output. Just like EPSPs, however, plateau potentials are still subject to considerable attenuation along the dendritic cable and thus have a strong effect only in

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their direct neighbourhood². This leads to a division of complex dendritic arbors into functional subunits [34, 35, 36], 62 which we here refer to as dendritic segments. How local plateau potentials in these segments interact within a dendritic 63 tree depends on its morphology. In particular, the depolarizing effect on other directly connected dendritic segments is effectively raising their resting potential for the whole duration of the plateau potential, thus lowering the amount 65 of coinciding spikes required to initiate a plateau potential there [37]. As [38] demonstrates, this local nonlinear 66 interaction of dendritic segments due to NMDAR-gated channels can allow neural dendrites to become selective to 67 specific sequences of synaptic inputs. While their work uses a biophysical, spatially extended neuron model to explain this behaviour, we instead derive a much simplified model composed of discrete dendritic segments. This helps explain 69 how local interactions between connected segments lead to cascades of plateau potentials, which in turn allow the 70 detection of specific long-lasting sequences within the dendritic tree.

Each segment of a dendritic tree tends to receive strongly correlated volleys of spikes on clustered synaptic inputs from some subpopulation of neurons [39, 40]. We suppose, that such incoming spike volleys constitute elementary events that convey relevant information. Then, the morphology of the dendritic tree then determines how this information is processed and retained in memory, and thereby endows the ADSP neuron with an intricate computational function.

The interaction of active dendritic processes realizes event-based computation.

We construct an abstract mathematical model of active dendritic sequence processing, that is firmly rooted in the previous biological observations. Conceptually, the complex dynamics of dendritic membrane potentials are reduced to the interactions of two kinds of events, EPSPs and actively generated plateau potentials, in a tree structure of dendritic segments. Since both of these events result in localized stereotypical effects on the dendritic membrane potential, we abstractly model them simply as rectangular pulses of unit magnitude and fixed duration τ_{synapse} and τ_{dendrite} , respectively. Because the qualitative behaviour of the dendritic arbor is thus explained purely in terms of the locations and times at which EPSPs and plateau potential are initiated in its dendritic segments, our model concisely describes dendritic computation.

Only those incoming spikes that are successfully transmitted by the probabilistic synapses induce EPSPs in the postsynaptic segment, which sum up and constitute the total *synaptic input* into the segment. This input is particularly strong when a *volley* of multiple spikes occurs in a time-window short enough for their EPSPs to overlap. In addition to synaptic input, the electric coupling between directly connected dendritic segments provides another source of *dendritic input*.

When both the synaptic and dendritic input into a segment exceed critical thresholds, the segment enters a prolonged plateau state. For the whole duration of the plateau, all other directly connected segments receive depolarizing dendritic input. Segments of the dendritic tree therefore act as coincidence detectors that respond to highly synchronized volleys 92 of spikes with plateau potentials. The precise thresholds for synaptic and dendritic input depend on the segment's 93 location within the dendritic tree. While a large volley of spikes alone suffices to trigger a plateau in the outermost 94 segments of the dendritic tree, internal segments require the additional dendritic input due to plateau potentials in connected segments. For segments that lie at branching points in the dendritic tree, more than one of their neighbours 96 may have to be in a plateau state concurrently to have a sufficient effect. If the soma, which lies at the root of the 97 dendritic tree, receives sufficient synaptic and dendritic input, a somatic action potential, rather than a plateau potential, 98 is generated. 99

Since the small effects of EPSPs remain confined to the postsynaptic dendrite segment, they only affect the neuron's behaviour indirectly by contributing to the generation of local plateau potentials. It is the plateau potentials and their interaction across neighbouring segments that drives the dendritic membrane potential, and therefore implements an event-based framework of dendritic computation on two distinct timescales orders of magnitude apart. On a fast timescale, the combined effect of a volley of coincident spikes initiates a localized plateau potential. On a much slower timescale, the interaction of these plateaus provides an ephemeral memory of the recent history. The computation we have described here is fully formalized in terms of synaptic spikes and plateau events as provided in the Methods section.

In **Fig. 1** we describe an exemplary ADSP neuron that receives input from five populations of neurons on five segments (**Fig. 1a**). Each segment, if sufficiently excited, responds to a spike volley in its respective input populations by emitting a plateau event at the time of the volley (**Fig. 1b**). The morphology of the dendritic tree determines how these plateaus interact along the dendritic tree. For example, segment C will only activate if both segments A and B are already active once segment C receives a spike volley. We formalize the relative timing requirement for these three segments by the expression $(A + B) \rightarrow_2 C$, which indicates that all two child branches A and B must be simultaneously active to enable the parent segment C, allowing it to emit a plateau in response to a spike-volley. We read this as "A and B,

²Unlike EPSPs, this attenuation cannot be circumvented by synaptic scaling as for dendritic democracy.

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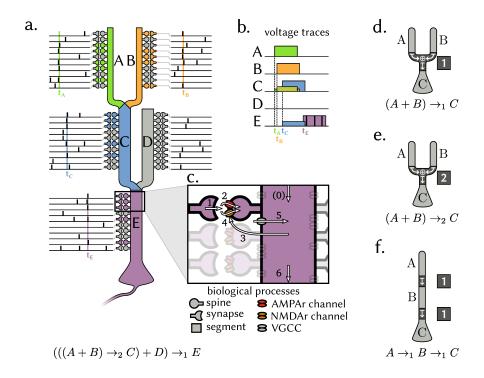


Figure 1: Schematic representation of a complex dendritic tree and its function. a A neuron receives on each of its 5 dendritic segments 10 synaptic connections from a corresponding neural population. Sufficiently many coincident spikes (here ≥ 6 out of 10) from population A lead the corresponding dendritic segment to generate a plateau potential (t_A) . Similarly, coincident spikes from population B induce a plateau in a parallel branch (t_B) . A third segment requires simultaneous input from both of these segments in addition to coincident synaptic input from population C, in order to fire a plateau of its own (t_C) . On another branch, a fourth segment receives its input from population D but does not trigger a plateau. A somatic spike is triggered when coincident synaptic input from population E arrives (t_E) during dendritic input from either of its two upstream segments (in this case C). **b** Local membrane potentials show a cascade of plateau potentials. c The steps involved in the generation of a plateau: The membrane potential is already elevated due to a plateau potential in a neighbouring segment (0). Presynaptic input arrives at a synapse (1), which leads to a postsynaptic EPSP via AMPAr mediated ion channels (2). Once the local membrane potential is sufficiently depolarized due to coincident EPSPs and prior depolarization, voltage gated, NMDAr mediated ion channels open, causing additional depolarization (4) which can be further facilitated by the opening of voltage gated calcium channels (5). This strong depolarization initiates a longer lasting plateau potential in the dendritic segment, which has a modest depolarizing effect on other neighbouring segments (6). Different dendritic morphologies correspond to different computed functions, indicated in the respective formula under each schematic illustration. d If activating one of two dendritic branches with input from either population A or B, followed by a somatic spike initiated by input from population C, is sufficient to produce a spike, the neuron implements the operation $(A+B) \to_1 C$, which constitutes an "or"-operation between population A and B. e If simultaneous input from A and B is required, the neuron calculates an "and"-operation between inputs A and B. f A simple neuron that requires sequential activation of first A "and then" B before C.

and then C" (see also **Fig. 1d**). If the threshold was lowered, such that input from either segment A or B alone would suffice, the expression would correspondingly become $(A+B) \to_1 C$, which translates to "A or B, and then C" (see also **Fig. 1 e**). Generally, the expression $(X_1+X_2+\ldots+X_n)\to_m Y$ translates to "At least m out of the n segments X_1,X_2,\ldots,X_n must be simultaneously active to enable segment Y". By chaining multiple segments together, these timing relations and nonlinear combinations can be arbitrarily nested, as for example in **Fig. 1f** that shows a neuron implementing $A \to_1 B \to_1 C$, which we read as "A, and then B, and then C". Using this formal notation, we express the complex ADSP neuron example in **Fig. 1a** as $(((A+B) \to_2 C) + D) \to_1 E$, a computation on spike volleys originating from the input populations associated with segments A,\ldots,E .

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The interaction between connected dendritic segments facilitates cascades of plateau potentials along the dendritic tree, 123 as illustrated in Fig. 1b. Starting in a distal segment, a leaf-node in our diagrams, a spike volley can initiate a plateau, 124 which then provides dendritic input for the parent segment. Next, that segment responds to an incoming spike volley 125 with a plateau of its own, in turn providing dendritic input to yet another segment. Whenever such a continuous chain 126 of plateau potentials proceeds all the way to the soma, it culminates in a somatic action potential. 127

This signals to other neurons, that a specific sequence of spike volleys has been detected – on a timescale that may 128 be as long as the number of segments times the plateau duration, i.e. hundreds of milliseconds. The precise timing between spike volleys is not prescribed exactly, as long as the distance between two successive volleys does not exceed 130 the duration of one plateau potential. This invariance is critical whenever the precise timing of the individual events can 131 vary, e.g. due to external circumstances such as varying movement speeds along a path in navigation tasks or due to 132 133 neural mechanisms such as sleep replay [41], because it allows the neuron to generalize over all such perturbations. The branching morphology of a dendritic tree therefore determines the computation performed by the neuron, which allows 134 even single neurons to detect complex compositions of sequential patterns. This event-based computation is what we 135 call active dendritic sequence processing (ADSP). 136

Results 137

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Dendritic processing allows the rapid detection of long, time-invariant patterns 138

To demonstrate the implications of such neuronal sequence detection, we return to the example of a rat navigating an 139 environment. We assume that the rat has an internal representation of its environment, tiled by the receptive fields of 140 distinct populations of place cells. While the animal resides within such a receptive field, the corresponding population 141 emits spike volleys with a magnitude that is largest when the animal is close to the center of the receptive field. Different 142 paths lead the animal through some of these receptive fields in different order, and result in different sequences of spike 143 vollevs. 144

Each individual spike volley consists of several coincident spikes, the EPSPs of which have to be integrated and 145 thresholded on a millisecond time-scale to detect sufficiently significant events in the presence of noise. To detect 146 whether the animal has taken a specific path through the environment, only specific sequences of such significant spike volleys must be detected on a much slower behavioural time-scale. These two distinct timescales pose a challenge for 148 conventional spiking neuron models, which is further exacerbated by the fact, that the precise timing of the spike-volleys 149 can vary substantially, depending e.g. on the speed with which the animal traverses its environment. While a solution to 150 this problem may be found on a population level, we illustrate in Fig. 2 how a single neuron can implement a solution very elegantly with just three active dendritic segments.

To simulate the rat's behaviour, we generate random movement trajectories through the environment by a stochastic 153 process (see Methods section). Each place-cell population fires spike-volleys with a magnitude determined by the 154 population's tuning-curve, a two-dimensional Gaussian function centered at the population's preferred location on a 155 hexagonal grid. In this example, we are interested in paths that traverse three specific receptive fields, respectively color-coded in blue, orange and purple, and hence look at a neuron that consists of a chain of three dendritic segments, 157 each receiving input from just one of these place-cell populations (Fig. 2b). The only trajectories that effectively 158 drive the neuron to spike are those that sequentially traverse the three receptive fields in the correct order Blue \rightarrow_1 159 Orange \rightarrow_1 Purple (**Fig. 2a**). 160

During the example path shown in solid black, the three place cell populations are activated in the correct order over the course of 200ms and emit sufficiently large spike volleys to trigger a cascade of plateau potentials that lead the neuron to emit a somatic spike Fig. 2b. To illustrate how reliable of a detector an individual neuron can be — even when its synaptic inputs are stochastic with a transmission probability of 0.5 —, we systematically evaluate the probability of the neuron to fire in response to different paths with varying directions and lateral offsets. For an ideal straight 200ms long path through the center of all three place cell populations, the firing probability of the neuron is around 75%. When the orientation of the path is varied, this probability sharply decreases to 0%, indicating that the neuron is both highly sensitive and highly specific for paths with this orientation (Fig. 2c). Similarly, when the path is shifted orthogonally to the movement direction, the response probability falls quickly, confirming that the neuron is sensitive to the absolute location of the path as well as its direction (Fig. 2d).

A remarkable feature of this mechanism is, that it is invariant to changes in the precise timing of the individual volleys 171 as long as two consecutive segments are activated within one plateau duration $au_{dendrite}$ of each other. The ADSP Neuron 172 can therefore detect paths of any duration from 0ms to $N\tau_{\text{dendrite}}$ ms, where N=3 is the number of consecutive 173 segments. We believe this source of timing-invariance to be a highly beneficial feature for generalization that helps 174 explain phenomena, where the same sequence of events must be detected across multiple timescales.

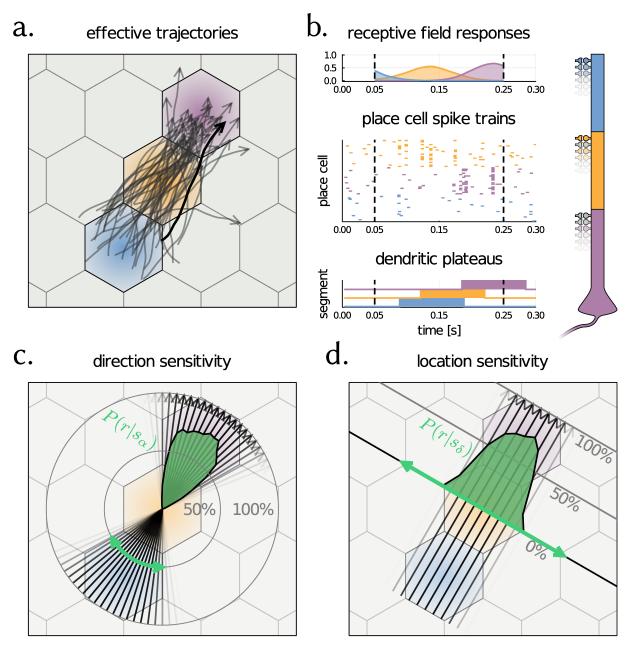


Figure 2: A simple neuron with three dendritic segments arranged as shown to the right of panel b can detect directed paths on a timescale of 300ms. **a.** The receptive fields of place cell populations tile the environment through which the animal moves in a hexagonal grid. Random trajectories are generated through a stochastic process with randomized initial positions, velocities and angular heading to simulate the animal's movements. **b.** While the animal follows the black trajectory through space, the response of the place cell populations' tuning curves show the sequential activation of the populations over time (top panel). The stimulus s are generated spikes (middle panel) that lead to a temporal sequence of dendritic plateaus (bottom panel) and results in a somatic spike as the response r. **c. and d.** The neuron responds with high probability to exactly those paths that traverse the desired receptive fields in the correct direction and with little lateral offset. In the experiment, a change of rotation leads to paths s_{α} (**c** black) or a shift orthogonal to the movement direction leads to paths s_{δ} (**d** black) as indicated by the green arrows. The empirical probability of firing responses $P(r|s_{\alpha})$ and $P(r|s_{\delta})$ respectively are shown in superimposed density plots in green in polar (**c**) and Cartesian (**d**) coordinates and show a highly specific pattern detector.

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6 Plateaus integrate evidence on long timescales

In the previous example, specific paths are recognized by memorizing the sequential activation of different neural populations on a slow behavioural time-scale. A seemingly different, yet in fact closely related problem is the integration of individually unreliable bits of evidence over time. Consider, for example, a population of neurons that extract some relevant feature of a stimulus, such as the local movement direction in a visual moving dots stimulus. If we assume a retinotopic mapping, neighbouring neurons are highly correlated, and whenever the local movement direction is apparent, we expect a couple of neighbouring neurons coding for that direction to produce a volley of spikes. However, these events are unlikely to occur at the exact same point in time throughout the entire input space. The decision, whether or not the visual flow is in a certain direction, therefore requires that a neuron can integrate many such pieces of evidence, each indicated by a spike volley event, over a longer time-scale. Despite the all-or-none response of dendritic plateaus, a neuron with sufficiently many dendritic segments can in fact approximate such a smooth integration of evidence on timescales of hundreds of milliseconds!

We give an example of evidence integration using dendritic plateau potentials in a simplified experiment, in which a neuron with 1000 dendritic compartments receives input from a population of 1000 input neurons through a total of 20,000 stochastic synapses (**Fig. 3**). The weak signal to be integrated by the ADSP neuron is encoded into spike volleys of 10 simultaneous spikes from adjacent neurons of the input population. Each dendritic segment of the ADSP neuron is connected to a different set of 20 adjacent neurons in the input population, and a total of 300 dendritic segments are required be in simultaneous plateau states for the neuron to emit a somatic spike.

Because each spike volley is likely to activate a different dendritic segment, we expect the number of simultaneously active dendritic compartments to reflect the average rate of incoming spike volleys during a time-interval of one plateau duration. This corresponds to a filtering of the time-varying rate by a rectangular filter, and, for a brief interval after stimulus onset, represents an ideal integrator. We observe this exact behavior by driving the rate, at which spike volleys are generated by the input population, to three different levels for brief time-intervals (**Fig. 3b**, orange line). The number of co-activated dendritic segments (blue line) closely follows the theoretical prediction of an ideal rectangular filter (black dashed line) until saturation. In particular, during the rising flanks right after stimulus onset (**Fig. 3c, d and e.**), we see the number of co-active segments rise with a slope proportional to the intensity of the stimulus until it saturates after 100ms. The neuron begins firing spikes once sufficiently many segments are active (red line). This is exactly the behavior expected for evidence integration: The ADSP neuron will fire sooner if the amount of evidence encoded in the stimulus is stronger, and will not fire at all if it remains sub-critical.

Interestingly, the stochasticity of synaptic transmission helps to further decorrelate the partially overlapping input to different dendritic segments, and can regulate the total amount of evidence required to reach the neuron's physiologically fixed spiking threshold. Also, while the example here makes use of just a single "layer" of dendritic segments directly driving the soma, this idea can be extended to deeper chains of multiple segments, such as in the previous example, to allow for the integration of evidence and non-linear combination thereof on timescales even longer than one plateau duration.

Dendritic morphology determines computational function

In the two previous examples, we assume that each dendritic segment is driven by well-timed volleys of coincident spikes, the magnitudes of which represent the magnitude of an underlying signal. But in theoretical neuroscience, the function of a neuron is often analyzed in a rate-based framework, which relates only the average firing rate of a neuron to the average firing rates of its spiking inputs.

Applying this sort of analysis to our proposed neuron model reveals, how different morphologies of dendritic arbors give rise to different non-linear computations. A dendritic segment driven by independent Poisson spike-trains originating from some population A of 25 neurons respond by triggering plateau potentials at a rate $\varrho(r_A)$ that continuously depend on the fixed firing-rate r_A of the populations' neurons. Here, 8 coincident spikes are required to trigger a plateau. As each plateau lasts for $100 \, \mathrm{ms}$, ϱ saturates at a rate of $10 \, \mathrm{plateau}$ specification by an identical but independent population of neurons, we analyze the relative contributions of the populations B and C in the same way. In these experiments, we hold the firing rate $r_A = 25 \, \mathrm{constant}$. For a neuron $C \to_1 \, B \to_1 \, A$, whose segments are sequentially chained together, a spike is generated if and only if both C and B are activated, and in the correct order. The resulting contour-plot, which shows how the output firing rate of this neuron scales with both r_C and r_B , illustrates that both a high firing rate of population C and B are required to result in a high firing rate of the neuron (Fig. 4d). This is similar to the neuron with two parallel segments $(C + B) \to_2 A$ (Fig. 4e), only that simultaneous activation of both segments, not sequential activation, is required. The shape of this function closely matches an idealized "and" operation (Fig. 4b), the firing rate of which can be derived as just the product of the rates at which plateaus are triggered in all dendritic segments:

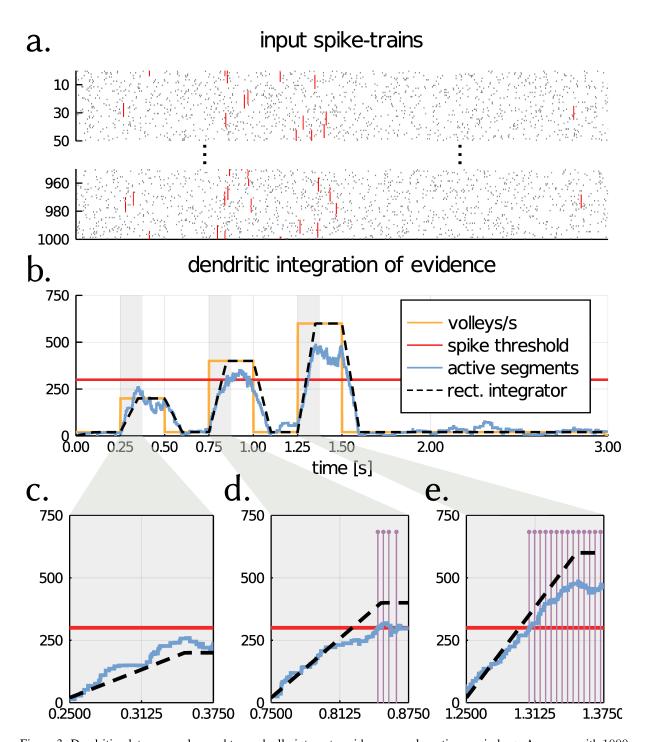


Figure 3: Dendritic plateaus can be used to gradually integrate evidence over long time periods. **a.** A neuron with 1000 dendritic segments is driven by 1000 incoming spike-trains. Embedded in these spike-trains are spike volleys of 10 coincident spikes each, spread across 10 neighbouring neurons (shown in red). **b.** The rate of spike volleys is determined by an input signal (organge line). Each segment receives input from 20 consecutive neurons through stochastic synapses with transmission probability p=0.5, and requires 5 coincident spikes to trigger a plateau potential. The total number of co-activated dendritic segments (blue line) follows the convolution of the stimulus signal with a rectangular filter of length 100ms (black dashed line). **c-e.** For increasing levels of stimulation, the number of co-activated segments rises faster and saturates at a higher level, crossing the threshold required for spike initiation (horizontal red line) at an earlier point in time or not at all, resulting in a sequence of spikes (vertical purple lines).

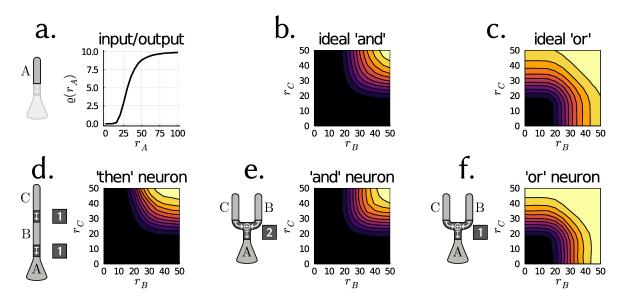


Figure 4: A rate-based analysis reveals well-known computational primitives. **a.** A single dendritic compartment that receives independent Poisson-spike trains at a fixed rate r_A from a population of 25 neurons responds with plateaus at a rate that can be expressed as a non-linear sigmoidal function $\varrho(r_A)$. For multiple dendritic segments, each of which receives input from an identical but independent population A,B or C, the neuron's computation depends on the dendritic morphology. **d and e.** If both segments C and B are required to enable a somatic spike, the neuron's firing rate is proportional an **b** idealized "and" operation between the two inputs. **f.** If either of the two segments suffices, **c** the firing rate instead resembles an idealized "or" operation.

$$\varrho(A,B,C) \propto \tau_{\text{dendrite}} \varrho(r_A) f_{\text{and}}(B,C)$$
 where $f_{\text{and}}(B,C) = \tau_{\text{dendrite}}^2 \varrho(r_C) \varrho(r_B)$

Here, $\varrho(A, B, C)$ is the firing rate of the neuron, and $f_{\rm and}(B, C)$ is the factor due to the segments B and C.

For a different dendritic morphology $(C+B) \to_1 A$, where a plateau in either segment C or B is sufficient (**Fig. 4f**), we see a response that closely resembles an idealized "or" operation (**Fig. 4c**):³

$$f_{\rm or}(B,C) \propto \tau_{\rm dendrite} \rho(C) + \tau_{\rm dendrite} \rho(B) - f_{\rm and}(B,C)$$

For a derivation of $f_{\rm and}$ and $f_{\rm or}$ see the Methods section. This rate-based functional description offers a very useful abstraction of the neurons' behaviours, but it necessarily neglects questions of timing. As we saw in the previous sections, depending on the morphology, a dendritic arbor can impose stringent requirements on the order in which different segments can be activated. For example, while both neurons $C \to_1 B \to_1 A$ and $(C+B) \to_2 A$ require strong input from both input population B and B and hence show the same "and"-like response in the rate-coding paradigm, the former imposes the constraint that the input from population B must arrive before that from population B while the latter does not. Rather than an "and"-like operation, neuron B and B in fact implemented an "and then" operation. This is apparent when looking at the joint probability density of the relative timing of dendritic plateaus in the respective segments directly preceding a somatic spike (Fig. 5). In particular, if we only consider the unambiguous cases of one dendritic plateau each occurring in each segment within a brief window before a somatic spike (shown by the white dots (Fig. 5)), we observe that for neuron B and B and a dendritic plateau in segment B can occur at most 100ms before the somatic spike and is preceded by a dendritic plateau in segment B by at most another 100ms for a maximum total delay of 200ms. In contrast for neuron B and B and plateau in either segment within a 100ms window suffices to trigger a somatic spike. For neuron B and B and plateau in either segment within a 100ms window suffices to trigger a somatic spike.

³ As the last equation shows, referring to this operation as an "or" is justified in the sense that the resulting rate is proportional to the addition of the segments' individual plateau-firing-rates minus the "and" operation applied to both, which generalizes the Boolean operation to real values.

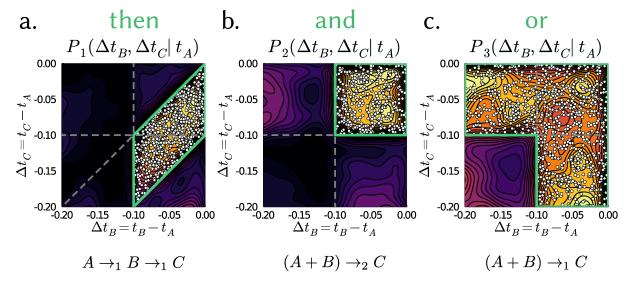


Figure 5: Dendritic morphology imposes timing constraints not revealed by rate-based analysis. For the neurons shown in figure 4, the joint probability distribution of relative timings $\Delta t_B, \Delta t_C$ of dendritic plateaus directly preceding a somatic spike at t_A show a distinct temporal structure (contour-plots). **a.** For the "then" neuron, a plateau in segment C must precede a plateau in segment C by at most 100ms, which in turn must occur at most 100ms before a somatic spike can be triggered. This is evident by the fact that all unambiguous cases, where exactly one plateau in each segment C and C and C was observed before a somatic spike, fall into the corresponding parallelogram-shaped domain (white dots). **b.** The "and" neuron shows a similar rate-response to the "then" neuron, but requires both inputs to occur within 100ms before the somatic spike. **c.** The "or" neuron only requires either of the populations C to trigger a plateau within 100ms before a somatic spike.

Discussion

In this theoretical study we showed how a well-known biological phenomenon, dendritic plateau potentials, can drastically improve the computational capabilities of spiking neurons, turning them into powerful spatio-temporal pattern detectors. Due to the long-lasting memory provided by these plateau potentials, it becomes possible for individual neurons to integrate evidence or distinguish specific sequences of input on a timescale of hundreds of milliseconds – an order of magnitude larger than commonly observed membrane time constants [42]. In our model, the morphology of a neural dendrite determines its computational function and, when viewed in a conventional rate-coding paradigm, allows an individual neuron to implement a wide range of nonlinear behaviours in a modular and intuitive way.

This is in line with the two-layer neuron model proposed in [43], which used a detailed biophysical simulation of a pyramidal neuron to investigate the nonlinear effect on the neuron's firing rate due to synaptic input at different dendritic branches. Using a diverse array of stimuli, they showed that a two-layer network of sigmoidal subunits provides a substantially better approximation of the neuron's firing rate than a linear point-neuron. They speculated, however, that the prediction could be improved further, if the nonlinear interactions between the branches were considered, which we did here. We also investigated the use of dendritic plateau potentials as long-lasting memory traces, which our results revealed to be particularly important for evidence integration and the detection of temporal sequences. Remarkably, our drastically simplified and inherently event-based model could qualitatively reproduce properties of the model in [43], such as the sigmoidal input-output firing rate response of each dendritic segment and the linear-nonlinear combination thereof at the soma (see methods section).

But on the fast time-scale of individual spikes, our model differs substantially from this and other rate-based point-neuron models, since it relies on the detection of volleys of coincident spikes on a millisecond time-scale as the basic units of information, which are then integrated on the slower time-scale of dendritic plateau potentials. Our model is more closely related to recent work by [44], which proposed the use of active coincidence detection in dendritic segments to model prolonged effects of basal dendrites on the soma. A similar line of reasoning can also be found in [45], which presented a very elegant two-compartment neuron model and corresponding learning rule with one somatic and one dendritic compartment. Both models assign a specific functional role to the (basal) dendrite segments, namely to predict subsequent activation at the soma from their local synaptic inputs, which allows individual neurons to learn to predict state-transitions ("prospective coding"). Longer sequences are then detected by networks of such

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laterally connected neurons, endowing the networks with a form of temporal sequence-memory ("hierarchical temporal memory"). In our work, we have focused on a more mechanistic model that heavily relies on biological phenomena observed in single neurons. This allowed us to describe a neuron's computational capability concretely as that of a sophisticated pattern detector with long-lasting memory, and to illustrate how these mechanisms at play would appear under a rate based analysis. We believe our results offer a very appealing explanation of spike-based computation that has wider implications in neuroscience and raises several important questions, which we briefly discuss in the following:

What is the role of inhibition for dendritic computation?

Our model only takes into account excitatory synapses, but has clear implications for the role of inhibition. The 283 all-or-none response of dendritic plateau potentials in our model implies that the only significant effect an inhibitory 284 synapse can have on the far-away soma is by either reducing the likelihood of plateaus, preventing the generation of 285 plateaus altogether, or by disrupting already ongoing plateau potentials. In the first two cases, an inhibitory synapse's 286 post-synaptic potential must be either well-timed to coincide with the volley of excitatory spikes or exhibit a longer 287 time-scale. Experiments suggest that inhibition can affect the ability of dendrites to generate active plateaus and prevent them [46]. The disruption of ongoing plateaus has also been reported and analyzed [47] and requires no such precise 289 timing a-priori, as long as the spike occurs within the plateau's duration. Inhibition may, however, exhibit different 290 effects depending on when during the plateau processes it is received. In all cases, the likely effect is shunting, rather 291 than substractive, inhibition.

Shunting inhibition can provide an efficient mechanism to improve the computational capabilities of the neurons described above, for example as it would allow individual neurons to exclusively respond to a sequence $a \to b$ but not to the sequence $a \to b \to c$, which is impossible for a neuron with purely excitatory synapses. Inhibition may therefore play an important and distinct role in ADSP neuron that warrants further investigation.

297 What are the implications of this model for plasticity?

We discussed a fundamental mechanism of dendritic computation and its capabilities, but did not cover the important topic of learning and plasticity. Nevertheless, the model presented here imposes constraints on potential plasticity mechanisms. Due to the long-lasting plateau potentials, a synaptic input can have a relevant causal effect for a somatic spike at a much later time. This makes the temporal assignment of credit for spiking outputs to synaptic inputs fundamentally difficult. The timing-invariance shown by our model and the dependency on the complex nonlinear dynamics within a dendritic tree further exacerbate this problem.

The most prominent example of synaptic learning is spike-time dependent plasticity [48], which tunes synaptic efficacy based on the relative timing of pre- and post-synaptic activity. Since the active dendritic processes discussed here both dominate the post-synaptic membrane potential as well as local Ca^{2+} concentration, they have a major effect on Hebbian plasticity [49, 50].

This is at odds with the common assumption, that backpropagating action potentials (bAPs) from the soma into the dendrite act as the primary post synaptic signal driving synaptic plasticity [51]. Since dendritic plateau potentials strongly depolarize dendrite segments for an extended period of time and should similarly "backpropagate" throughout the dendritic tree, it seems unlikely to us that bAPs are the primary factor for synaptic plasticity in neurons with active dendritic processes. Resolving this inconsistency is an important, but open research question.

Additionally, our model is based on binary stochastic synapses, and which segment the synapse terminates on plays a more important role than its efficacy. We therefore believe that structural plasticity mechanisms are particularly relevant for this kind of model. Furthermore, homeostatic plasticity mechanisms, e.g. scaling synaptic transmission probabilities[52], are in our view important to ensure that only sufficiently large spike-volleys, but not randomly correlated inputs, can reliably trigger plateau potentials.

318 Is neuronal computation based on plateau processes?

Dendritic processes are thought to implement solutions to a number of specific computational problems in neurons [53], often distributed across many functional dendritic compartments [54, 55]. Based on convincing biological evidence for the mechanism of plateau generation and the interaction of such plateaus, we have argued that they are indeed the primary building block for the implementation of behaviorally highly relevant computations. How can this claim be experimentally verified or falsified?

Direct experimental verification, that computation in single neurons is well described by our proposed ADSP neuron model requires simultaneous measurement of synaptic inputs and local membrane potentials along a single neuron's dendrite on a fine temporal and spatial resolution over a long-time span.

As a first step, since our model is driven by incoming spike volleys from multiple intact neuron populations, *in vivo* measurements could verify the existence of patterns of spike-volleys over different timescales using newly developed

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statistical techniques [56, 57]. 329

that achieves this without reliance on active dendritic processes. This may be the case either for neurons incapable of generating plateaus in the first place, or if plateau-generating processes have been pharmacologically disabled. 333 Thirdly, we predict single neurons that use active dendritic sequence processing to have spatio-temporal receptive 334 fields on long temporal timescales, but with high tolerance to variations in the precise timing of individual plateaus, 335 qualitatively described in Fig. 2. Because of this invariance, we propose to go beyond linear analysis such as spike-336 triggered averages and instead measure both somatic response, as well as the timing of plateaus across the dendritic 337 tree to find structures in the joint distributions as demonstrated in Fig. 5. Experimentally, spatio-temporal receptive 338 fields of this kind could also be found by systematically varying stimuli, and should disappear when plateau-generating 339 processes are disrupted.

Secondly, a key part of the model, the detection and integration of information across two timescales, one on the order

of a few milliseconds, the other on the order of a hundred milliseconds or more, can be refuted for any type of neuron

While we have based our analysis on NMDAr-mediated plateaus in pyramidal cells [20], the same computational 341 principle may be found in other neuron types, as well. For example, Purkinje cells in the cerebellum also generate 342 localized Ca²⁺ events in response to coincident input on individual dendritic segments [58, 59], and thalamo-cortical 343 neurons respond to strong synaptic input by localized plateaus in distal dendritic branches [60]. This indicates that the 344 underlying ADSP mechanism, possibly implemented through diverse means in a case of convergent evolution, may be 345 very general and ubiquitous in the brain. 346

In summary, we have presented and analyzed an intentionally simple model of neural computation based solely on the interaction of coincident spikes and dendritic plateau potentials. This revealed, how the morphology of the dendritic tree can implement and compose a wide range of non-linear computational functions. Two key features of this computational mechanism are its invariance to exact timings of inputs and its ability to operate on timescales much longer than post-synaptic potentials. We have highlighted the importance of the combination of these two features in two behaviorally relevant tasks: the detection of sequences and the integration of weak signals on long timescales. However, when analyzed from the usual perspective of rate-coding, these computational properties are hard to detect. To this end, we have therefore an alternative set of analyses to identify whether computation in single neurons is indeed based on dendritic plateaus.

Methods 356

Formal description of the event-based framework for computation in active dendrites 357

Mathematically, we approximate both EPSPs and plateau potentials by rectangular pulses with fixed duration τ_{synapse} 358 and τ_{dendrite} , respectively. Here, we chose $\tau_{\text{synapse}} = 5 \text{ms}$ and $\tau_{\text{dendrite}} = 100 \text{ms}$ for all experiments if not stated otherwise. 359 The dynamics of each dendritic segment can then be fully described in terms of the arrival times of incoming spikes 360 as well as the times at which plateau potentials are initiated within the segment itself or in other directly connected 361 segments. For some segment i, the synaptic input X_i and the dendritic input Y_i take the form of equations (1) and (2), 362 respectively:

$$X_i(t) = \sum_{j \in S_i} \sum_k \chi_{i,j,k} \cdot \mathbf{1}_{[s_k^j, s_k^j + \tau_{\text{synapse}}]}(t) \quad \text{where } \chi_{i,j,k} \sim \text{Bernoulli}(\omega_{i,j})$$
 (1)

$$X_{i}(t) = \sum_{j \in S_{i}} \sum_{k} \chi_{i,j,k} \cdot \mathbf{1}_{[s_{k}^{j}, s_{k}^{j} + \tau_{\text{synapse}}]}(t) \quad \text{where } \chi_{i,j,k} \sim \text{Bernoulli}(\omega_{i,j})$$

$$Y_{i}(t) = \sum_{j \in D_{i}} \sum_{k} \mathbf{1}_{[t_{k}^{j}, t_{k}^{j} + \tau_{\text{dendrite}}]}(t)$$

$$(2)$$

$$t_{m+1}^{i} = \min\left\{t \in \mathbb{R} \mid t \geq t_{m}^{i} + \tau_{\text{dendrite}}, X_{i}(t) \geq \theta_{i}^{\text{syn}} \text{ and } Y_{i}(t) \geq \theta_{i}^{\text{den}}\right\},\tag{3}$$

where $\mathbf{1}_{[a,b]}$ represents a unit pulse during the time interval [a,b], and s_k^j and t_j^i are the times of spikes arriving from some presynaptic neuron j and the plateau onset times on segment i, respectively. The random variable $\chi_{i,j,k}$ represents the independent probabilistic transmission of every spike k from source j via a synapse to dendritic segment i, where the transmission occurs with the synapse specific probability $\omega_{i,j}$. The sets S_i and D_i respectively identify the segment's synaptic connections to other neurons and which other dendritic segments it is directly coupled to, and therefore reflect the morphology of the neuron's dendritic tree. Equation (3) states that, if the segment is not in a plateau state already, a new plateau is initiated as soon as both synaptic and dendritic inputs exceed their respective thresholds θ_i^{syn} and θ_i^{den} .

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Implementation of the navigation experiments

To simulate the stochastic movements of a rat, random paths are generated with time-varying location l(t) = $(X(t),Y(t)) \in \mathbb{R}^2$ as solutions of the following system of stochastic differential equations:

$$\begin{split} dX &= \cos(2\pi A)Vdt\\ dY &= \sin(2\pi A)Vdt\\ dA &= 0.25dW_A\\ dV &= 10.0(0.25-V)dt + 0.1dW_V \end{split}$$

A represents the angular heading of the animal, V represents its velocity in $\frac{m}{s}$ and W_A , W_V represent independent standard Brownian motion processes. Each path is generated with a randomized initial position within a rectangular domain of $10cm \times 9.5cm$, a random angular heading and a random velocity according to the marginal stationary distribution of V in the equation above, and is simulated for a fixed duration of 200ms. Three populations of place cells, each 20 neurons strong, are centered on a hexagonal grid with center-to-center distance of $r \approx 2.9cm$. Each population randomly emits spike volleys following a homogeneous Poisson process with rate $\lambda = 50Hz$. The magnitude of each spike volley is determined by the population's mean activity at the time, which depends on the animal's location within the environment through a receptive field tuning curve. The tuning curves model the probability of each individual neuron within the population to participate in a given spike volley by the bell-curves $f_i(x) = \exp(-\frac{x-\mu_i}{2\sigma^2})$ with coefficient $\sigma = 9.7mm$, centered on the tiles of the hexagonal grid. The total number of spikes emitted during a 383 volley from population i at time t is therefore a random variable distributed according to a Binomial distribution with population size n=20 and probability $p=f_i(l(t))$. Additionally, each neuron in the population emits random spikes at a rate of 5Hz to emulate background activity. Each spike is transmitted through stochastic synapses independently with probability 0.5.

Each of the simulated neuron's dendritic segments receives spiking input from the 20 neurons of one population and 388 requires at least 5 coincident spikes to trigger a plateau potential. The three segments are connected in a chain that 389 requires sequential activation by spike volleys from the input populations in correct order to fire a spike. A random 390 path is considered to be accepted by the neuron, if the neuron responds with a spike at any point in time during the 391 corresponding simulation run. 392

To evaluate the rotation and location sensitivity of the neuron, we also generate straight paths with constant movement 393 speed $v=\frac{3r}{200 \mathrm{ms}} \approx 43 \mathrm{cm/s}$ that are either rotated around the center of the environment by an angle α or offset from the center by a distance Δx orthogonal to the optimal movement direction. For each angle or offset, respectively, the 395 empirical firing probability of the neuron in response to that path is estimated by simulating the path and the neuron's 396 responses 500 times each. 397

Implementation of the evidence-integration experiments 398

The input to the evidence-integrating neuron is generated by superimposing spike volleys onto 1000 independent Poisson processes with a constant firing rate of 10Hz. The volleys times are generated by a Poisson process with a time-varying rate $\lambda(t)$ representing the incoming "evidence". Here, $\lambda(t) = 200 \text{Hz} \cdot (\mathbf{1}_{[0.25,0.5]}(t) + 2 \cdot \mathbf{1}_{[0.75,1.0]}(t) + 2 \cdot \mathbf{1}_{[0.75,1.0]}(t)$ $3 \cdot \mathbf{1}_{[1,25,1.5]}(t)) + 20$ Hz. Each volley consists of simultaneous spikes from a randomly chosen set of ten input neurons with consecutive indices (wrapping around from 1000 to 1). Since each EPSP is assumed to last for a duration of 5ms, volleys and individual spikes are discarded if they occur less than 5ms after a preceding volley or spike. Each of the neuron's 1000 dendritic segments receives synaptic input via stochastic synapses with transmission probability 0.5 from 20 consecutive input neurons. As the number of input neurons and dendritic segments matches in this example, there is exactly one dendritic segment for every group of 20 consecutive input neurons, and each input neuron projects to exactly 20 dendritic segments. The total number of the neuron's synapses in this example is therefore 20000. Over time, the number of simultaneously active dendritic compartments as well as the times of generated somatic spikes is recorded. As a reference, the convolution $(\lambda \star \Pi)(t)$ of the time-varying rate-function λ with a rectangular filter Π of length 100ms and unit-integral is calculated.

Implementation of the rate-based analysis

For the rate-based analysis, four different neurons are constructed. First, a neuron consisting of a single dendritic compartment is driven by a total of 25 independent Poisson spike-trains with constant firing rate r_A . As in all 414

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other experiments, the duration of each spike is set to $au_{synapse} = 5 \mathrm{ms}$, the duration of a plateau potential is set to $au_{dendrite} = 100 \mathrm{ms}$. By systematically varying r_A and, for each choice, recording the number of plateau potentials generated during a simulation time-interval of $250 \mathrm{s}$ we can estimate the smooth function $\varrho(r_A)$, which relates the firing rate of the input population A to the resulting rate at which plateau potentials are generated.

For each of the three morphologies representing the $C \to_1 B \to_1 A$ neuron, the $(C+B) \to_2 A$ neuron and the $(C+B) \to_1 A$ neuron, we systematically vary the input firing rates of both populations B and C independently while keeping the firing rate of population A fixed at a constant 25Hz. For each combination, we again record the number of somatic spikes generated over a time-interval of 250s. As a reference for these two-dimensional functions, we use an idealized "and" and "or" function defined as:

$$f_{\text{and}}(B,C) = \tau_{\text{dendrite}}^2 \varrho(r_C) \varrho(r_B) \tag{4}$$

$$f_{\text{or}}(B,C) = \tau_{\text{dendrite}}\varrho(C) + \tau_{\text{dendrite}}\varrho(B) - f_{\text{and}}(B,C)$$
(5)

$$= 1 - (1 - \tau_{\text{dendrite}} \varrho(C))(1 - \tau_{\text{dendrite}} \varrho(B)) \tag{6}$$

At a firing rate r_X , a segment driven by population X is in a plateau state at a given point in time with probability $\tau_{\text{dendrite}}\varrho(r_X)$, therefore the probability that a segment driven by population C is active at the time that an input from population B arrives, which could in turn activate the next segment, is $\tau_{\text{dendrite}}\varrho(r_C)$. The probability that this second segment is still active, when yet another volley from population A arrives to possibly trigger a somatic spike is also $\tau_{\text{dendrite}}\varrho(r_B)$. Therefore the neuron's firing rate is proportional to $\tau_{\text{dendrite}}^2\varrho(r_C)\varrho(r_B)$. Similarly, the probability that two parallel upstream segments driven by populations C and B are simultaneously active at a given point in time is $\tau_{\text{dendrite}}^2\varrho(r_C)\varrho(r_B)$. In contrast, the probability that either upstream segment is active at a given point in time is just the probability that not both are simultaneously inactive, i.e. $1-(1-\tau_{\text{dendrite}}\varrho(C))(1-\tau_{\text{dendrite}}\varrho(B))$. This expression has the nice alternative form c+b-cb, where $c=\tau_{\text{dendrite}}\varrho(C)$, $b=\tau_{\text{dendrite}}\varrho(B)$ and $cb=f_{\text{and}}(B,C)$, which generalizes the Boolean "or" operation to real-valued firing rates. When identifying true with 1 and false with 0, the truth-table of this expressions matches that of the logic expression "c or b".

To evaluate timing requirements for each of these three neuron morphologies, we run another simulation at constant input rates $r_A = r_B = r_C = 25 \mathrm{Hz}$ for a duration of 1h of simulated time. We record the time of each plateau-initiation-event in both upstream segments driven by population C and B for a time-interval of 200ms preceding each somatic spike. If there is exactly one plateau-event from each segment in such a time-interval, we record this as an *unambiguous* pair of plateau events. If there is more than one plateau-event on either of the dendritic segments, we record all pairs of plateau-events in that time-interval composed of one plateau event for each segment. We refer to these latter pairs as *ambiguous*. Using these ambiguous pairs, we estimate the joint probability distribution $P_i(\Delta t_B, \Delta t_C|t_A)$ over relative times Δt_B and Δt_C between a plateau triggered by population B or C and a somatic spike triggered at time t_A by population A. For a more reliable estimate of the timing constraints, we consider only the unambiguous pairs, which evidently fall into distinct domains of these joint probability distributions that uniquely characterize the precise timing requirements of the respective neuron morphologies. This can be seen in figure 5. E.g. for the $C \to_1 B \to_1 A$ neuron, all plateaus triggered by population C must precede those triggered by B, but cannot precede them by more than one plateau duration of 100ms, therefore they fall into a parallelogram below the diagonal. For the $(C + B) \to_2 A$ neuron, on the other hand, both plateau events must independently occur within 100ms before a somatic spike, and hence fall into the upper quadrant of the joint density.

450 Code availability

All simulations are implemented in a custom developed package in the Julia programming language [61], publicly available via the code repository hosted at https://github.com/jleugeri/ADSP.jl. Further documentation of the simulator and implementation details can be found there.

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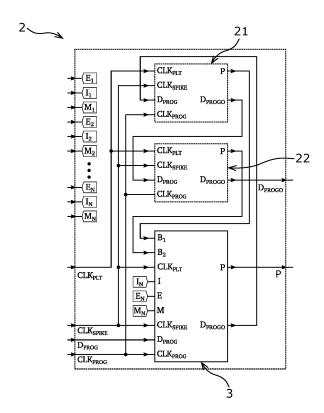
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Prüfungsantrag gemäß § 44 PatG ist gestellt.

Die folgenden Angaben sind den vom Anmelder eingereichten Unterlagen entnommen.

(54) Bezeichnung: Neuromorpher Musterdetektor und neuromorphe Schaltkreisanordnung hiermit

(57) Zusammenfassung: Die vorliegende Erfindung betrifft einen neuromorphen Musterdetektor (2), welcher ausgebildet ist, wenigstens zwei 1-Bit Eingangssignale (E_1 - E_N) eines zu erkennenden Musters zu erhalten, mit wenigstens zwei Vergleichsschaltungen (3), welche jeweils ausgebildet sind, eines der 1-Bit Eingangssignale (E_1 - E_N) zu erhalten, die Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen 1-Bit Eingangssignals (E_1 - E_N) innerhalb eines vorbestimmten Zeitraums zu zählen, die Anzahl der gezählten Zustände mit einem vorbestimmten Schwellwert der jeweiligen Vergleichsschaltung (3) zu vergleichen und bei Überschreiten des Schwellwerts auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen.



Beschreibung

[0001] Die vorliegende Erfindung betrifft einen neuromorphen Musterdetektor gemäß dem Patentanspruch 1 sowie eine neuromorphe Schaltkreisanordnung gemäß dem Patentanspruch 13.

[0002] Zur Verarbeitung ihrer Informationen können analog erfasste Signale, welche z.B. sensorisch erfasste Informationen repräsentieren können, in digitale Signale gewandelt und dann verarbeitet werden. Die Erfassung der analogen Signale kann üblicherweise mittels elektrischer Spannung erfolgen, welche einen zeit- und wertkontinuierlichen Verlauf, d.h. eine durchgängigen Verlauf der elektrischen Spannung über der Zeit, aufweist. Ein derartiges elektrisches Analogsignal kann mittels eines Analog-Digital-Umsetzers in ein digitales Signal in Form eines zeit- und wertdiskreten Verlaufs gewandelt werden, um die Information der digitalen Signalverarbeitung zugänglich zu machen. Ein derartiges digitales Signal kann auch als binäres Signal bezeichnet werden und zwei unterschiedliche Zustände in Form von unterschiedlich hohen elektrischen Spannungspegeln aufweisen, so dass über die Länge bzw. Dauer des Signalverlaufs zwischen niedrigen und hohen Spannungspegeln unterschieden werden kann. Die niedrigen Spannungspegel können als "low"-Zustände und die hohen Spannungspegel als "high"-Zustände bezeichnet werden. Hierdurch können die Zustände "0" und "1" dargestellt werden.

[0003] Die digitale Signalverarbeitung mittels entsprechender elektronischer Bauelemente wie z.B. digitale Signalprozessoren und Mikroprozessoren bietet dabei Vorteile und Möglichkeiten, welche mit analog arbeitender Elektronik gar nicht oder lediglich mit hohem Aufwand umsetzbar wären. Dabei werden die digitalen Signale üblicherweise nicht als die binären Signale eines Verlaufs von Nullen und Einsen in Form von niedrigen und hohen Spannungszuständen verarbeitet sondern als Werte etc. z.B. in Folgen von acht Bits, auch Byte genannt, dargestellt, gespeichert und durch Software verarbeitet. Die entsprechenden Algorithmen, welche die Verarbeitung der digitalen Signale durchführen, werden hierzu als Programmcode einer geeigneten Programmiersprache umgesetzt und z.B. auf einem Mikroprozessor oder auf einer CPU (Central Processing Unit) als serielle Abfolge der Programmierschritte ausgeführt. Mit anderen Worten werden in der digitalen Signalverarbeitung üblicherweise die Instruktionen numerischer Algorithmen von Prozessorarchitekturen auf Binärzahlen implementieren, was die sequentielle Abarbeitung in arithmetisch-logischen Einheiten (ALU) und die Verwendung einer Speichereinheit bedingt.

[0004] Zur Verarbeitung digitaler sowie analoger Informationen in Form digitaler Signale können auch sog. künstliche neuronale Netze bzw. Netzwerke ver-

wendet werden, bei denen mittels künstlicher Neuronen die Funktionsweisen biologischer Neuronen bzw. biologischer neuronaler Netze bzw. Netzwerke nachgebildet werden. Die einzelnen künstlichen Neuronen arbeiten dabei zeitlich parallel zueinander, vergleichbar der Vorbilder der biologischen Neuronen. Da eine derartige Arbeitsweise mit den sequentiell arbeitenden Prozessoren strukturbedingt jedoch nicht möglich ist, kann die parallele Arbeitsweise der künstlichen Neuronen auch bei Verwendung mehrerer paralleler Prozessoren bzw. Prozessorkerne nur unzureichend implementiert werden. Dies erschwert die tatsächliche Implementierung parallel arbeitender Verfahren zur digitalen Signalverarbeitung mit künstlichen neuronalen Netzen.

[0005] Zur Implementierung von künstlichen neuronalen Netzen bzw. Netzwerken werden daher auch neuromorphe Schaltkreise verwendet, welche jeweils ein biologisches Neuron als elektronische Schaltung abbilden und durch ihr Zusammenwirken das künstliche neuronale Netz bzw. Netzwerk ergeben. Die einzelnen neuromorphen Schaltkreise können dabei tatsächlich parallel zueinander arbeiten und hierdurch die Signalverarbeitung beschleunigen bzw. die als Vorbild dienenden biologischen Neuronen besser nachbilden.

[0006] Typischerweise wird das Verhalten des einzelnen künstlichen Neurons dynamischen Systemen aus den theoretischen Neurowissenschaften wie z.B. dem Leaky-Integrate-and-Fire-Modell nachempfunden, durch digitale Arithmetik approximiert und der Datenaustausch zwischen den künstlichen Neuronen durch die Übertragung von Paketen realisiert. Dies erfordert jedoch den Einsatz vieler Recheneinheiten bzw. vieler arithmetisch-logischer Einheiten und stellt hohe Anforderungen an das Paket-Routing zwischen den einzelnen Recheneinheiten. Im speziellen Bereich der Spiking-Neuromorphic-Hardware wird dabei pro künstlichen Neuron und pro Zeitschritt lediglich ein binäres Signal erzeugt.

[0007] In einem parallel signalverarbeitenden neuromorphen Netz bzw. Netzwerk der Digitaltechnik sollten somit folgende technische Probleme gelöst bzw. folgende technische Eigenschaften realisiert werden:

- Künstliche Neurone sollte Eingangssignale von vielen anderen künstlichen Neuronen integrieren können. Dies erfordert einen Mechanismus, um Eingangssignale aufzuaddieren und mit einem kritischen Grenzwert vergleichen zu können. In bestehenden Ansätzen der digitalen Signalverarbeitung wird dies mittels ALUs durch Ganzzahlarithmetik realisiert.
- Das Einsatzgebiet von digitaler Signalverarbeitung ist häufig durch das Erfordernis der Echtzeitfähigkeit ausgezeichnet, d.h. durch die Fähigkeit des Betriebssystems der Recheneinheit

bzw. der Recheneinheiten, digitale Signale innerhalb einer vorbestimmbaren Frist sicher verarbeiten zu können. Die Einhaltung einer Reaktion auf das digitale Signal innerhalb dieser Frist muss in diesem Fall sichergestellt sein.

So sind die Zeitskalen, auf welche ein analoges Signal in der Außenwelt relevante und zu verarbeitende Charakteristika aufweist, nicht fest und zum Teil auf schnellen oder langsamen Skalen variiert. Daher müssen die Zeitskalen der Verarbeitung digitaler Signal im integrierten Schaltkreis von denen in der Außenwelt entkoppelt werden. Klassische Ansätze der digitalen Signalverarbeitung in z.B. Mikrokontrollern umgehen dieses Problem, indem Zwischenergebnisse im dedizierten Arbeitsspeicher abgelegt werden.

[0008] In neuromorphen Ansätzen wird Information meist stattdessen lokal im Zustand der einzelnen Neuronen gehalten. Ggfs. kann die Rate, mit der sich der Zustand des Neurons pro Zeitschritt ändert, skaliert und auf die relevante Zeitskala des Eingangssignals abgestimmt werden. Die Verarbeitung langsamer Signale mit einem schnellen Takt erfordert daher einen hoch aufgelösten internen Zustand der Neuronen.

- Um komplexere Funktionalitäten wie das Erkennen von Mustern mittels neuromorpher Schaltkreise abzubilden, müssen viele Neuronen sinnvoll miteinander verschaltet werden. Dies wird gegenwärtig durch verschiedene Mesh- und Crossbar-Routing-Systeme implementiert, welche bestimmte Konfigurationen zulassen und Output-Signale dem Input verschiedener Neuronen zuordnen. Die Verbindungen zwischen einzelnen Neuronen sind dabei meist unterschiedlich gewichtet, was einen entsprechenden Mechanismus zur verbindungsspezifischen Konfiguration und Signalübertragung erfordert.
- Um mit verrauschten Eingangssignalen umgehen zu können, sollte als Ausgangssignal nicht nur das gewünschte Signal, z.B. ob ein gegebenes Muster erkannt wurde oder nicht, sondern auch ein Maß der zugehörigen Unsicherheit generiert werden. Dies kann von bestehenden Ansätzen lediglich mittelbar unter Rückgriff auf bestimmte Netzwerkarchitekturen realisiert werden, ist aber nicht in der Hardware selbst angelegt.

[0009] Somit weisen die bestehenden Ansätze spike-basierter neuromorpher Hardware, welche auf gepulsten neuronalen Netzen (Englisch: spiking neural networks - SNN) beruhen, verschiedene Nachteile auf. So erfordert die Verwendung von Ganzzahlarithmetik und Paket-Routing den Einsatz von Mikroprozessoren, was die technische Komplexität der Hardware erhöhen und aufgrund ihrer sequentiellen Operation zu Latenzen führen kann. Auch kann die

Beschränkung auf einfache generische Neuronenmodelle mit gewichteten Verbindungen, welche nicht für die Analyse von kontinuierlichen Signalströmen entwickelt wurden, zur Verwendung von notwendigerweise großen Netzwerken führen, deren interne Kommunikation viel Platz-, Energie- und bzw. oder Zeitressourcen beanspruchen kann.

[0010] Eine Aufgabe der vorliegenden Erfindung ist es, einen neuromorphen Schaltkreis bereitzustellen, um die zuvor genannten technischen Probleme zu lösen bzw. die zuvor genannten technischen Eigenschaften zu realisieren. Insbesondere soll ein zu erkennendes Muster in einem binären Eingangssignal schneller und bzw. oder zuverlässiger als bisher bekannt erkannt werden können. Zumindest soll eine Alternative zu bekannten derartigen neuromorphen Schaltkreisen bereitgestellt werden.

[0011] Die Aufgabe wird erfindungsgemäß durch einen neuromorphen Musterdetektor mit den Merkmalen des Patentanspruchs 1 sowie durch eine neuromorphe Schaltkreisanordnung mit den Merkmalen des Patentanspruchs 13 gelöst. Vorteilhafte Weiterbildungen sind in den Unteransprüchen beschrieben.

[0012] Somit betrifft die Erfindung einen neuromorphen Musterdetektor, welcher ausgebildet ist, wenigstens zwei 1-Bit Eingangssignale eines zu erkennenden Musters zu erhalten, mit wenigstens zwei Vergleichsschaltungen, welche jeweils ausgebildet sind, eines der 1-Bit Eingangssignale zu erhalten, die Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen 1-Bit Eingangssignals innerhalb eines vorbestimmten Zeitraums zu zählen, die Anzahl der gezählten Zustände mit einem vorbestimmten Schwellwert der jeweiligen Vergleichsschaltung zu vergleichen und bei Überschreiten des Schwellwerts auf die erfolgte bzw. auf die erfolgreiche Erkennung des zu erkennenden Musters hinzuweisen. Der neuromorphe Musterdetektor ist vorzugsweise mittels Digitaltechnik umgesetzt.

[0013] Mit anderen Worten werden wenigstens zwei 1-Bit Datenströme, welche gemeinsam ein zu erkennendes Muster in Form einer parallelen Bitfolge enthalten, dem erfindungsgemäßen neuromorphen Musterdetektor in Form einer neuromorphen Schaltung zugeführt. Über eine vorbestimmte Anzahl von Bit, welche dem vorbestimmten Zeitraum entsprechen, werden nun die "high"-Zustände oder die "low"-Zustände, d.h. die hohen Signalpegel oder die niedrigen Signalpegel, gezählt. Diese Anzahl wird fortlaufend mit einem Schwellwert verglichen. Wird dieser Schwellwert überschritten, so wird hieraus geschlussfolgert, dass zu erkennende Muster in dem jeweiligen 1-Bit Datenstrom der jeweiligen Vergleichsschaltung erkannt zu haben. Dies wird von dem neuromorphen Musterdetektor nach außen angezeigt, z.B. über ein entsprechendes Ausgangssignal.

[0014] Auf diese Art und Weise kann erfindungsgemäß vergleichsweise einfach mittels einer neuromorphen Schaltung eine Mustererkennung in einem digitalen Signal erfolgen.

[0015] Gemäß einem Aspekt der Erfindung ist die eine Vergleichsschaltung der anderen Vergleichsschaltung erstrangig untergeordnet, wobei die übergeordnete Vergleichsschaltung ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der übergeordneten Vergleichsschaltung überschritten und zeitgleich von der erstrangig untergeordneten Vergleichsschaltung auf die erfolgte Erkennung des zu erkennenden Musters hingewiesen wird.

[0016] Dies kann es ermöglichen, die Entscheidung der übergeordneten Vergleichsschaltung von der Entscheidung der untergeordneten Vergleichsschaltung, das vorbestimmte Muster erkannt zu haben oder nicht, abhängig zu machen.

[0017] Gemäß einem weiteren Aspekt der Erfindung weist der neuromorphe Musterdetektor wenigstens eine weitere Vergleichsschaltung auf, welche parallel zu der untergeordneten Vergleichsschaltung angeordnet ist, wobei die übergeordnete Vergleichsschaltung ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der übergeordneten Vergleichsschaltung überschritten und zeitgleich von den erstrangig untergeordneten Vergleichsschaltungen jeweils auf die erfolgte Erkennung des zu erkennenden Musters hingewiesen wird.

[0018] Dies kann es ermöglichen, die Entscheidung der übergeordneten Vergleichsschaltung von der Entscheidung der beiden untergeordneten Vergleichsschaltungen, das vorbestimmte Muster erkannt zu haben oder nicht, abhängig zu machen.

[0019] Gemäß einem weiteren Aspekt der Erfindung weist der neuromorphe Musterdetektor wenigstens eine weitere Vergleichsschaltung auf, welche zweitrangig untergeordnet zu der erstrangig untergeordneten Vergleichsschaltung angeordnet ist, wobei die erstrangig untergeordnete Vergleichsschaltung ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der erstrangig untergeordneten Vergleichsschaltung überschritten und zeitgleich von der zweitrangig untergeordneten Vergleichsschaltung auf die erfolgte Erkennung des zu erkennenden Musters hingewiesen wird.

[0020] Dies kann es ermöglichen, die Entscheidung der erstrangig untergeordneten Vergleichsschaltung von der Entscheidung der zweitrangig untergeordneten Vergleichsschaltung, das vorbestimmte Muster erkannt zu haben oder nicht, abhängig zu machen.

[0021] Dabei können die zuvor beschriebenen Möglichkeiten der Anordnung von mehr als zwei Vergleichsschaltungen auch miteinander kombiniert werden, indem wenigstens zwei erstrangige und wenigstens eine zweitrangige Vergleichsschaltung verwendet und wie zuvor beschrieben miteinander und bzw. oder seitens der übergeordneten Vergleichsschaltung in Abhängigkeit gesetzt werden.

[0022] Gemäß einem weiteren Aspekt der Erfindung bilden die wenigstens drei Vergleichsschaltungen einen Binärbaum mit wenigstens zwei Ebenen. Unter einem Binärbaum, auch binärer Baum genannt, wird eine besondere Unterart eines Baumes verstanden, wie er in der Informatik für hierarchische Datenstrukturen verwendet wird. Der Ausgangspunkt, wie hier die übergeordnete Vergleichsschaltung, wird als Wurzel oder auch Binärbaumwurzel bezeichnet, von welcher sich der Binärbaum in verschiedenen Ebene wie hier der erstrangigen und zweitrangigen Vergleichsschaltungen einzeln oder paarweise verzweigt, bis der jeweilige Ast an einem Binärbaumblatt endet.

[0023] Entsprechend können die Eigenschaften und Vorteile derartiger hierarchischer Datenstrukturen auf die erfindungsgemäße neuromorphe Schaltung übertragen und dort genutzt werden.

[0024] Gemäß einem weiteren Aspekt der Erfindung sind die Vergleichsschaltungen identisch ausgebildet. Dies kann die Umsetzung vereinfachen, da der Entwurf der neuromorphen Schaltung mit geringerem Aufwand ausfallen kann, in dem das Design der Vergleichsschaltung mehrfach verwendet wird. Auch kann dies die Vergrößerung der Schaltung des neuromorphen Musterdetektors vereinfachen und hierdurch eine Skallierbarkeit ermöglichen.

[0025] Gemäß einem weiteren Aspekt der Erfindung wird bei Überschreiten des Schwellwerts ein 1-Bit Ausgangssignal der jeweiligen Vergleichsschaltung auf den "high"-Zustand, ansonsten auf den "low"-Zustand, gesetzt, oder umgekehrt. Dies kann es ermöglichen, dass Hinweisen der jeweiligen Vergleichsschaltung auf die erfolgte Erkennung des zu erkennenden Musters einfach umzusetzen.

[0026] Gemäß einem weiteren Aspekt der Erfindung sind die Vergleichsschaltungen ausgebildet, jeweils ein 1-Bit Steuersignal zu erhalten und in Reaktion auf einen "high"-Zustand oder auf einen "low"-Zustand des jeweiligen 1-Bit Steuersignals das 1-Bit Ausgangssignal der jeweiligen Vergleichsschaltung auf den "low"-Zustand zu setzen. Hierdurch kann eine Möglichkeit geschaffen werden, die entsprechende Vergleichsschaltung mittels des jeweiligen 1-Bit Steuersignals wieder zurückzusetzen. Mit anderen Worten kann die Vergleichsschaltung von außen resetted werden. Dies kann es insbesondere ermög-

lichen, alle Vergleichsschaltungen zurückzusetzen, um anschließend mit dem Erkennen eines neuen Musters beginnen zu können, ohne dass der zuvor erfolgte Vorgang auf dessen Ergebnis Auswirkungen haben kann.

[0027] Gemäß einem weiteren Aspekt der Erfindung gibt der vorbestimmte Schwellwert der Anzahl der Zustände der jeweiligen Vergleichsschaltung vor, wann das zu erkennende Muster als erkannt angesehen wird. Mit anderen Worten kann durch die Höhe des Schwellwerts in Relation zur Länge bzw. Kürze des vorbestimmten Zeitraums bzw. der vorbestimmten Anzahl von Bit des Eingangssignals vorbestimmt werden, wie deutlich eine Übereinstimmung zwischen dem jeweiligen Eingangssignal und dem vorbestimmten Muster vorliegen muss, um das vorbestimmte Muster im jeweiligen Eingangssignal als erkannt anzusehen. Dies kann für jedes zu erkennende Muster und für jede Vergleichsschaltung vorgegeben werden. Dies kann über die Konfiguration der Vergleichsschaltungen erfolgen.

[0028] Gemäß einem weiteren Aspekt der Erfindung weisen die Vergleichsschaltungen jeweils einen Schiebefensterdetektor auf, welcher jeweils ausgebildet ist, das jeweilige 1-Bit Eingangssignal zu erhalten und die Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen 1-Bit Eingangssignals innerhalb des vorbestimmten Zeitraums zu zählen. Dies kann die Umsetzung dieser Funktion der Vergleichsschaltungen einfach und bzw. oder zuverlässig ermöglichen.

[0029] Gemäß einem weiteren Aspekt der Erfindung erfolgt das Zählen der Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen 1-Bit Eingangssignals innerhalb des vorbestimmten Zeitraums mittels eines bidirektionalen Schieberegisters des jeweiligen Schiebefensterdetektors. Dies kann die Umsetzung dieser Funktion der Vergleichsschaltungen einfach und bzw. oder zuverlässig ermöglichen.

[0030] Gemäß einem weiteren Aspekt der Erfindung erhalten die Vergleichsschaltungen, vorzugsweise deren Schiebefensterdetektor, jeweils ein Taktsignal zur Steuerung der Verarbeitung der Pulse und ein Taktsignal zur Steuerung der Länge der Plateaus, wobei die beiden Taktsignale unterschiedlich sind. Unter einem Puls bzw. Spike ist der Zustand eines Signals im Zustand "high" nach und vor einem Zustand "low" zu verstehen. Unter einem Plateau eines Signals ist die Zeitdauer bzw. die Signallänge im Zustand "high" zu verstehen. Mit anderen Worten ist unter eine Plateau eine Funktion vergleichbar einem volatilen Cache-Zwischenspeicher zu verstehen, welcher für eine konfigurierbare Zeit, d.h. die Zeitdauer des Plateaus, ein Zwischenergebnis speichert. Auf diese Art und Weise kann die Mustererkennung der Vergleichsschaltungen und damit auch des neuromorphen Musterdetektors in Abhängigkeit von wenigstens zwei unterschiedlichen Taktsignalen erfolgen.

[0031] Die vorliegende Erfindung betrifft auch eine neuromorphe Schaltkreisanordnung mit einer Mehrzahl von neuromorphen Musterdetektoren wie zuvor beschrieben, wobei jeder neuromorphe Musterdetektor ausgebildet ist, das gleiche 1-Bit Eingangssignal zu erhalten, ein unterschiedliches 1-Bit Zufallszahlensignal zu erhalten, das jeweilige 1-Bit Eingangssignal mit dem entsprechenden 1-Bit Zufallszahlensignal zu verändern, und die Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen veränderten 1-Bit Eingangssignals innerhalb eines vorbestimmten Zeitraums zu zählen.

[0032] Unter einem 1-Bit Zufallszahlensignal ist ein Signal mit einer Bitfolge zu verstehen, welche zufällig erzeugt wurde. Dies kann deterministisch oder nichtdeterministisch erfolgen. Ein deterministisch erzeugtes 1-Bit Zufallszahlensignal kann auch als Pseudo-Zufallszahlensignal bezeichnet werden. Dabei kann die Verwendung eines pseudo-zufälligen 1-Bit Zufallszahlensignals vorteilhaft sein, da dies einfacher als ein nicht-deterministisches 1-Bit Zufallszahlensignal erzeugt werden und zur Erzielung der entsprechenden Eigenschaften und Vorteile ausreichend sein kann.

[0033] Somit können mehrere der zuvor beschriebenen neuromorphen Musterdetektoren parallel zueinander angeordnet und verwendet werden, um jeweils das gleiche vorbestimmte Muster in dem gleichen Eingangssignal zu erkennen. Hierbei können die beiden 1-Bit Datenströme jeweils unterschiedlich stochastisch verändert werden, so dass das gleiche Muster jeweils in unterschiedlichen Eingangssignalen der einzelnen neuromorphen Musterdetektoren erkannt werden muss. Dies kann eine Aussage über die Zuverlässigkeit der Mustererkennung erlauben, da die gleichen Eingangssignal mit dem zu erkennenden Muster durch die 1-Bit Zufallssignale unterschiedliche verfremdet bzw. gestört jeweils identisch durch die neuromorphen Musterdetektoren bearbeitet werden.

[0034] Diesbezüglich sei angemerkt, dass ein (pseudo-)zufälliges Maskieren eines Datenstroms in mehrere sich zufällig unterscheidende Datenströme auch unabhängig von einer neuromorphen Schaltkreisanordnung wie zuvor beschrieben und insbesondere unabhängig von einer Mehrzahl von neuromorpher Musterdetektoren wie zuvor beschrieben umgesetzt und angewendet werden kann. Dies kann es ermöglichen, die entsprechenden Eigenschaften und Vorteile auch unabhängig umzusetzen und anzuwenden.

[0035] Gemäß einem Aspekt der Erfindung weist wenigstens eine Vergleichsschaltung, vorzugsweise weisen alle Vergleichsschaltungen jeweils, ein Und-Gatter auf, welches ausgebildet ist, das jeweilige 1-Bit Eingangssignal und das entsprechende 1-Bit Zufallszahlensignal zu kombinieren. Hierdurch kann die Veränderung der gleichen Eingangssignal durch die unterschiedlichen stochastischen 1-Bit Zufallszahlensignal umgesetzt werde.

[0036] Gemäß einem weiteren Aspekt der Erfindung ist die neuromorphe Schaltkreisanordnung ausgebildet, die Anzahl der 1-Bit Ausgangssignale der jeweiligen Vergleichsschaltung, welche zeitgleich im "high"-Zustand oder im "low"-Zustand sind, zu erfassen und aus dem Verhältnis der Anzahl von 1-Bit Ausgangssignalen im "high"-Zustand oder im "low"-Zustand und der Anzahl der neuromorphen Musterdetektoren einen Grad der Übereinstimmung zwischen 1-Bit Eingangssignal und zu erkennendem Muster zu bestimmen. Hierdurch kann diese Information bestimmt und zur Verfügung gestellt werden.

[0037] Gemäß einem weiteren Aspekt der Erfindung weist wenigstens eine Vergleichsschaltung, vorzugsweise weisen alle Vergleichsschaltungen jeweils, einen Zeitmultiplexer auf, welcher ausgebildet ist, parallele Ausgangssignale der neuromorphen Musterdetektoren zu einem 1-Bit-Ausgangssignalder neuromorphen Schaltkreisanordnung zusammenzuführen. Auf diese Art und Weise kann ein einziger resultierenden 1-Bit Datenstrom als Ausgangssignal der neuromorphen Schaltkreisanordnung erzeugt werden.

[0038] Ein Ausführungsbeispiel und weitere Vorteile der Erfindung werden nachstehend im Zusammenhang mit den folgenden Figuren rein schematisch dargestellt und näher erläutert. Darin zeigt:

- **Fig. 1** eine schematische Darstellung eines Symbols eines Schaltkreises einer Population der **Fig. 2**;
- Fig. 2 eine schematische Darstellung eines Schaltkreises der Population der Fig. 1;
- **Fig. 3** eine schematische Darstellung eines Symbols eines Schaltkreises eines Neurons der **Fig. 4**;
- Fig. 4 eine schematische Darstellung eines Schaltkreises des Neurons der Fig. 3;
- **Fig. 5** eine schematische Darstellung eines Schaltkreises eines Binärbaumzweigs;
- Fig. 6 eine schematische Darstellung eines Schaltkreises eines Abschlusszweigs;
- Fig. 7 eine schematische Darstellung eines Symbols eines Schaltkreises eines Segments der Fig. 8;

- **Fig. 8** eine schematische Darstellung eines Schaltkreises des Segments der **Fig. 7**;
- **Fig. 9** eine schematische Darstellung eines Symbols eines Schaltkreises eines Schiebefensterdetektors der **Fig. 10**;
- Fig. 10 eine schematische Darstellung eines Schaltkreises des Schiebefensterdetektors der Fig. 9;
- **Fig. 11** eine schematische Darstellung eines Symbols eines Schaltkreises eines Zeitmultiplexers der **Fig. 12**; und
- **Fig. 12** eine schematische Darstellung eines Schaltkreises des Zeitmultiplexers der **Fig. 11**.

[0039] Fig. 1 zeigt eine schematische Darstellung eines Symbols eines Schaltkreises einer Population 1 der Fig. 2. Fig. 2 zeigt eine schematische Darstellung eines Schaltkreises der Population 1 der Fig. 1.

[0040] Unter einer Population 1 im Sinne von Computersoftware wird eine Anordnung von gleichen Computerprogrammen verstanden, welche gemeinsam die Population 1 bilden. Wird dies auf neuromorphe Schaltkreise übertragen, so kann die o.g. Population 1 mittels neuromorpher Schaltkreise als neuromorphe Schaltkreisanordnung 1 gebildet werden, indem mehrere neuromorphe Musterdetektoren 2, welche auch als Neuronen 2 bezeichnet werden können, gleicher Struktur in Form von identisch ausgebildeten neuromorphen Schaltkreisen miteinander zur Population 1 verschaltet werden, siehe Fig. 2.

[0041] Die Population 1 besteht dabei gemäß dem dargestellten Ausführungsbeispiel aus einer Anzahl K von Neuronen 2, von welchen in der Fig. 2 das erste, das zweite und das K-te Neuron 2 von links nach rechts dargestellt sind. Jedes Neuron 2 erhält denselben eingehenden Datenstrom E als Eingangssignal E, welches aus einer Anzahl N von einzelnen 1-Bit Eingangssignalen E₁-E_N besteht. Das Eingangssignal E enthält ein zu erkennendes Muster, welches auch als Pattern bezeichnet werden kann.

[0042] Jedes Neuron **2** erhält ferner parallel dasselbe Steuersignal **I**, welches aus einer Mehrzahl von einzelnen 1-Bit Steuersignalen I₁-I_N besteht. Das Steuersignal I kann zum Zurücksetzen von Vergleichsschaltungen **2**, auch Segmente **3** genannt, innerhalb der Neuronen **2** verwendet werden, wie weiter unten näher beschrieben werden wird.

[0043] Ferner erhält jedes Neuron 2 eine Anzahl N von binären Zufallszahlensignalen $\mathbf{M}_{1,1}$ - $\mathbf{M}_{K,N}$, welche deterministisch erzeugt und für jedes Neuron 2 unterschiedlich pseudo-zufällig sind. Genauer gesagt wird der Population 1 für jedes der N Eingangssignale \mathbf{E}_1 - \mathbf{E}_N und für jedes der K Neuronen 2 ein zufälliges 1-Bit Zufallssignal $\mathbf{M}_{1,1}$ - $\mathbf{M}_{K,N}$ zur Verfügung gestellt.

[0044] Des Weiteren erhält jedes Neuron 2 drei unterschiedliche Taktsignal CLK_{PLT}, CLK_{SPIKE} und CLK_{PROG}. Das Taktsignal CLK_{PLT} ist ein Taktsignal zur Steuerung der Länge der Plateaus der Vergleichsschaltungen 3, wie weiter unten noch näher erläutert werden wird. Das Taktsignal CLK_{SPIKE} ist ein Taktsignal zur Steuerung der Verarbeitung von Spikes, d.h. von Pulsen, der Vergleichsschaltungen 3, wie ebenfalls weiter unten noch näher erläutert werden wird. Das Taktsignal CLK_{PROG} ist ein Taktsignal eines Konfigurationssignals D_{PROG} bzw. D_{PROGO}, wie ebenfalls weiter unten noch näher erläutert werden wird.

[0045] Ein Konfigurationssignal D_{PROG} der Population 1 wird als Eingangssignal dem ersten Neuron 2 zugeführt, dort zur Konfiguration des ersten Neurons 2 verwendet und als Konfigurationssignal D_{PROGO} von dem ersten Neuron 2 an das zweite Neuron 2 ausgegeben. Das zweite Neuron 2 erhält somit das Konfigurationssignal D_{PROG} als Eingangssignal usw. Das Konfigurationssignal D_{PROGO} als Ausgangssignal des letzten K-ten Neurons 2 ist das Konfigurationsausgangssignal D_{PROGO} der Population 1.

[0046] Jedes der K Neuronen **2** erzeugt ein binäres Ausgangssignal P_1 - P_K , welche parallel einem Zeitmultiplexer **5** als dessen Eingangssignale S_1 - S_K zugeführt werden. Die Verarbeitung dieser Eingangssignale S_1 - S_K zu einem Ausgangssignal **O** des Zeitmultiplexers **5**, welches auch das Ausgangssignal **O** der gesamten Population **1** darstellt, wird weiter unter beschrieben.

[0047] Fig. 3 zeigt eine schematische Darstellung eines Symbols eines Schaltkreises eines Neurons 2 der Fig. 4. Fig. 4 zeigt eine schematische Darstellung eines Schaltkreises des Neurons 2 der Fig. 3. Fig. 5 zeigt eine schematische Darstellung eines Schaltkreises eines Binärbaumzweigs 21, 22. Fig. 6 zeigt eine schematische Darstellung eines Schaltkreises eines Abschlusszweigs 20.

[0048] Jedes Neuron 2 besteht im Wesentlichen aus einem rekursiv eingebetteten, binären Baum, auch Binärbaum genannt, mit einem ersten Binärbaumzweig 21, einem zweiten Binärbaumzweig 22 sowie dem zuvor bereits erwähnten Segments 3, siehe z.B. Fig. 4. Jeder der beiden Baumzweige 21, 22 kann in jeder Ebene des binären Baums entweder ein weiteres Neuron 2 mit zwei weiteren Binärbaumzweigen 21, 22 und einem Segment 3, siehe Fig. 5, oder ein Abschlusszweig 20 mit lediglich einem Segment 3, siehe Fig. 6, sein. Die beiden Binärbaumzweige 21, 22 können auch als innere Knoten des Binärbaums 21, 22 oder als Nested Branches 21, 22 bezeichnet werden. Der Abschlusszweig 20 kann auch als Binärbaumblatt 20 oder als Terminal Branch 20 bezeichnet werden. Das Neuron 2 selbst kann daher auch als Binärbaumwurzel 2 bezeichnet werden. Mit anderen Worten wird jeder Binärbaumzweige 21, 22 entweder aus einem weiteren Neuron 2, welches seinerseits wieder zwei Binärbaumzweige 21, 22 aufweist, oder aus einem Abschlusszweig 20 gebildet.

[0049] Dabei besitzt das jeweilige Segment 3, welches die Wurzel des Binärbaums bildet, die gleiche Struktur wie die Binärbaumzweige 21, 22 der weiteren Ebenen des binären Baums mit dem Unterschied, dass das Segment 3 der Wurzel des Binärbaums statt dem Taktsignal CLK_{PLT} das Taktsignal CLK_{SPIKE} erhält. Die Binärbaumzweige 21, 22 erhalten das Taktsignal CLK_{PLT}. Dies führt dazu, dass das Ausgangssignal P₁-P_k des Neurons 2 als kurze Spikes mit dem Taktsignal CLK_{SPIKE} und nicht lange Plateaus mit dem Taktsignal CLK_{PLT} aufweist.

[0050] Das Konfigurationssignal D_{PROG} des Neurons 2 wird jedem Binärbaumzweig 21, 22, jedem Abschlusszweig 20 sowie jedem Segment 3 zugeführt.

[0051] Jedes Segment 3 jeder Ebene des binären Baums erhält eines der 1-Bit Eingangssignale $\mathbf{E_1}$ - $\mathbf{E_N}$ sowie das entsprechende 1-Bit Steuersignale $\mathbf{I_1}$ - $\mathbf{I_N}$ und das entsprechende Zufallszahlensignal $\mathbf{M_{1,1}}$ - $\mathbf{M_{K,N}}$. Die Funktion des Segments 3 wird weiter unten erklärt werden. Auch erhält jedes Segment 3 jeder Ebene des binären Baums die Taktsignale $\mathbf{CLK_{PLT}}$, $\mathbf{CLK_{SPIKE}}$ und $\mathbf{CLK_{PROG}}$ zu den zugehörigen Signalen.

[0052] Der Abschlusszweig 20 besteht lediglich aus einem Segment 3 mit zwei konstanten Eingangssignalen $\mathbf{B_1}$, $\mathbf{B_2}$, welche beide den Zustand "high" aufweisen. Ferner erhält das Segment 3 ebenfalls eines der 1-Bit Eingangssignale $\mathbf{E_1}$ - $\mathbf{E_N}$ sowie das entsprechende 1-Bit Steuersignale $\mathbf{I_1}$ - $\mathbf{I_N}$ und das entsprechende Zufallszahlensignal $\mathbf{M_{1.1}}$ - $\mathbf{M_{K.N}}$.

[0053] Fig. 7 zeigt eine schematische Darstellung eines Symbols eines Schaltkreises eines Segments 3 der Fig. 8. Fig. 8 zeigt eine schematische Darstellung eines Schaltkreises des Segments 3 der Fig. 7.

[0054] Das Segment 3, welches wie zuvor beschrieben jeweils identisch in jedem Neuron 2 mehrfach auf verschiedenen Ebenen des binären Baums verwendet wird, erhält stets die Ausgangssignale P der Binärbaumzweige 21, 22 derselben Ebene als Eingangssignal B₁, B₂. Die beiden Eingangssignale B₁, B₂ sind parallel sowohl auf ein erstes Oder-Gatter 30 als auch auf ein erstes Und-Gatter 31 geschaltet. Die Ausgangssignale der beiden ersten Gatter 30, 31 können entweder ein konstantes "high"-Signal oder ein konstantes "low"-Signal sein, welche parallel einem 4-fach Multiplexer 32 zugeführt werden. Zusätzlich zu den beiden Ausgangssignalen der beiden ersten Gatter 30,31 werden ein konstantes "low"-Signal und ein konstantes "high"-Signal parallel dem 4-fach Multiplexer 32 zugeführt. Dabei wird das Ausgangs-

signal des 4-fach Multiplexers **32** von einem ersten 2-bit SIPO Schieberegister **33** (SIPO: serial-input-parallel-output) gewählt und einem zweiten Und-Gatter **34** zugeführt.

[0055] Die zwei 1-Bit Eingangssignale E_1 - E_N und M_{1.1}-M_{K.N} des jeweiligen Segments 3 werden von einem dritten Und-Gatter 35 verschaltet, dessen Ausgangssignal in einen Schiebefensterdetektor 4 als dessen Eingangssignal D_{IN} geschaltet wird, welcher auch als Slider 4 bezeichnet werden kann und weiter unten näher erläutert werden wird. Der Schiebefensterdetektor 4 wird durch den Datenstrom des Konfigurationssignals **D**_{PROG} mit dem zugehörigen Taktsignal CLK_{PROG} konfiguriert. Das Ausgangssignal des Konfigurationssignals **D**_{PROGO} des Schiebefensterdetektors 4 ist wiederum das Eingangssignal des ersten 2-bit SIPO Schieberegisters 33, welches seinerseits durch die steigende Flanke des Taktsignals CLK_{PROG} weitergeschoben wird. Das erste 2-bit SIPO Schieberegister 33 erzeugt dabei parallel zu der zuvor beschriebenen Auswahl des Ausgangssignals des 4fach Multiplexers 32 das Ausgangssignal des Konfigurationssignals **D**_{PROGO} des Segments **3**.

[0056] Der Schiebefensterdetektor 4 erzeugt parallel zu dem Konfigurationssignals D_{PROGO} des Schiebefensterdetektors 4 ferner ein Ausgangssignal D_{OUT} , welches das zweite Eingangssignal des zweiten Und-Gatters 34 ist, dessen Ausgangssignal einen 1-Bit Flipflop 36 zu jeder steigenden Flanke in den "high"-Zustand versetzt. Das Ausgangssignal des 1-Bit Flipflops 36 ist auch das Ausgangssignal P des jeweiligen Segments 3.

[0057] Zu jeder steigenden Flanke des Taktsignals **CLK**_{PLT} wird ein zweites N-bit SIPO Schieberegister 37 um einen Schritt geschoben, wodurch das aktuelle Ausgangssignal des 1-Bit Flipflops 36 ausgelesen wird. Das letzte Bit des parallelen Ausgangssignals des zweites N-bit SIPO Schieberegisters 37 bildet ein Eingangssignal eines zweiten Oder-Gatters 38. Das andere Eingangssignal des zweiten Oder-Gatters 38 ist das entsprechende 1-Bit Steuersignale I₁-I_N. Wenn eines der beiden Eingangssignale des zweiten Oder-Gatters 38 den "high"-Zustand aufweist, ist auch das Ausgangssignal des zweiten Oder-Gatters 38 "high" und die steigende Flanke schaltet den Zustand des 1-Bit Flipflops 36 zurück sowie setzt hierdurch alle Bits des zweites N-bit SIPO Schieberegisters 37 auf "low", d.h. in den "low"-Zustand.

[0058] Fig. 9 zeigt eine schematische Darstellung eines Symbols eines Schaltkreises eines Schiebefensterdetektors 4 der Fig. 10. Fig. 10 zeigt eine schematische Darstellung eines Schaltkreises des Schiebefensterdetektors 4 der Fig. 9.

[0059] Der Schiebefensterdetektor 4 dient dazu zu erkennen, ob die Anzahl der "high"-Bits, d.h. der Bits

im "high"-Zustand, in seinem Eingangssignal D_{IN} innerhalb der letzten N-bits, d.h. innerhalb eines vorbestimmten Zeitraums, welcher durch das Konfigurationssignal D_{PROG} konfigurierbar ist, des 1-Bit Eingangssignals D_{IN} einen konfigurierbaren Schwellwert übersteigt. Hierzu wird das 1-Bit Eingangssignal D_{IN} des Schiebefensterdetektors 4 einem ersten Und-Gatter 40 als dessen erstes Eingangssignal zugeführt.

[0060] Das Ausgangssignal des ersten Und-Gatters 40 wird als Eingangssignal in ein erstes N-bit SIPO Schieberegister 41 geleitet, welches das serielle Eingangssignal parallelisiert und mit jeder steigenden Flanke im Taktsignal CLK_{IN} einen Schritt weitergeschoben wird. Das N-te parallele Ausgangssignal des ersten N-bit SIPO Schieberegisters 41 ist das linksschiebende Eingangssignal SL in ein zweites bidirektionales M-bit SIPO-Schieberegister 42. Das Ausgangssignal des ersten Und-Gatters 40 selbst ist das rechtsschiebende Eingangssignal SR des zweiten bidirektionalen M-bit SIPO-Schieberegisters 42.

[0061] Zu jeder steigenden Flanke des Taktsignals CLK_{IN} wird das zweite bidirektionale M-bit SIPO-Schieberegister 42 einen Schritt in die Richtung nach rechts geschoben, falls das rechtsschiebende Eingangssignal SR "high" und das linksschiebende Eingangssignal SL "low" ist. Wenn das rechtsschiebende Eingangssignal SR "low" und das linksschiebende Eingangssignal SL "high" ist, wird das zweite bidirektionale M-bit SIPO-Schieberegister 42 hingegen in die Richtung nach links geschoben. Ansonsten bleibt das zweite bidirektionale M-bit SIPO-Schieberegister 42 unverändert.

[0062] Falls das zweite bidirektionale M-bit SIPO-Schieberegister 42 in die Richtung nach rechts geschoben wird, wird ein "high"-Bit von links eingefügt. Falls hingegen das zweite bidirektionale M-bit SIPO-Schieberegister 42 in die Richtung nach links geschoben wird, wird ein "low"-Bit von rechts eingefügt. Das letzte Bit des parallelen Ausgangssignals des zweiten bidirektionalen M-bit SIPO-Schieberegisters 42 wird invertiert als zweites Eingangssignal des ersten Und-Gatters 40 genutzt.

[0063] Ein M+1fach Multiplexer 43 wird von K 1-Bit Eingangssignalen konfiguriert und generiert so entweder ein konstantes "high"-Ausgangssignal oder selektiert einen der M parallelen Ausgangssignale des zweiten bidirektionalen M-bit SIPO-Schieberegisters 42. Das selektierte Signal ist das Ausgangssignal des M+1fach Multiplexers 43 und des gesamten Schiebefensterdetektors 4.

[0064] Welches der M+1 <= 2^K (M plus 1 kleinergleich 2 hoch K) Eingangssignale mittels des M +1 fach Multiplexers 43 ausgesucht wird, wird von dem parallelen Ausgangssignals eines dritten K-bit

SIPO-Schieberegisters **44** festgelegt, welches von einem Bitstrom des Eingangssignals **D**_{PROG} mit einem dazugehörigen Taktsignals **CLK**_{PROG} betrieben wird. Das letzte parallele Ausgangssignal des dritten K-bit SIPO-Schieberegisters **44** ist der zusätzliche Konfigurationssignal **D**_{PROGO} als Ausgangssignal des Schiebefensterdetektors **4**, um mehrere Segmente **3** bzw. Neuronen **2** in Serie verschalten zu können.

[0065] Fig. 11 zeigt eine schematische Darstellung eines Symbols eines Schaltkreises eines Zeitmultiplexers 5 der Fig. 12. Fig. 12 zeigt eine schematische Darstellung eines Schaltkreises des Zeitmultiplexers 5 der Fig. 11.

[0066] Der Zeitmultiplexer 5 ist in der Lage, eine Folge von K 1-Bit parallelen Eingangssignalen $\mathbf{S_1}$ - $\mathbf{S_K}$ in ein serielles 1-Bit Ausgangssignal \mathbf{O} zu enkodieren. Die steigende Flanke eines der K Eingangssignale $\mathbf{S_1}$ - $\mathbf{S_K}$ setzt ein korrespondierendes Flipflop 50 einer Anzahl \mathbf{K} von identischen und parallel zueinander angeordneten Flipflops 50 in den "high"-Zustand. Die Ausgangssignale der Flipflops 50 sind jeweils eines der beiden Eingangssignale eines jeweils korrespondierenden Und-Gatters 51 einer Anzahl \mathbf{K} von identischen und parallel zueinander angeordneten Und-Gattern 51.

[0067] Zu jeder steigenden Flanke des Taktsignals CLK_{SPIKE} wird ein selbst initialisierter K-bit Ringzähler 52 weitergeschoben, dessen parallele Ausgangssignale jeweils das zweite Eingangssignal der Und-Gatter 51 sowie das zurücksetzende Signal, d.h. das Reset-Signal, für die Flipflops 50 sind. Zu jedem Zeitpunkt ist genau ein Bit des Ringzählers 52 im Zustand "high" während alle anderen Bit des Ringzählers 52 im Zustand "low" sind. Zur fallenden Flanke des Reset-Signals wird das jeweilige Flipflop 50 in den "low"-Zustand geschaltet.

[0068] Während beide Eingangssignale eines der Und-Gatter 51 im Zustand "high" sind, ist auch das Ausgangssignal dieses Und-Gatters 51 im Zustand "hoch", ansonsten im Zustand "low". Wenn eines der K Und-Gatter 50 ein Ausgangssignal im Zustand "hoch" hat, ist das Ausgangssignal eines Oder-Gatters 53 ebenfalls im Zustand "high", sonst im Zustand "low". Zur steigenden Flanke des Taktsignals CLK_{SPIKE} wird das Ausgangssignal des Oder-Gatters 53 für einen Taktzyklus in einem D-Flipflop 54 zwischengespeichert. Das Ausgangssignal des D-Flipflops 54 ist das Ausgangssignal des Zeitmultiplexers

[0069] Das Taktsignal CLK_{PROG} ist an den Datenstrom des Konfigurationssignals D_{PROG} zur Konfiguration der Segmente 3 gekoppelt und hat lediglich die Funktion, die Segmente 3 innerhalb des jeweiligen Neurons 2 sowie die Neuronen 2 innerhalb der Population 1 untereinander zu synchronisieren.

[0070] Das Taktsignal CLK_{SPIKE} steuert die Verarbeitung von sog. "Spikes", d.h. von Pulsen als "high"-Zustände. Zum einen wird mit der Frequenz des Taktsignales CLK_{SPIKE} das Ausgangssignal der Population 1 in der Zeit multiplexed. Zum anderen wird das Taktsignal CLK_{SPIKE} im zweiten bidirektionalen M-bit SIPO-Schieberegister 42 des Schiebefensterdetektors 4 genutzt, um dieses Eingangssignal synchronisiert zu verarbeiten. Somit ist der ausgehende Datenstrom des zweiten bidirektionalen M-bit SIPO-Schieberegisters 42 an das Taktsignal CLK_{SPIKE} gebunden. Auch hängen alle eingehenden Datenströme des Eingangssignals E, des Kontrollsignals I sowie der binären Zufallszahlensignal M, welche zur Mustererkennung dienen, an dem Taktsignal CLK_{SPIKE}.

[0071] In der Verarbeitung des Eingangssignals E, des Kontrollsignals I sowie der binären Zufallszahlensignal M zur Mustererkennung gilt insbesondere, dass das Zeitfenster des Schiebefensterdetektors 4 N * 1/f(CLK-spike) ist, also durch den Horizont des n bidirektionalen M-bit SIPO-Schieberegister 42 des Schiebefensterdetektors 4 und durch die Frequenz des Taktsignals gegeben ist. In der Anwendung lässt sich durch die Wahl der Frequenz die Population 1 auf die Zeitskalen anpassen, auf denen Teilmuster erkannt werden sollen, wobei ein Teilmuster das ist, was ein Segment 3 alleine durch den Schiebefensterdetektor 4 erkennt.

[0072] Das Taktsignal CLK_{PLT} steuert ausschließlich die Länge der Plateaus in den einzelnen Segmente 3, also die Zeitdauer bzw. Signallänge, für die ein einzelnes Segment 3 sich die Erkennung eines Teilmusters zusammen mit ausreichendem Signal aus dem binären Baum merkt: Cache für das Zwischenergebnis. Im Speziellen wird das asynchron geschaltete "high"-Ausgangssignal des Segments 3 nach mindestens N * 1/f(CLK_{PLT}) und nach maximal (N+1) * 1/f(CLK_{PLT}) wieder ausgeschaltet. Die Spanne ergibt sich dadurch, dass das Zählen im Schiebefensterregister 4 zum Ausschalten vom Anschalten des Ausgangssignals entkoppelt ist. Damit lässt sich über die Wahl von N die zeitliche Präzision auf Kosten von Bauteilen und über die gemeinsame Wahl von N und der Frequenz des Taktsignals CLK_{PLT} die Zeitskala regeln, auf der Zwischenergebnisse und Teilmuster gespeichert werden. Das Taktsignal CLK_{PLT} stellt somit eine zweite Zeitskala in der Mustererkennung dar.

[0073] Die Kombination der Taktsignale CLK_{SPIKE} und CLK_{PLT}, um Teilmuster auf zwei unabhängig wählbaren Zeitskalen zur Mustererkennung zu kombinieren, stellt eine Besonderheit der Neuronen 2 dar. Isoliert kontrolliert jedes Taktsignal CLK_{SPIKE} und CLK_{PLT} wie bisher üblich einen Teil des Neurons 2 über Flankensteuerung. Genauer betrachtet werden jedoch erfindungsgemäß die Segmente 3 innerhalb des Neurons 2 von den verschiedenen Taktsignalen CLK_{SPIKE} und CLK_{PLT} gesteuert und dies zur Im-

plementierung von Algorithmen zur Mustererkennung verwendet.

[0074] Die zuvor beschriebene Population 1 kann dazu verwendet werden, mit niedriger Latenz Muster in kontinuierlichen, digitalen Datenströmen (Bitstreams) zu erkennen. Da aufgrund von Störsignalen oder zeitlicher Impräzision niemals dieselben Muster in gleicher Form auftreten, können dabei auch ungefähre Übereinstimmungen erkannt und der Grad der Übereinstimmung quantifiziert werden. Dabei sind die zu erkennenden Muster konfigurierbar, d.h. können vorbestimmt werden.

[0075] Hierzu werden die zuvor beschriebenen Neuronen 2 als mehrere Musterdetektoren in Gruppen in Form von Populationen 1 zusammengefasst. Jedes einzelne Neuron 2 ist hier eine hierarchische Struktur der Segmente 3, welche untereinander verknüpft sind und, je nach problemspezifischer Konfiguration, jeweils eigene Eingangssignale verarbeiten. Wenn ein komplexes Muster als Eingangssignal alle Segmente 3 in der richtigen zeitlichen Sequenz aktiviert, erzeugt das jeweilige Neuron 2 in seinem Ausgangssignal ein positives Bit, d.h. ein Ausgangssignal mit dem Zustand "high"; sozusagen "feuert" das Neuron 2 bzw. das Neuron 2 erzeugt einen Pulse bzw. einen "Spike".

[0076] Die Wahrscheinlichkeit, mit der ein einzelnes Neuron 2 feuert, reflektiert dabei den Grad der Übereinstimmung zwischen dem geforderten, d.h. dem konfigurierten vorbestimmten, und dem gesehenen, d.h. der Population 1 zugeführten, Muster. In einer Population 1 lesen alle K gleichkonfigurierten Neuronen 2 den gleichen Datenstrom als Eingangssignal E und versuchen, das gleiche Muster in dem Eingangssignal E zu erkennen, erhalten jedoch durch eine pseudo-zufällige Maskierung der Eingangssignale E mit den binären Zufallszahlensignalen M stochastisch voneinander verschiedene Eingangssignale E. Dies bedeutet, dass auf jedes Muster W von K Neuronen 2 reagieren, wobei W den Grad der Übereinstimmung zwischen dem zugeführten Muster und dem konfigurierten vorbestimmten Muster abbildet. Die technische Umsetzung kommt hierbei gänzlich ohne Mikroprozessoren aus und ist gänzlich in den zuvor beschriebenen Schaltkreisen umsetzbar.

[0077] Hierzu wird der eingehende Datenstrom E, in welchem ein Muster erkannt werden sollen, als getaktetes binäres Signal E in Form von N 1-Bit Eingangssignalen $\mathbf{E_1}$ - $\mathbf{E_N}$ auf mehreren parallelen Leitungen gelegt und der Population 1 zugeführt. Dies gilt ebenso für das Kontrollsignal I und die binären Zufallszahlensignale \mathbf{M} .

[0078] Innerhalb der Population 1 werden auf der Eingangsseite die gleichen Eingangssignale E und Kontrollsignale I an jedes Neuron 2 geleitet, wo die Eingangssignale **E** mit den neuronenspezifischen binären Zufallssignalen **M** maskiert werden. Die einzelnen Ergebnisse der Neuronen **2** werden dann im Zeitmultiplexer **5** zusammengeführt, um einen einzelnen Datenstrom **O** als Ausgang der Population **1** zu generieren, welcher wie gefordert die Qualität des erkannten Musters in den eingehenden Datenstrom **E** widerspiegelt.

[0079] Jedem einzelnen Neuron 2 innerhalb Population 1 kommt dabei die Aufgabe zu, das konfigurierbare, vorbestimmte Muster im jeweiligen 1-Bit Eingangssignal E₁-E_N der N 1-Bit Eingangssignale E₁-E_N zu erkennen. Hierzu sind die Neuronen 2 jeweils aus den einzelnen N Segmenten 3 aufgebaut, von denen jedes eines der eingehenden N 1-Bit Eingangssignale E₁-E_N verarbeitet. Jedes Segment 3 reagiert dabei auf ein relevantes Signal in seinem zugeordneten 1-Bit Eingangssignal E₁-E_N, d.h. das k-te Segment 3 auf ein relevantes Signal im 1-Bit Eingangssignal E_k, indem das Segment 3 für eine bestimmte Zeit eingeschaltet, d.h. in den "high"-Zustand versetzt, wird. Untereinander sind diese Segmente 3 in dem binären Baum derart verschaltet, dass jedes einzelne Segment 3 nur dann durch das jeweilige 1-Bit Eingangssignal $\mathbf{E_1}$ - $\mathbf{E_N}$ eingeschaltet werden kann, wenn - je nach Konfiguration - Null, Eins oder Zwei der untergeordneten Binärbaumzweige 21, 22 oder Abschlusszweige 20 im binären Baum bereits eingeschaltet sind. Wie lange ein Segment 3 eingeschaltet ist, wird durch das Taktsignal CLK_{PLT} festgelegt, welches nicht an das Taktsignal CLK SPIKE des Eingangssignals E gekoppelt ist.

[0080] In jedem Neuron 2 ist diese Verschachtelung in dem Binärbaum abgebildet. Ein Neuron 2 hat für jedes Segment 3 ein jeweils zugeordnetes 1-Bit Eingangssignal E₁-E_N der N 1-Bit Eingangssignale E₁-E_N, mit welchem das jeweilige Segment 3 für eine feste Zeit eingeschaltet, d.h. in den Zustand "high" gebracht, werden kann (Plateau). Jedes Segment 3 hat ebenso ein jeweils zugeordnetes 1-Bit Kontrollsignal I₁-I_N der N 1-Bit Kontrollsignale I₁-I_N, mit welchem das Segment 3, falls es bereits in den Zustand "high" ist, durch das jeweils zugeordnete 1-Bit Kontrollsignal I₁-I_N als externes Signal wieder ausgeschaltet, d.h. in den "low"-Zustand gebracht, werden kann. Einzelne Segmente 3 bekommen das konfigurierbare Taktsignal CLK_{PLT}, welches die zeitliche Dauer bestimmt, für die ein Segment 3 eingeschaltet ist. Das Segment 3 an der Wurzel der Baumstruktur, d.h. in der obersten Ebene es binären Baums, generiert kurze Pulse, auch Spikes genannt, mit derselben Taktung des Taktsignals CLK_{SPIKE} wie die Eingangssignale E anstatt längere Plateaus zu erzeugen, wie in den übrigen Segmenten 3.

[0081] Jedes der Segmente 3 wird zunächst durch die Eingangssignale B1, B2 anderer im binären Baum untergeordneter Segmente 3 getrieben, sofern die-

se existieren. Hier kann konfiguriert werden, ob Null, Eins oder Zwei Segmente 3 eingeschaltet sein müssen. Weiter wird der eingehende Datenstrom E in dem Schiebefensterdetektor 4 verarbeitet, welches für kurze Zeit eingeschaltet ist, falls die Anzahl der gesetzten Bits in einem festen Zeitfenster einen kritischen Pegel überschreitet. Das Eingangssignal in den Schiebefensterdetektor 4 wird vorher mit dem binäres Zufallszahlensignal M maskiert. So ist die Antwort jedes Schiebefensterdetektors 4 auf dasselbe Eingangssignal E stochastisch und unterscheidet sich, wie oben beschrieben, von anderen Schiebefensterdetektoren 4 in der Population 1, welche auf das gleiche Eingangssignal E abweichend reagieren.

[0082] Falls sowohl das jeweils zugeordnete 1-Bit Eingangssignal E₁-E_N das konfigurierte Kriterium der untergeordneten Binärbaumzweige 21, 22 bzw. Abschlusszweige 20 erfüllt und der Schiebefensterdetektor 4 des Segments 3 im zugeordneten 1-Bit Eingangssignal E₁-E_N ein Signal erkannt hat, schaltet sich das Segment 3 ein. In dem zweiten bidirektionalen M-bit SIPO-Schieberegister 42 wird dieser Zustand für eine feste Anzahl an Takten des Taktsignals **CLK**_{PLT} gehalten, worauf sich das Segment 3 selbst wieder ausschaltet. Weiter kann das jeweils zugeordnete 1-Bit Kontrollsignal I₁-I_N den zweiten bidirektionalen M-bit SIPO-Schieberegister 42 zurücksetzen und das Segment 3 frühzeitig ausschalten. Intern ist das Ausgangssignal des Segments 3 lediglich indirekt über den Schiebefensterdetektor 4 an ein Taktsignals CLK_{SPIKE} gebunden. Das Segment 3 reagiert ansonsten mit einer zu vernachlässigen Verzögerung der einzelnen Bauteile.

[0083] Dem Schiebefensterdetektor 4 kommt dabei die Aufgabe zu, zu detektieren, falls die Anzahl der gesetzten Bits, d.h. der Bits im "high"-Zustand, in einem festen vorbestimmten Zeitfenster des getakteten Datenstroms einen kritischen Wert überschreitet. Hierzu wird seitens des Schiebefensterdetektors 4 pro Segment 3 die Anzahl der eingehenden Pulse, d.h. der Bits im Zustand "high", in dem festen Zeitfenster in dem jeweiligen Datenstrom D_{IN} gezählt. Jedes eingehende gesetzte Bit, d.h. Bit im Zustand "high", schiebt das zweite bidirektionale M-bit SIPO-Schieberegister 42 vorwärts und wird gleichzeitig in einer durch das zweite bidirektionale M-bit SIPO-Schieberegister 42 und durch das Taktsignal **CLK**_{SPIKE} des Datenstrom **E** implementierten Delayline gespeichert. Nach diesem Delay wird das zweite bidirektionale M-bit SIPO-Schieberegister 42 wieder zurückgeschoben.

[0084] Konfigurierbar ist, an welcher Stelle im zweiten bidirektionalen M-bit SIPO-Schieberegister **42** ein Bit gesetzt sein muss, um ein Ausgangssignal zu erzeugen. So signalisiert das zweite bidirektionale M-bit SIPO-Schieberegister **42**, wann mehr als ein konfigurierbarer, vorbestimmter Schwellwert 1-Bit-Signa-

le in dem durch das Taktsignal ${\sf CLK_{SPIKE}}$ und durch die Länge der Delayline festgelegten Zeitfenster im Datenstrom ${\sf D_{IN}}$ zu finden waren.

[0085] Der Schiebefensterdetektor 4 dekodiert somit 1-Bit Signale, welche genau der Kodierung des Ausgangssignals der Population 1 entsprechen. Die Anzahl der Pulse in kurzer Zeit kodiert dabei die Stärke des Signals. Der Schwellwert im zweiten bidirektionalen M-bit SIPO-Schieberegister 42 legt fest, wann ein Signal stark genug war.

[0086] Das Konfigurationssignal D_{PROG} , welches sich durch alle Bauteile der Population 1 zieht, ermöglicht die Konfiguration der Population 1 und aller enthaltener Bauteile.

[0087] Im Vergleich zu bekannten Lösungen zur digitalen Signalverarbeitung und Mustererkennung, besonders im Bereich neuromorpher Technologien, ergeben sich eine Reihe von Vorteilen aus den zuvor beschriebenen Neuronen 2 sowie der hieraus aufgebauten Population 1.

[0088] So ermöglicht der zuvor beschriebene Ansatz die Erkennung von konfigurierbaren Mustern auf verschiedenen Zeitskalen und ist somit tolerant gegenüber Störungen im Signal oder im Timing. Dies erlaubt den Einsatz in erschwerten Bedingungen, z.B. im Verbund mit unpräziser Sensorik oder mit Signalen mit hoher Variabilität.

[0089] Auch kann durch die Nutzung stochastischer Eingangssignale als die binären pseudo-zufälligen Zufallszahlensignale **M** nicht nur ein gegebenes Muster erkannt werden, sondern es kann auch der Grad der Übereinstimmung quantifiziert werden.

[0090] Sowohl das Eingangssignal **E** als auch das Ausgangssignal **O** der Population **1** sind kompatibel, um mit weiteren Populationen zu kommunizieren, und erlauben somit die Verschaltung zu großen Netzen.

[0091] Die Informationsverarbeitung erfolgt gänzlich ohne den Einsatz von Mikroprozessoren oder Paket-Routing, was technisch einfacher umsetzbar ist, ein hohes Maß an Parallelisierung ermöglicht und zu niedrigen Latenzen führt.

[0092] Die Kommunikation zwischen Populationen 1 und das An- bzw. Ausschalten von Segmenten 3 ist an zwei verschiedene Taktsignale, nämlich die Taktsignale CLK_{PLT} und CLK_{SPIKE}, gebunden. Hierdurch wird das Taktsignal CLK_{SPIKE}, auf welchem Muster im Datenstrom E als Eingangssignal E erkannt werden sollen, von dem Taktsignal CLK_{PLT} entkoppelt. Auf diese Art und Weise können im bestimmungsgemäßen Gebrauch im Datenstrom E Teilmuster erkannt werden, welche auf einer von dem Datenstrom

CLK_{PROG}

 D_{IN}

D_{OUT}

D_{PROG}

D_{PROGO}

E, E₁-E_N

i

Κ

Ν

0

P₁-P_k

S₁-S_K

 $M, M_{1.1}-M_{K.N}$

 I, I_1-I_N

Taktsignal des Konfigu-

rationssignals D_{PROG},

Eingangssignal eines

Ausgangssignal eines

Schiebefensterdetektors

Konfigurationssignal als

Konfigurationssignal als

Eingangssignal der Po-

pulation 1; eingehender

Anzahl der Eingangs-

Anzahl der Neuronen

1-Bit bzw. binäres Zu-

Anzahl der Segmente

Zeitmultiplexers 5 bzw.

1-Bit Ausgangssignale

schlusszweige 20, der

und der Segmente 3

Eingangssignale des

der Neuronen 2, der Ab-

Binärbaumzweige 21, 22

Ausgangssignal des

der Population 1

fallszahlensignale

signale des Abschluss-

Eingangssignal

Ausgangssignal

Datenstrom

Kontrollsignal

Zählindex

zweigs 20

Schiebefensterdetektors

D_{PROGO}

E entkoppelten Zeitskala, nämlich dem Taktsignal **CLK**_{SPIKE}, mit anderen Teilmustern des Datenstroms **E** verbunden werden.

[0093] Zum Beispiel können viele Pulse im Datenstrom E in sehr kurzer Zeit übertragen werden und auf ein wichtiges Ereignis wie zum Beispiel das Überschreiten eines kritischen Wertes eines Temperatursensors hinweisen. Ein zweites Ereignis wie zum Beispiel das Überschreiten eines kritischen Wertes eines Beschleunigungssensors kann ebenfalls schnell mittels des Taktsignals **CLK**_{SPIKE} übertragen werden. Beide Ereignisse können als Teil eines Muster "kritische Temperatur und dann kritische Beschleunigung" dann aber auf einer Zeitskala, welche von dem Taktsignal **CLK**_{SPIKE} entkoppelt und durch das Taktsignal CLK_{PLT} zum Beispiel deutlich langsamer definiert ist, kombiniert werden. Durch die Kombination beider Taktsignale CLK_{SPIKE}, CLK_{PLT} kann ein Segment 3 auf die speziellen externen Timing-Anforderrungen der Anwendung angepasst werden.

[0094] Werden die Neuronen 2 mit einer höheren Komplexität in Form eines binären Baums mit zahlreichen Ebenen umgesetzt, so können mehr Informationen im internen Zustand der Neuronen 2 verarbeitet und gespeichert werden. Daher sind für dieselbe Leistung weniger individuelle Neuronen 2 erforderlich, was die Größe der resultierenden Population 1 und damit die Komplexität der notwendigen Kommunikationsinfrastruktur deutlich reduzieren kann.

[0095] Weitere Ausgestaltungen der Erfindung, welche von dem betrachteten Ausführungsbeispiel abweichen, sind vorstellbar. Jedes einzelne der oben genannten Bauteile kann in seinem Funktionsumfang erweitert oder in der Umsetzung angepasst werden. Auch können mehrere Populationen **1** zu Netzen verschaltet werden, die eingesetzt werden könnten, um komplexere Probleme zu lösen.

Bezugszeichenliste		S ₁ S _K	Zeitmultiplexers 5
B ₁	Eingangssignal eines Segments 3 seitens ei- nes ersten Binärbaum- zweigs 21	SL	linksschiebendes Eingangssignal des zweiten bidirektionalen (M-bit SIPO-) Schieberegisters 42 des Schiebefenster-
B_2	Eingangssignal eines		detektors 4
	Segments 3 seitens ei- nes zweiten Binärbaum- zweigs 22	SR	rechtsschiebendes Ein- gangssignal des zweiten bidirektionalen (M-bit SI-
CLK _{PLT}	Taktsignal zur Steue- rung der Länge der Pla- teaus der Vergleichs- schaltungen 3		PO-) Schieberegisters 42 des Schiebefensterdetektors 4
		W	Grad der Übereinstim-
CLK _{SPIKE}	Taktsignal zur Steue- rung der Verarbeitung der Spikes der Ver- gleichsschaltungen 3		mung zwischen zuge- führtem Muster und kon- figurierten vorbestimm- ten Muster

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1	neuromorphe Schalt- kreisanordnung; Popula-	52	selbst initialisierter (K- bit) Ringzähler	
	tion	53	Oder-Gatter	
2	neuromorpher Muster- detektor; Neuron; Neu-	54	D-Flipflop	
	ronen-Schaltkreis; Binär- baumwurzel	Patentansprüche		
20	Abschlusszweig; Bi- närbaumblatt; Terminal Branch	1. Neuromorpher Musterdetektor (2), welcher ausgebildet ist, wenigstens zwei 1-Bit Eingangssignale ($\rm E_1$ - $\rm E_N$) eines zu erkennenden Musters		
21	erster Binärbaumzweig; erster innerer Knoten des Binärbaums; erster Nested Branch	zu erhalten, mit wenigstens zwei Vergleichsschaltungen (3), welche jeweils ausgebildet sind, eines der 1-Bit Eingangssignale (E ₁ -E _N) zu erhalten, die Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen 1-Bit Eingangssignals (E ₁ -E _N) innerhalb eines vorbestimmten Zeitraums zu zählen, die Anzahl der gezählten Zustände mit einem vorbestimmten Schwellwert der jeweiligen Vergleichsschaltung (3) zu vergleichen und bei Überschreiten des Schwellwerts auf die erfolgte Erkennung des zu erkennenden Musters hinzuwei-		
22	zweiter Binärbaum- zweig; zweiter innerer Knoten des Binärbaums; zweiter Nested Branch			
3	Vergleichsschaltung; Segment			
30	erstes Oder-Gatter	sen.		
31	erstes Und-Gatter	2. Neuromorpher Mu	sterdetektor (2) nach An-	
32	(4-fach) Multiplexer	spruch 1, dadurch gekennzeichnet, dass die ei-		
33	erstes (2-bit SIPO-) Schieberegister	ne Vergleichsschaltung (3) der anderen Vergleichsschaltung (3) erstrangig untergeordnet ist, wobei die übergeordnete Vergleichsschaltung (3) ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der übergeordneten Vergleichsschaltung (3) überschritten und zeitgleich von der erstrangig untergeschaltung (3) ausgebildet in der erstrangig untergeschaltung (3) ausgebildet ist, nur dann auf die erfolgte Erstrangig untergeschaltung (3) ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden in der erstrangig untergeschaltung (3) ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der übergeschaltung (3) ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der übergeschaltung (3) überschriften und zeitgleich von der erstrangig untergeschaltung (4) ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der übergeschaltung (4) ausgebildet ist, nur der erstrangig untergeschaltung (4) ausgebilde		
34	zweites Und-Gatter			
35	drittes Und-Gatter			
36	(1-Bit) Flipflop			
37	zweites (N-bit SIPO-) Schieberegister	ordneten Vergleichsschaltung (3) auf die erfolgte Er- kennung des zu erkennenden Musters hingewiesen wird.		
38	zweites Oder-Gatter	2 Nouremember Mu	otordotoktor (2) nach An	
4	Schiebefensterdetektor; Slider	3. Neuromorpher Musterdetektor (2) nach Anspruch 2, gekennzeichnet durch wenigstens eine weitere Vergleichsschaltung (3), welche parallel zu der untergeordneten Vergleichsschaltung (3) angeordnet ist, wobei die übergeordnete Vergleichsschaltung (3) ausgebildet ist, nur dann auf die erfolgte Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der übergeordneten Vergleichsschaltung (3) überschritten und zeitgleich von den erstrangig untergeordneten Vergleichsschaltungen (3) jeweils auf die erfolgte Erkennung des zu erkennenden Musters hingewiesen wird.		
40	erstes Und-Gatter			
41	erstes (N-bit SIPO-) Schieberegister			
42	zweites bidirektionales (M-bit SIPO-) Schiebere- gister			
43	((M+1)-fach) Multiplexer			
44	drittes (K-bit SIPO-) Schieberegister			
5	Zeitmultiplexer; Time Multiplexer	4. Neuromorpher Musterdetektor (2) nach einem der Ansprüche 2 oder 3, gekennzeichnet durch wenigstens eine weitere Vergleichsschaltung (3), welche zweitrangig untergeordnet zu der erstrangig untergeordneten Vergleichsschaltung (3) angeordnet ist, wobei die erstrangig untergeordnete Vergleichsschaltung (3) ausgebildet ist, nur dann auf die erfolg-		
50	Flipflops			
51	Und-Gatter			

- te Erkennung des zu erkennenden Musters hinzuweisen, falls der Schwellwert der erstrangig untergeordneten Vergleichsschaltung (3) überschritten und zeitgleich von der zweitrangig untergeordneten Vergleichsschaltung (3) auf die erfolgte Erkennung des zu erkennenden Musters hingewiesen wird.
- 5. Neuromorpher Musterdetektor (2) nach Anspruch 3 oder 4, **dadurch gekennzeichnet**, dass die wenigstens drei Vergleichsschaltungen (3) einen Binärbaum mit wenigstens zwei Ebenen bilden.
- 6. Neuromorpher Musterdetektor (2) nach einem der vorangehenden Ansprüche, **dadurch gekennzeichnet**, dass die Vergleichsschaltungen (3) identisch ausgebildet sind.
- 7. Neuromorpher Musterdetektor (2) nach einem der vorangehenden Ansprüche, **dadurch gekennzeichnet**, dass bei Überschreiten des Schwellwerts ein 1-Bit Ausgangssignal (P₁-P_k) der jeweiligen Vergleichsschaltung (3) auf den "high"-Zustand, ansonsten auf den "low"-Zustand, gesetzt wird, oder umgekehrt.
- 8. Neuromorpher Musterdetektor (2) nach Anspruch 7, **dadurch gekennzeichnet**, dass die Vergleichsschaltungen (3) ausgebildet sind, jeweils ein 1-Bit Steuersignal (I_1 - I_N) zu erhalten und in Reaktion auf einen "high"-Zustand oder auf einen "low"-Zustand des jeweiligen 1-Bit Steuersignals (I_1 - I_N) das 1-Bit Ausgangssignal (P_1 - P_k) der jeweiligen Vergleichsschaltung (3) auf den "low"-Zustand zu setzen.
- 9. Neuromorpher Musterdetektor (2) nach einem der vorangehenden Ansprüche, dadurch gekennzeichnet, dass der vorbestimmte Schwellwert der Anzahl der Zustände der jeweiligen Vergleichsschaltung (3) vorgibt, wann das zu erkennende Muster als erkannt angesehen wird.
- 10. Neuromorpher Musterdetektor (2) nach einem der vorangehenden Ansprüche, **dadurch gekennzeichnet**, dass die Vergleichsschaltungen (3) jeweils einen Schiebefensterdetektor (4) aufweisen, welcher jeweils ausgebildet ist, das jeweilige 1-Bit Eingangssignal (E_1 - E_N) zu erhalten und die Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen 1-Bit Eingangssignals (E_1 - E_N) innerhalb des vorbestimmten Zeitraums zu zählen.
- 11. Neuromorpher Musterdetektor (2) nach Anspruch 10, **dadurch gekennzeichnet**, dass das Zählen der Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen 1-Bit Eingangssignals (E_1 - E_N) innerhalb des vorbestimmten Zeitraums mittels eines bidirektionalen Schieberegisters (42) des jeweiligen Schiebefensterdetektors (4) erfolgt.

- 12. Neuromorpher Musterdetektor (2) nach einem der vorangehenden Ansprüche, dadurch gekennzeichnet, dass
- die Vergleichsschaltungen (3), vorzugsweise deren Schiebefensterdetektor 4, jeweils ein Taktsignal CLK_{SPIKE} zur Steuerung der Verarbeitung der Pulse und ein Taktsignal CLK_{PLT} zur Steuerung der Länge der Plateaus erhalten,

wobei die beiden Taktsignale ${\rm CLK_{SPIKE}}$ und ${\rm CLK_{PLT}}$ unterschiedlich sind.

13. Neuromorphe Schaltkreisanordnung (1) mit einer Mehrzahl von neuromorphen Musterdetektoren (2) nach einem der vorangehenden Ansprüche, wobei jeder neuromorphe Musterdetektor (2) ausgebildet ist,

das gleiche 1-Bit Eingangssignal (E_1 - E_N) zu erhalten, ein unterschiedliches 1-Bit Zufallszahlensignal ($M_{1,1}$ - $M_{K,N}$) zu erhalten,

das jeweilige 1-Bit Eingangssignal (E_1 - E_N) mit dem entsprechenden 1-Bit Zufallszahlensignal ($M_{1,1}$ - $M_{K,N}$) zu verändern, und

die Anzahl der "high"-Zustände oder der "low"-Zustände des jeweiligen veränderten 1-Bit Eingangssignals (E_1 - E_N) innerhalb eines vorbestimmten Zeitraums zu zählen.

- 14. Neuromorphe Schaltkreisanordnung (1) nach Anspruch 13, **dadurch gekennzeichnet**, dass wenigstens eine Vergleichsschaltung (3), vorzugsweise alle Vergleichsschaltungen (3) jeweils, ein Und-Gatter (35) aufweist, welches ausgebildet ist, das jeweilige 1-Bit Eingangssignal (E₁-E_N) und das entsprechende 1-Bit Zufallszahlensignal (M_{1,1}-M_{K,N}) zu kombinieren.
- 15. Neuromorphe Schaltkreisanordnung (1) nach Anspruch 13 oder 14, **dadurch gekennzeichnet**, dass

die neuromorphe Schaltkreisanordnung (1) ausgebildet ist.

die Anzahl der 1-Bit Ausgangssignale (P_1 - P_k) der jeweiligen Vergleichsschaltung (3), welche zeitgleich im "high"-Zustand oder im "low"-Zustand sind, zu erfassen und

aus dem Verhältnis der Anzahl von 1-Bit Ausgangssignalen (P_1 - P_k) im "high"-Zustand oder im "low"-Zustand und der Anzahl der neuromorphen Musterdetektoren (2) einen Grad (W) der Übereinstimmung zwischen 1-Bit Eingangssignal (E_1 - E_N) und zu erkennendem Muster zu bestimmen.

16. Neuromorphe Schaltkreisanordnung (1) nach einem der Ansprüche 13 bis 15, **dadurch gekennzeichnet**, dass wenigstens eine Vergleichsschaltung (3), vorzugsweise alle Vergleichsschaltungen (3) jeweils, einen Zeitmultiplexer (5) aufweist, welcher ausgebildet ist, parallele Ausgangssignale (P₁-P_K) der neuromorphen Musterdetektoren (2) zu einem 1-Bit-

Ausgangssignal (O) der neuromorphen Schaltkreisanordnung (1) zusammenzuführen.

Es folgen 7 Seiten Zeichnungen

Anhängende Zeichnungen

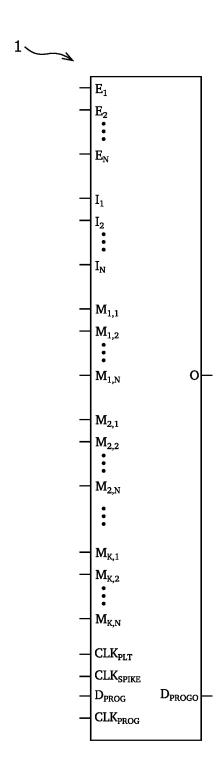


FIG. 1

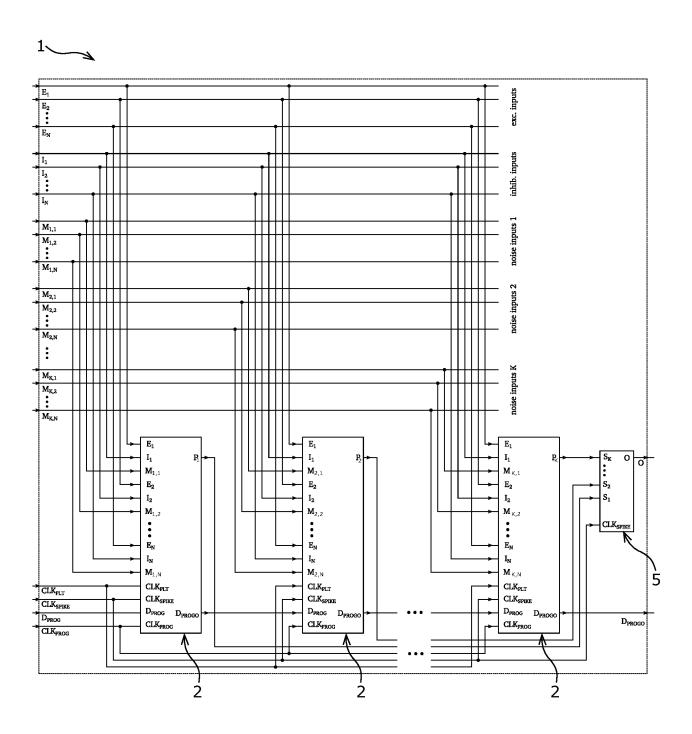


FIG. 2

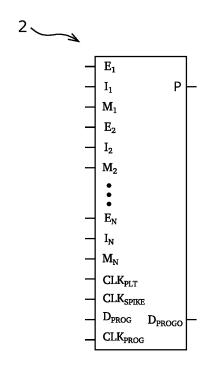
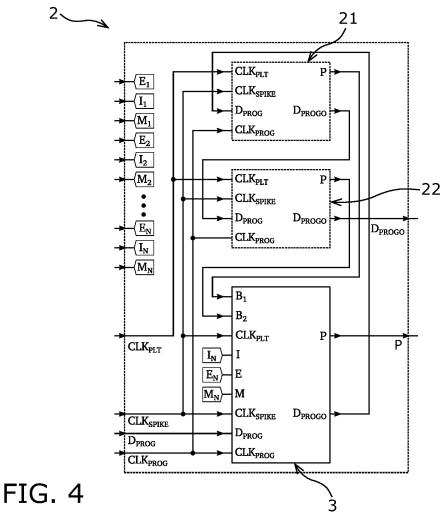


FIG. 3



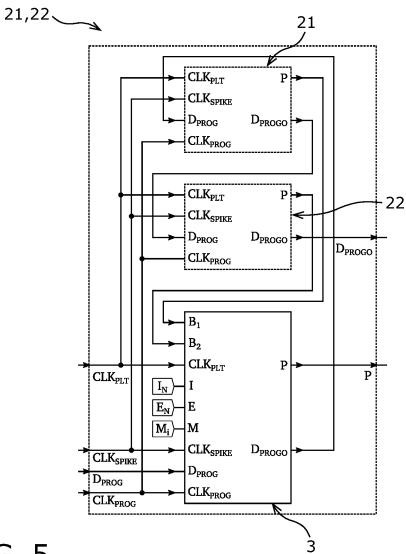


FIG. 5

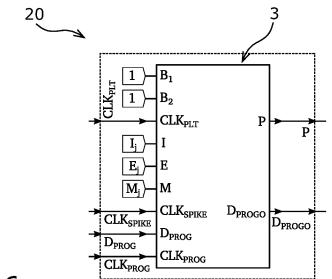


FIG. 6



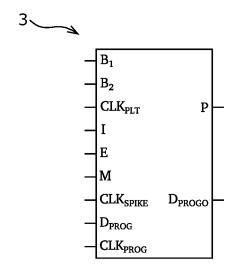


FIG. 7

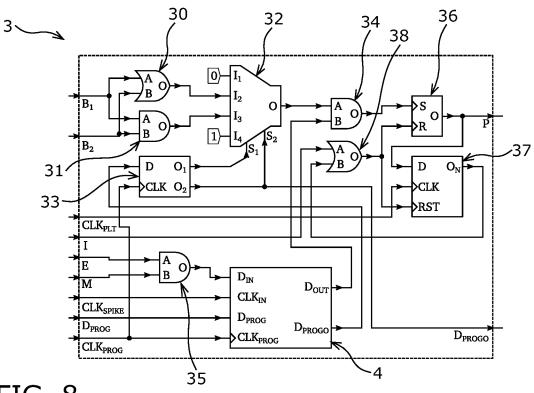


FIG. 8



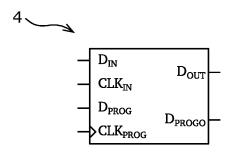


FIG. 9

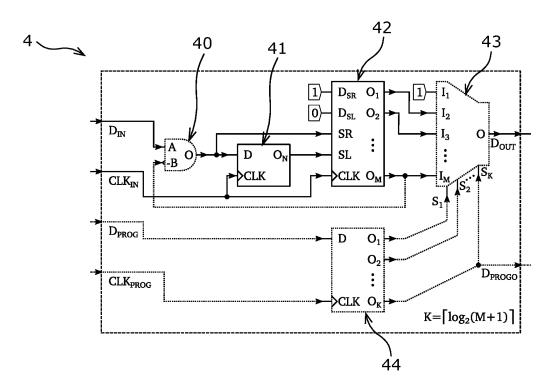


FIG. 10

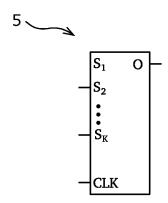


FIG. 11

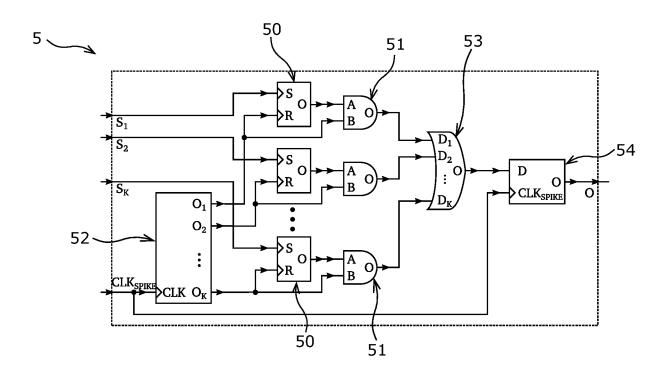


FIG. 12