

Master of Science in Engineering with Specialisation in Data Science

Master Thesis

On Improving Pattern Recognition in Deep Learning Systems through Self-Organization

**Incorporating Findings from Neuroscience about Natural Intelligence into
Modern Deep Learning Architectures**

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September 7, 2022

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Dedicated to the dedicated.

– Pascal Sager

*"Max Planck said, 'Science progresses one funeral at a time.'
The future depends on some graduate student who is deeply
suspicious of everything I have said."*

– Geoffrey Hinton, University of Toronto, 2017.

Zusammenfassung

Abstract

Preface

This thesis is the last hurdle before I will hold the title “Master of Science”. To me, science means the systematic analysis of the real or virtual world through observations and experiments as well as the further development of existing technology. I am lucky enough to be able to apply the knowledge and methodologies I learned during my studies to research projects at the Centre for Artificial Intelligence (CAI) of the Zurich University of Applied Sciences (ZHAW). I was mentored and supported during my studies by Prof. Dr. Thilo Stadelmann. He uses to say (also in accordance with his [blog-post](#)) “Great methodology delivers great theses”. It is always desirable to have an excellent outcome such as a system that can execute a task and thereby achieves or even overcomes state-of-the-art performance. However, in my opinion, it is equally or even more important to reason why and how something works, to justify choices, and to show limitations. I wrote my Thesis with these thoughts in mind and hope that the readers are able to follow my reasoning.

In the Introduction section, the work is motivated and in the subsequent chapter the fundamentals of Deep Learning and its limitations is described. Afterwards, it is motivated why methodologies inspired by neuroscience could overcome these limitations. This thesis aims at a target audience with a background in Deep Learning. Consequently, the concepts of Deep Learning are only roughly described. Since Neurocomputing may be rather unknown to the target audience, a more extensive overview about this field is given.

I would like to thank a couple of colleagues and friends. First I think of my mentor Prof. Dr. Thilo Stadelmann who got me excited about AI years ago and later introduced me to research. He always encouraged creative ideas and helped to link different research areas to address problems with methodologies from other fields. Thanks to his support, help, and guidance, I have grown personally as well as professionally. Further thanks go to Dr. Jan Deriu. Especially at the beginning of my thesis, he steered my thoughts in one direction. His unconventional thinking has led to the questioning of many methods that have stood the test of time for decades (this was also the inspiration for Geoffrey Hinton’s quote on the page before, although I wouldn’t presume to say that this thesis will change the future). To Prof. Dr. Christoph von der Malsburg for his seemingly endless patience in introducing me to neuroscience. He can build the bridge between the two diverging fields of Deep Learning and Neuroscience, inspires to think outside the box.

The biggest thanks, however, goes to my family, who made this journey possible for me. My parents, who supported and encouraged me in every way. My younger brother who inspired me to study. My wife and son, who have been understanding and supportive and have always been the perfect counterbalance to the daily routine of studying. Without the support of my family, I would never have been able to embark on this academic path.

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Introduction

1

Mankind has always tried to simplify its life through technological progress. Thousands of years before Christ the wheel was invented, later the pulley, the printing press, and the steam locomotive. At the beginning of the 19th century, the first generators were built to produce electricity, and in 1835, the first light bulb was invented by James Bowman Lindsay¹. In the course of time more and more complex electrical circuits were produced and finally in 1941 the first digital calculating machine, the computer was developed. This calculating machine is able to perform various arithmetic operations on the basis of commands. Through the development of transistors, more and more complex computers could be produced, which are able to execute various tasks. The definition of these tasks are captured in software. The computer has been so successful in fact that it has spawned its own scientific discipline, Computer Science. Nowadays it is impossible to imagine life without computers: almost every household and company owns computers. Computers facilitate various tasks, from communication to the acquisition of knowledge. For all these tasks, software developers have created appropriate tools. Software development is the process of writing a script with a programming language to specify how the computer should behave for a given input. Simply put: A software program tells the computer what to do if the user enters a command. This works very well if the tasks is clearly defined and can be described precisely. However, there exists tasks which are almost impossible to program. For example, writing a script that detects cats in images is almost impossible because we cannot describe how a cat looks like².

Therefore, computer scientists came up with the idea to not just program such tasks but to let the computer learn them instead. Machine learning (ML) are algorithms that are able to learn and adapt without following explicit instructions such as program code. Instead, they use statistical models to analyse data, to find patterns in the data and to draw inferences out of it. Machine learning has become an indispensable part of our everyday lives. For example, we use it for machine translation, transport and logistics organization, product recommendations, fraud detection, self-driving cars, unlocking smartphones, improving video games, speech recognition, and much more. A sub-branch of Machine Learning, namely Deep Learning, has made waves in the last decade. Breakthroughs are being made with this technology are made almost monthly and allows us to execute more and more tasks.

However, such DL system are usually good at one ore a few closely related tasks. In fact, they have they have some crucial flaws by definition (c.f. Section Section 2.2) which cannot be resolved for sure in the current DL framework. One of the Godfathers of Deep Learning is the Turing Award winner Geoffrey Hinton. Especially his contribution to back propagation (c.f. Section Section 2.1) has shaped the field. Back propagation is *the* core learning algorithm of DL systems and was developed in 1986. More than

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1: and not, as so often falsely claimed, by Thomas Alva Edison

2: at least not on a pixel-level basis so that we can simply compare a given image with our description

[1]: Inc. (2017)

30 years later, in 2017, Hinton says that he is “deeply suspicious” about back propagation and in his view we have to “throw it all away and start again” to improve current systems fundamentally [1]. Considering what DL systems have achieved, this seems a bit extreme. However, it also shows that the current learning algorithm of such systems has serious flaws.

3: there exist many additional (sub-)fields studying the brain such as cognitive science, cognitive psychology, neurology, and neuropsychology

Mankind is often inspired by nature when it comes to developing novel systems. The development of artificial intelligence (AI) and Deep Learning has been strongly influenced by, according to our perception, one of (if not the) most intelligent systems on planet Earth, the human brain. The brain is studied by the scientific field of neuroscience³. From neuroscience or related fields come various insights on how the human nervous system and especially the brain works. Much of this knowledge has been gained through observations and experiments on humans or other living creatures. Often these findings are only what can be measured from the outside, the real core, the mechanism that causes intelligence, remains unknown.

While Deep Learning is clearly inspired by the insights of Neuroscience, the two fields have little in common in their present form. The implementation of neuroscientific findings has emerged as a subfield in its own right, called Neurocomputing (c.f. Section 2.4). In general, neurocomputing is closer related to the functionality of the human brain than Deep Learning. Neurocomputing has produced many promising algorithms that can overcome some weaknesses faced by deep learning systems. Although neurocomputing has also achieved significant breakthroughs, most systems used in everyday life are still based on the principle of Deep Learning.

TODO: deep learning or Deep Learning -> make uniform

1.0.1 Motivation

Despite the fact that Deep Learning has achieved incredible performance on a variety of tasks it is still questionable whether the current methodology is enough to achieve real (or at least more advanced) intelligence (c.f. Section 2.2). Many of the current limitations are tackled by approaches from the field of neurocomputing (c.f. Section 2.4). However, even though algorithms from the field of neurocomputing have interesting properties, it usually does not perform as good as Deep Learning and often works on specific or small data only.

Neuroscience provides a source of inspiration for AI algorithms, independent of mathematical models. So far, neuroscientists have studied the building blocks of the brain, their responsibilities in the overall process as well as how they interconnect and communicate. The “algorithm” that makes the brain intelligent remains unknown. Furthermore, it is not known which observed functionality in the brain is really part of this algorithm and which are just consequences of it. For example, it is hard to tell if the dynamics in the human brain are necessary to achieve intelligence or if this is just nature’s way of implementing it.

Renowned scientists [2] put forward the hypothesis that the core of natural intelligence lies within the self-organization of the brain. Self-organization is the process by which individual units organize their global behavior by local interactions amongst themselves. There is no central control unit that orchestrates the units. In the context of the brain, self-organization primarily affects how neurons are interconnected. The brains of living beings already have certain structures from birth, which are specified by the DNA. In the course of time, these structures change. For example, neurons or connections between neurons may be added or removed. Thus, the brain adapts to its environment over time.

[2]: Malsburg et al. (2022)

Such a mechanism has not been successfully implemented in deep learning nor neurocomputing systems. In deep learning system, the architecture is usually predefined and only the parameters (i.e. weight and biases, c.f. Section 2.1) are updated. In neurocomputing, on the other hand, the theory behind self-organization is often associated with Hebbian Learning (c.f. Section 2.4.1). this dynamic learning of connections is often associated with Hebbian Learning (c.f. Section 2.4.1). The theory behind this learning rule can be summarized as “cells that fire together wire together”[3]. However, Hebbian learning is usually rather used to update the weight between connections instead of to add or to remove connections. Connections are therefore not removed or added but can be “turned off” and “turned on” if the activation function of a neuron uses a threshold below which signals are no longer transmitted⁴. In this case, the architecture of the network is to a large part still predefined and self-organization can for example not be used to add additional layers.

[3]: Löwel et al. (1992)

4: if the weights get small enough, the threshold limit cannot be reached and the neurons output is 0

This thesis aims to combine the strengths of deep learning and neurocomputing with recent findings of neuroscience. Especially the hypothesis that self-organization is key for natural intelligence is incorporated in the learning process of modern artificial neural networks. While in classical Deep Learning the architecture is predefined, in this thesis we attempt to learn not only the parameters but also the architecture itself. The architecture is built by self-organization. This means that local communication between the building blocks defined by an initial architecture takes place and that new connections can be created or existing connections can be removed. The challenge is how this re-wiring of the connections can be learned. It seems certain that the existing learning algorithm back-propagation must either be extended or replaced. However, proposing a concrete algorithm based on the rather abstract findings from Neuroscience is part of this thesis.

1.0.2 Organization of Thesis

The remainder of the thesis is organized as follows: In chapter 2 we present the fundamentals necessary to understand this thesis. We provide an overview about how deep learning works and what the most common research areas in neurocomputing are. Chapter 3 presents the work related to this thesis.

TODO....

TODO: Input Thilo -> mache den Bezug von diesem Teil auf die schlussendliche Arbeit viel deutlicher -> was wird wie gebraucht?

Machine Learning uses mathematical functions to map an input to an output. These functions usually extract patterns from the input data to build a relationship between input and output. The term Machine Learning stems from the fact that we use *machines* to correlate the input and the output to a function (i.e. to *learn* a function) during a training period. A sub-branch of Machine Learning is Deep Learning (DL). DL algorithms are able to learn hidden patterns within data to make predictions. They benefit from the accelerated computing power and big data made available in the last decade. Deep Learning is considered state-of-the-art for many learning tasks especially for high-dimensional data. Typical high-dimensional data are texts, audio recordings, 2D as well as 3D images, and videos. Deep Learning models use artificial neural networks to learn the mapping function between given input and output data. In the following, the fundamentals of Deep Learning is explained. Only those aspects that are relevant for the understanding of the rest of the thesis are discussed.

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2.1 Artificial Neural Networks

The idea for artificial neural networks (ANN) stems from biology and aims to capture the interaction of brain cells (neurons) with a mathematical model. A first model for a neuron was proposed by McCulloch and Pitts in 1943 [4]. Similar to how a neuron of the human brain transmits electrical impulses through the nervous system, the artificial neuron of McCulloch and Pitts receives multiple input signals and transforms them into a output signal. A neuron takes an input vector $\mathbf{x} = (x_1, \dots, x_n)$ where $x_i \in \{0, 1\}$ and maps it to an output $\hat{y} \in \{0, 1\}$. The mapping from the input to the output is done by using an aggregation function g that sums up the input vector \mathbf{x} and a activation function f that outputs 1 if the output of g is greater than a threshold θ and 0 otherwise.

[4]: McCulloch et al. (1943)

$$g(\mathbf{x}) = g(x_1, \dots, x_n) = \sum_{i=1}^n x_i \quad (2.1)$$

$$\hat{y} = f(g(\mathbf{x})) = \begin{cases} 1, & \text{if } g(\mathbf{x}) \geq \theta \\ 0, & \text{otherwise} \end{cases} \quad (2.2)$$

In 1958, Rosenblatt [5] developed the Perceptron which works with real numbers as input. The input vector $\mathbf{x} = (x_1, \dots, x_n)$ where $x_i \in \mathbb{R}^n$ is multiplied with a weight vector $\mathbf{w} = (w_1, \dots, w_n)$ where $w_i \in \mathbb{R}^n$ with the same length.

[5]: Rosenblatt (1958)

$$g(\mathbf{x}) = g(x_1, \dots, x_n) = \sum_{i=1}^n w_i \cdot x_i \quad (2.3)$$

The output $\hat{y} \in \{0, 1\}$ is similar to the McCulloch and Pitts neuron 1 if the aggregated value is greater than a threshold θ and 0 otherwise as described in equation ?? The equations (2.3) and (3.1) can be rewritten as

$$\hat{y} = f(g(\mathbf{x})) = \begin{cases} 1, & \text{if } \sum_{i=1}^n w_i \cdot x_i - \theta \geq 0 \\ 0, & \text{otherwise} \end{cases} \quad (2.4)$$

Later, the step-function f was replaced with other functions so that the output could also be a real number $\hat{y} \in \mathbb{R}$. Often used functions are

$$\begin{aligned} \text{Sigmoid: } \sigma(z) &= \frac{1}{1 + e^{-z}} \\ \text{Rectified linear unit (ReLU): } (z)^+ &= \max(0, z) \\ \text{Hyperbolic tangent (tanh): } \tanh(z) &= \frac{e^z - e^{-z}}{e^z + e^{-z}} \end{aligned} \quad (2.5)$$

By convention, instead of using the threshold $-\theta$ often a bias b is used which leads to:

$$\begin{aligned} z = g(\mathbf{x}) &= \mathbf{w} \cdot \mathbf{x} = \sum_{i=1}^n w_i \cdot x_i + b \\ \hat{y} &= f(z) \end{aligned} \quad (2.6)$$

However, the brain consists of multiple neurons which are connected through synapses. Therefore, ANNs consist not only of one neuron but combine multiple neurons in a network. These neurons are organized into layers. A shallow neural network consists of one hidden layer in which the input \mathbf{x} is fed to calculate the output $\hat{\mathbf{y}}$ ¹. The universal approximation theorem of Cybenko [6] proves that a shallow network with enough neurons can approximate any mapping function between inputs and outputs. However, very complex mapping functions may need too many hidden neurons. The neurons of the hidden layer extract features in the input space. Since only one layer is used, the features cannot be hierarchically organized and become complex enough.

A multi-layer perceptron (MLP), on the other hand, consists of multiple layers. The different layers extract increasingly complex features. In a MLP the input \mathbf{x} is fed into the first layer, each subsequent layer l gets the output of the previous layer $l - 1$ as input. In the following, we describe the mathematical model for fully connected layers, where all neurons of a layer are connected with the subsequent layer². For a MLP with m layers, we define the output of the aggregation g as $\mathbf{z}^{[l]}$ and the output the activation function as $\mathbf{a}^{[l]}$ for layer l . Furthermore, we use $\mathbf{w}^{[l]}$ for the weight vector and $b^{[l]}$ for the bias of layer l . Thus, the mathematical model of a MLP is defined as

1: $\hat{\mathbf{y}}$ has become a vector since multiple neurons produce multiple output values
[6] Cybenko (1989)

2: many modern network architectures are not fully connected and can have either missing or recurrent connections

$$\begin{aligned} z^{[l]} &= w^{[l]} a^{[l-1]} + b^{[l]} \\ a^{[l]} &= f(z^{[l]}) \end{aligned} \quad (2.7)$$

Since the input is fed into the first layer and the output is the result from the last layer $x = a^{[0]}$ and $\hat{y} = a^{[m]}$ holds true.

So far, only the forward-pass which is used to calculate the output \hat{y} was discussed. However, the model output \hat{y} will only be close to the target output y if the weights $w^{[l]}$ and biases $b^{[l]}$ are properly defined. These parameters are learned during a training period. The training can take place in a supervised, semi-supervised, self-supervised, unsupervised, or reinforcement learning based manner. In supervised learning, the output of the model \hat{y} for a given input x is compared to manually created target outputs y . Unsupervised learning, on the other hand, tries to find patterns in the input x and to cluster the samples into meaningful groups without using target labels. Semi-supervised learning is a hybrid approach of the the aforementioned principles that combines a small amount of labelled data with a large amount of unlabelled data. In self-supervised learning, the target outputs y are directly derived from the input data x (e.g. predict a masked part of the input x). Lastly, reinforcement learning algorithms aim to maximize a reward that they become from an environment based on some action they executed.

These learning principle have in common that a loss function \mathcal{L} can calculate a loss value based on the model output \hat{y} . For example, the mean square error (MSE) can be used for regression problems or the negative log-likelihood for classification problems. The chosen loss function is minimized iteratively with stochastic gradient descent (SGD)³ until the network converges. The idea behind stochastic gradient descent is to make use of the fact that the negative gradient of the loss value points to the direction of the steepest descent (i.e. in the direction where the loss gets smaller). SGD therefore updates the network parameters by taking a step of size η in the direction of their negative gradient

$$\begin{aligned} \Delta w^{[l]} &= -\eta \nabla_{w^{[l]}} \mathcal{L} \\ \Delta b^{[l]} &= -\eta \nabla_{b^{[l]}} \mathcal{L} \end{aligned} \quad (2.8)$$

The gradients of the weights $w^{[l]}$ and biases $b^{[l]}$ can efficiently be calculated with an algorithm called backpropagation [8], which is just a smart implementation of the chain rule⁴.

TODO: Describe CNN, Transformer, ... etc.???

3: There exist also other optimizer methods such as SGD with momentum, RM-Sprop, or Adam [7]

[8]: Rumelhart et al. (1986)

4: While a detailed discussion on backpropagation is out of scope for this thesis, we refer interested reader to the Deep Learning course by Andrew Ng [9]

2.2 Limitations of Deep Learning

The rise of Deep Learning over the past decade has only been possible because of major technological advances in hardware. Without the computational resources and the storage capacity of the systems created in the last decades no system could run today's algorithm. Moreover, much of the progress of recent years was possible due to the improved hardware. Moore's law [10] states that the number of transistors in a

dense integrated circuit doubles about every two years and is the only known physical process following an exponential curve. An analysis by OpenAI shows that since 2012 the amount of compute has even increasing exponentially with a doubling time of 3.4 months [11]. However, the exponential increase seems to come to an end since the size of transistors hit physical limitations. It is assumed that Moore's law will end by around 2025 [12]. Besides the progress in the field and the development of new technology, Deep Learning models also became better because the number of parameters and the size of datasets grew exponentially. Even the growth in the last five years is astonishing. While the state-of-the-art language model from 2018 [13] had around 94M parameters, the state-of-the-art in 2020 [14] already had 175B parameters. Training such a model on a single V100 GPU would take about 355 years and cost about 4.6M dollars [15]. A recent language model from Microsoft and Nvidia [16] even has 530B parameters. Only a few institutions with massive resources are able to train such big models. In general, inference on low-budget hardware such as smartphones or embedded hardware becomes prohibitive with the growing size of deep networks. Even though there exist techniques to shrink the model size after training such as quantization [17], model pruning [18], or model distillation [19] it is questionable if making models bigger is the best way to develop intelligent systems.

Another major issue of Deep Learning systems is that they suffer from catastrophic forgetting. If a model is trained on a specific task and afterwards trained (or fine-tuned) on another task, the model suffers a "catastrophic" drop in performance over the first task. The reason for this effect is that the model during training on the second task adjusts the parameters learned during the first task and therefore "forgets" the learned mapping functions. Just mixing all datasets or to learn all tasks in parallel in a current multi-task setup [20] doesn't seem feasible to achieve some kind of general intelligence as this involves too many different unrelated tasks. Catastrophic forgetting is also caused by the fact that learning is mostly done offline⁵. Online learning [21] and lifelong learning [22] are currently hot research topics. However, these methods have not yet been established.

Furthermore, there exists problems which may cannot be solved with the current principles used for Deep Learning. First of all, it is questionable if Deep Learning models can achieve *real* generalization⁶. With enough data, can achieve generalization in the sense that the model can interpolate within the data distribution. However, deep learning models fail to extrapolate. For example, convolutional neural networks (CNNs) do not generalize to different viewpoints unless they are added to the training data [23].

Second, Deep Learning is not able to learn abstract relationships in a few trials but requires many samples of it and is thus data hungry.⁷ Marcus Gary [24] argues that if he tells that a "schmister" is a sister over the age of 10 but under the age of 21, humans can immediately infer whether they or their best friends have any "schmister". However, modern DL systems lacks a mechanism for learning abstractions through explicit, verbal definition and require thousands or even more training samples.

Third, no DL model has been able to demonstrate causal reasoning in

[12]: Kumar (2015)

[13]: Peters et al. (2018)

[14]: Brown et al. (2020)

[16]: Shoenybi et al. (2020)

5: Offline in this context means that the model parameters are not adapted after training during inference time

6: Generalization refers to the ability of the model to adapt properly to previously unseen data from the same distribution

[23]: Madan et al. (2022)

7: Delme!!!

[24]: Marcus (2018)

a generic way. Deep Learning models find correlations between the inputs and the outputs, but not the causation. Other AI approaches such as hierarchical Bayesian computing or probabilistic graphical models are better at causal reasoning but cannot be well combined with Deep Learning models.

Lastly, Deep Learning models are to some extent too isolated since they have no embodiment and cannot interact with the world. For example, the human body provides needs, goals, emotions, and gut feeling⁸. In current Deep Learning systems emotions are totally absent and the goals are set externally. Deep Reinforcement Learning can be considered as a first step in the direction of dissolving this isolation, as they interact with a virtual environment. AI systems that interact with the real world do not work well so far. Moravec's paradox [25] states that "it is comparatively easy to make computers exhibit adult level performance on intelligence tests or playing checkers, and difficult or impossible to give them the skills of a one-year-old when it comes to perception and mobility".

8: one could argue that the body is therefore even a co-processor of the brain

[25]: Moravec (1995)

2.3 Biological Learning

The human brain comprises many interconnected areas processing everything in parallel. For example, Figure 2.1 illustrates the connections between different organizational units in the cerebral cortex which are responsible for vision. It can be seen that these areas are connected in a rather complex structure. Deep Learning architectures, on the other hand, are mostly unidirectional and the signal flows unidirectional from layer to layer⁹. However, the choice of the architecture influences how the model can learn the mapping function from input to output. It could be that the complex structure of our brain comprises an inductive bias which was learned over time through evolution.

9: Except for recurrent connections, skip-connections, or residual-connections

A learning system requires a mechanism that tells the system if something goes well or wrong so that it can learn from it. This is called the *credit assignment problem*. Backpropagation (c.f. Section ??) solves this problem by propagating the error backwards through the network. However, information flows in the brain only in one direction from the presynaptic neurons to the postsynaptic neurons. Therefore, backpropagation is not biologically plausible. Lillicrap et al. [27] shows that an additional set of random feedback weights is able to transmit useful gradients. Their work has reopened questions how the brain could process error signals and has dispelled some long-held assumptions about algorithmic constraints on learning.

[27]: Lillicrap et al. (2016)

Not only the structure of the network and the way how the feedback is calculated is different between biological learning and Deep Learning. Also the neurons themselves are different. While the artificial neuron doesn't have any dynamics (c.f. Equation (2.6)), biological neurons are highly dynamic: Biological neurons adapt their firing rate to constant inputs, they may continue firing after an input disappears, and can even fire when no input is active.

TODO: Add reference to reservoir computing

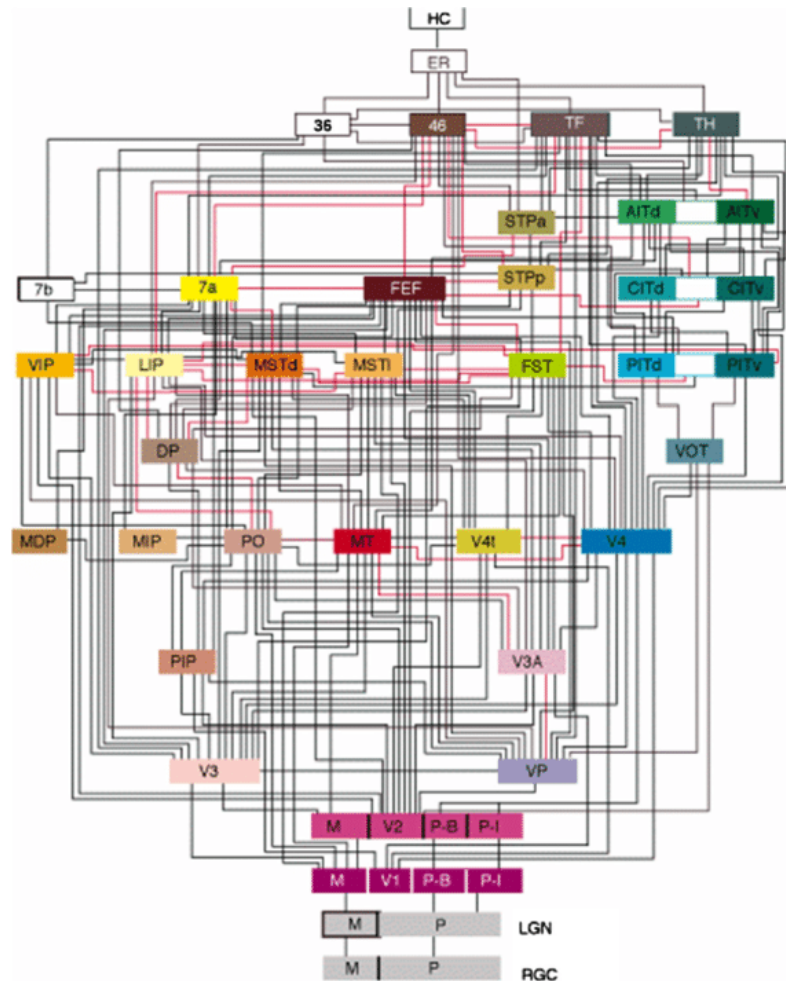


Figure 2.1: The organization of the visual system in the cerebral cortex. The image is from Felleman et al. [26].

Lastly, the neurons in the brain are self-organizing. This means that a group of elementary units such as neurons or a group of neurons perform similar rule of behavior on a sub-set of the available information. Such a system doesn't have a central supervision that orchestrates these units. Each unit applies similar deterministic functions to the information received. Two important principles of such systems are (i) localized learning which means that each unit adapt their behavior to the information they receive; and (ii) emergence which means that there is no explicit loss function that tells the system what to do.

2.4 Neurocomputing

2.4.1 Hebbian Learning

[28]: Hebb (1949)

Donald Hebb [28] describes how the connections between cells in the nervous system adapt as: "When an axon of cell A is near enough to excite a cell B and repeatedly or persistently takes part in firing it, some growth process or metabolic change takes place in one or both cells such that A's efficiency, as one of the cells firing B, is increased". This statement

is often simplified to the well-known phrase “Neurons that fire together wire together”.

Hebbian learning is based on this principle. The weight w_{ij} from neuron i to neuron j changes based on the pre-synaptic activity r_i of neuron i and post-synaptic activity r_j of neuron j

$$\Delta w_{ij} = \eta r_i r_j \quad (2.9)$$

where η is the learning rate. Thus, the weights between frequently co-activated neurons becomes strong which is called Hebbian plasticity.

In its original form, Hebbian learning had the problem that the connections could only become stronger but not weaker. Therefore, it is often extended based on the covariance of the activity between neurons. The covariance is positive if two neurons fire often together and negative if they do not often fire together. The following equation changes the weight relative to the covariance:

$$\Delta w_{ij} = \eta (r_i - \psi_i) \cdot (r_j - \psi_j) \quad (2.10)$$

where ψ_i and ψ_j are estimates of the expected pre- and post-synaptic activity¹⁰. The formulation above lacks of boundaries, i.e. the weights could grow to infinite. A simple solution is to enforce hard boundaries $w_{min} \leq w_{ij} \leq w_{max}$.

10: the expected activity can for example be estimated through a moving average function

Another solution to weaken the connections is given by the Bienenstock-Cooper-Monroe (BCM) learning rule which was introduced by Bienenstock et al. [29] and extended by Intrator and Cooper [30]. They propose a sliding threshold for long-term potentiation (LTP) or long-term depression (LTD) induction. When a pre-synaptic neuron fires and the post-synaptic neuron is in a lower activity state than the sliding threshold, it tends to undergo a LTD (i.e. the connection is weaken).

[29]: Bienenstock et al. (1982)

Around the same time, Oja [31] improved the learning rule of Equation (2.9) with an normalization term:

[31]: Oja (1982)

$$\Delta w_{ij} = \eta (r_i r_j - \alpha r_j^2 w_{ij}) \quad (2.11)$$

The parameter α is a constant value that determines the size of the norm of the weight vector. This update rule is also known as the Oja learning rule. Furthermore, he has found that a layer of multiple linear neurons converges to the first principle component of the input data. As all neurons only learn the first principle component, a network of multiple neurons in this setting seem not very useful. Differentiation between neurons can be achieved with several different methods. Two well known approaches are the winner-take-all competition (i.e. only the neuron with the most similar activity is selected for learning)¹¹ and a recurrent circuit that provides a competitive signal (i.e. the neurons compete with their neighbours to become active to learn).

11: in practice is k-winner-take-all often preferred where k instead of one neuron learns

It is known that independent neurons can encode more information and work better than dependent neurons [32]. Anti-Hebbian learning is a

[32]: Simoncelli et al. (2001)

method that adds a penalty for similarly active neurons and thus minimizes the linear dependency between neurons. Vogels et al. implemented this by switching the sign of the weight change [33].

[33]: Vogels et al. (2011)

There exists many further improvements for Hebbian learning which are not summarized in this thesis. For example, Joshi and Triesch [34] as well as Teichmann and Hamker [35] adapt the activation function of the neurons to enforce a certain activity distribution and to stabilize Hebbian learning even in multilayer neural networks.

[34]: Joshi et al. (2009)

[35]: Teichmann et al. (2015)

Similar to large parts of the brain, Hebbian learning is unsupervised and learns based on local information (i.e. neurons in close proximity). However, the brain is also largely recurrent and could guide neighbouring or preceding units. This assumption inspired supervised Hebbian learning. In supervised Hebbian learning, a subset of inputs which should evoke post-synaptic activity can be selected. Supervised Hebbian learning can be extended to top-down and bottom-up learning [36] which leads to a combination of supervised and unsupervised Hebbian learning.

[36]: Grossberg (1988)

2.4.2 Hopfield Networks

Hopfield networks [37] were introduced 1982 by J. Hopfield. They serve as associative (i.e. content-addressable) memory systems. Such systems are particularly useful to retrieve representations based on degraded or partial inputs. Auto-associative memories return for an input the most similar previously seen sample. A classical implementation of an auto-associative memory is the nearest neighbour algorithm [38]. This algorithm compares a given samples with the previously seen training data with a distance metric and returns the most similar sample¹². Memory networks [39] implement an auto-associative memory within the Deep Learning framework. Such networks convert an input x to a internal feature representation $I(x)$, update memories given the new input $m = G(m, I(x))$, and compute the output features $o = O(m, I(x))$. This process is applied during the training and inference phase. The only difference is that the parameters for the functions I , G , and O are only updated during training.

[37]: Hopfield (1982)

[38]: Fix et al. (1989)

12: or the k most similar samples in the case of the k -nearest neighbour (k-NN) algorithm

In a Hopfield network all neurons are connected, but there are no self-connections: $w_{ii} = 0$ where w_{ij} is the weight between neuron i and neuron j . Furthermore, the weights are symmetrical $w_{ij} = w_{ji}$. A Hopfield network in its original form works only with binary units. For consistency, this networks are called binary Hopfield networks in the following. The output of a neuron in a binary Hopfield network depends on the output of the other neurons within the network:

$$x_i = \sum_{j \neq i} w_{ij} y_j + b \quad (2.12)$$

$$y_i = \begin{cases} 1, & \text{if } x_i > 0 \\ -1, & \text{otherwise} \end{cases} \quad (2.13)$$

TODO in equation: check where to use y and where to use y with hat

Hopfield networks have their own dynamics and the output evolves over time. If the initial value y_i of a binary Hopfield network has a different sign than $\sum_{i \neq j} w_{ij} y_j + b$ the output will flip (i.e. change its sign). This will in turn influence all other neurons which may also flip. The term $y_i(\sum_{i \neq j} w_{ij} y_j + b)$ is negative if y_i is not equal to $\sum_{i \neq j} w_{ij} y_j + b$, otherwise it is positive. Since the neuron flips if the term $y_i(\sum_{i \neq j} w_{ij} y_j + b)$ is negative or stays the same if this term is positive, the change of this term can only be positive:

$$\Delta[y_i(\sum_{i \neq j} w_{ij} y_j + b)] \geq 0 \quad (2.14)$$

The negative sum of the term $y_i(\sum_{i \neq j} w_{ij} y_j + b)$ for the entire network is called the energy E of the network:

$$E(\mathbf{y}) = - \sum_i y_i \left(\sum_{j > i} w_{ji} y_j + b \right) \quad (2.15)$$

As we have shown in (2.14), the energy function $E(\mathbf{y})$ of the network can only decrease. Moreover, the energy function has a lower bound and thus the network reaches after a finite number of iterations a stable state. A stable pattern of the network (i.e. no neurons flip their sign) is a local minima of the energy function and is called a point attractor. A pattern is attracted by the closest stable pattern. Thus, the network can be used as associative memory because an input pattern can be matched to the closest stable pattern.

McEliece [40] has shown that a binary Hopfield network with N neurons has a capacity of $C = 0.138N$ (i.e. the number of patterns that can be stored. Hebbian learning (c.f. Section 2.4.1) can be used to store pattern in a Hopfield network. To store a pattern, the weights \mathbf{w} must be chosen in a way so that the desired local patterns $(\mathbf{y}^1, \dots, \mathbf{y}^P)$ are local minima of the energy function. By combining Hebbian learning with some smart mathematical transformations it can be shown that the weights can be directly learned with only one iteration over the training patterns¹³:

[40]: McEliece et al. (1987)

13: for the derivation of this equation refer to [37]

$$\mathbf{w} = \frac{1}{p} \sum_{k=1}^P \mathbf{y}^k \times (\mathbf{y}^k)^T - \mathbf{I} \quad (2.16)$$

where \mathbf{I} is the identity matrix. Later, Hopfield et al. [41] extended the binary Hopfield network so that it can either learn pattern during an awake cycle or forget patterns during a sleep cycle.

[41]: Hopfield et al. (1983)

One of the limiting factors of binary Hopfield networks is the capacity of $C = 0.138N$. The problems comes from the fact that the energy function is a quadratic function. More than three decades after the introduction of the binary Hopfield networks, Krotov and Hopfield [42] reformulated the energy function as a polynomial function to get polynomial capacity $C \approx N^{a-1}$ where a is the order of the function. Later, the energy function was reformulated as exponential function [43] and thus modern Hopfield networks have an exponential capacity of $C \approx 2^{\frac{N}{2}}$.

[42]: Krotov et al. (2016)

[43]: Demircigil et al. (2017)

[44]: Ramsauer et al. (2021)

The second limiting factor of binary Hopfield networks is that only binary patterns can be stored. Ramsauer et al. [44] extended the binary Hopfield network to continuous patterns by reformulating the energy function and the corresponding update rule. Continuous Hopfield networks can retrieve continuous patterns or even combination of several similar continuous patterns. The authors claim that a continuous Hopfield networks can replace fully-connected layers, attention layers, LSTM layers, support vector machines (SVM), and k-NN.

TODO: References for LSTM, attention, SVM, ...???

2.4.3 Spiking Neural Networks

14: the membrane potential is related to the electrical charge of the membrane of a biological neuron

[45]: Abbott (1999)

[46]: Brooke et al. (2003)

Biological neurons emit spikes. To transmit information, especially the firing rate (i.e. the number of spikes per second in Hz) and precise timing of the spikes are relevant. The amplitude and duration of the spike does not matter much. This behaviour has also been implemented in ANNs. So called Spiking neural networks (SNNs) incorporate the concept of time into their model. SNN do not transmit information in each forward-pass but rather transmit a signal when the membrane potential reaches a threshold value¹⁴. The neuron fires as soon as the threshold is reached and thereby influences the potential of other neurons. The most prominent model of a spiking neuron is the leaky integrate-and-fire (LIF) neuron [45]. The LIF neuron models the membrane potential with a differential equation. Incoming spikes can either increase or decrease the membrane potential. The membrane potential either decays over time or is reset to a lower value if the threshold value is reached and the neuron has fired. There exists different integrate-and-fire (IF) neurons models such as the Izhikevich quadratic IF [46] or the adaptive exponential IF [47]. While each of these model has different mathematical properties, the concept of a membrane potential that is increased or decreased through spikes from other neurons and decays over time or by emitting a spike is similar to the LIF.

[48]: Paugam-Moisy (2006)

Biological neurons have different dynamics. Some neurons fire regularly if they receive an input current, others slow down the firing rate over time or emit bursts of spikes. Modern models of spiking neurons can recreate this behaviour of biological neurons [48].

[49]: Bi et al. (2001)

The synaptic plasticity can be modeled with Hebbian learning (c.f. Section 2.4.1). The spike-timing dependent (STDP) plasticity rule [49] distinguishes the firing behaviour of pre-synaptic and post-synaptic neurons. If the pre-synaptic neuron fires before the post-synaptic neuron, the connection is strengthened, otherwise it is weakened.

[50]: Kheradpisheh et al. (2018)

For a long time, SNN only worked for very shallow networks. In 2018, Kheradpisheh et al. [50] has proposed a SNN based on the idea of CNNs called a deep spiking convolutional network. This network uses convolutional and pooling layers with IF neurons instead of classical artificial neurons and is trained with STDP. First, the image is fed into DoG cells. These cells apply the difference of Gaussians (DoG) feature enhancement algorithm. This algorithm subtracts a Gaussian blurred version of an image from the original image. By doing so, positive or negative contrast is detected in the input image. The higher the contrast is, the stronger is a cell activated and the earlier it emits a spike. Thus, the

order of the spikes depends on the order of the contrast. These spikes are forwarded to a convolutional layer. Deep spiking convolutional networks use two types of LIF neurons: On-center neurons fire when a bright area is surrounded by a darker area, off-center neurons do the opposite. Convolutional neurons emit a spike as soon as they detect their preferred visual feature¹⁵. Neurons that fire early perform the STDP update with a winner-takes-all mechanism. This means that the neurons within a layer compete with each other and those which fire earlier learn the input pattern. This prevents other neurons from firing and guarantees a sparse connection. Later convolutional layers detect more complex features by integrating input spikes from the previous layer. The features from the last convolutional layer are flattened and a Support Vector Machine is used to classify the features.

15: the location of the feature is not relevant as convolution layers are translation invariant

2.4.4 Reservoir Computing

As described in Section 2.3, biological neurons are highly dynamical while artificial neurons are not. Reservoir computing introduces such dynamics into an artificial network. Reservoir computing is an umbrella term for networks based on the concepts of Echo State Networks (ESN) [51] and Liquid State Machines (LSM) [52]. A reservoir is a fixed non-linear system that maps a input vector x to a higher dimensional computation space. After the input vector is mapped into computation space, a simple readout mechanism is trained to return the desired output based on the reservoir state. In principle, the system should be capable of any computation if it has a high enough complexity [53]. However, not every system is suited as reservoir. A good reservoir system distributes different inputs into different regions of the computation space [53].

[52]: Maass (2001) (2002)

[53]: Konkoli (2018)

A ESN is a set of sparsely connected recurrent neurons as visualized in Figure 2.2.

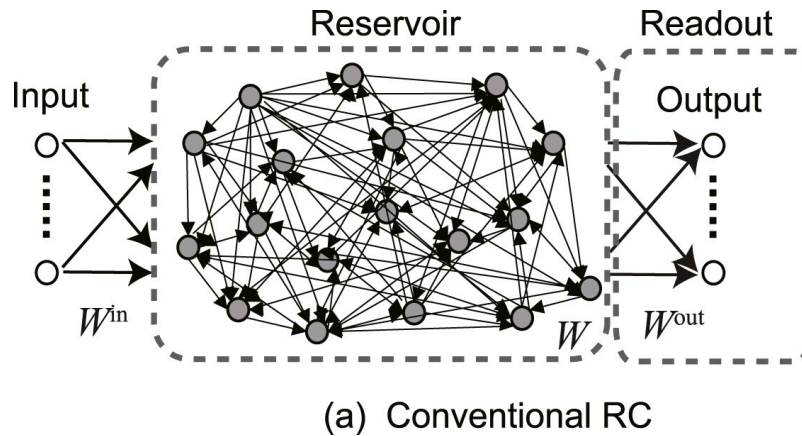


Figure 2.2: Structure of a Echo State Network. The image is from Tanaka et al. [54].

The reservoir consists of N nodes which are connected according a Erdős–Rényi graph model¹⁶ [55]. This graph model is represented by an adjacency matrix w ¹⁷ of size $N \times N$. The time varying input signal $x(t)$ is mapped to a sub-set of N/M graph nodes by multiplying it with $w_{in} \in \mathbb{R}^{N \times M}$ and the output by multiplying the reservoir state with $w_{out} \in \mathbb{R}^{M \times N}$. We refer interested reads to [56] to read more about the mathematical properties and how network is updated in detail.

[55]: Erdős et al. (1959)
The Erdős–Rényi model is a model for generating random graphs where all graphs on a fixed set of vertices and edges is equally likely

17: Figure 2.2 uses upper case letter W
[56]: Lukoševičius (2012)

In the original form of ESN, only the readout weights are learned, the rest is chosen randomly. The input $\mathbf{x}(t)$ brings the recurrent units in a initial state. The recurrent connections inside the reservoir create different dynamics in the network. The readout neurons linearly transform the recurrent dynamics into temporal outputs. The readout weights \mathbf{w}_{out} are trained to reproduce a target function $\mathbf{y}(t)$.

Liquid State machines use a spiking neural network instead of a graph of recurrent units as reservoir. The nodes of the spiking neural network are randomly connected together. Thus, every node receives time varying inputs from the inputs as well as from other nodes. The recurrent connections turns the varying input into a spatio-temporal pattern. Similar to ESN, the spatio-temporal patterns of activation are read out by a linear layer.

In general, reservoirs are universal approximators and can approximate any non-linear function given there are enough neurons in the reservoir. They generalize better and faster than equivalent MLP. The main drawback of current systems is that cannot deal well with high-dimensional inputs such as images.

3.1 Natural Intelligence

This thesis is inspired by the work "A Theory of Natural Intelligence" from von der Malsburg et al. [2]. Therefore, we dedicate this section to summarize their work in detail.

According [2], the process of learning is influenced by "nature", "nurture", and "emergence"¹. They point out that human genome (as of nature) only contain 1GB of information [57] and humans only absorb a few GB into permanent memory over a lifetime (as of nurture) but it requires about 1PB to describe the connectivity in human brain. Therefore, it is important to distinguish the amount of information to describe a structure from the amount of information needed to generate it. Similar, nature and nurture only require a few GB to construct, respectively instruct the entire human brain. Therefore, they argue that the human brain must be highly structured (i.e. nature and nurture "generate" the human brain by selecting from a set pre-structured patterns). The authors call the process of generating the highly structured network in the human brain the "Kolmogorov [58] Algorithm of the Brain"². Network self-organization is the only mechanism that has not yet been disproved by experiments as the brains Kolmogorov algorithm [59, 60]. This mechanism loops between activity and connectivity, with activity acting back on connectivity through synaptic plasticity until a steady state, called an attractor network, is reached. The consistency property of an attractor network means that a network has many alternative signal pathways between pairs of neurons [61]. Thus, the brain develops as an overlay of attractor networks called net-fragments [62]. Net-fragments consist of small sets of neurons, whereby each neuron can be part of several net fragments. The network self-organization has to start from an already established coarse global structure which is improved in a coarse-to-fine manner to avoid being caught in a local optima.

Also, von der Malsburg et al. [2] discuss scene representation (i.e. how a scene is represented in the brain) even though they point out that this is a contested concept [63]. Scene representation is a organization framework to put abstract interpretation of scene layouts, elements, potential actions, and emotional responses in relation. The details are not rendered as in photographic images but the framework supports the detailed reconstructions of narrow sectors of the scene. The basic goal if learning is to integrate a behavioral schema into the flow of scene representations. They propose the hypothesis that the network structure resulting from self-organization together with the neural activation in the framework of scene representation are the inductive bias that tunes the brain to the natural environment.

Finally, they discuss how net fragments can be used to implement such structures and processes using vision as an example. They point out that

3.1 Natural Intelligence	17
3.2 Self-Organization	18
Self-Organization in Spiking Neural Networks	20
3.3 Correlation within CNNs	21

1: nature refers to the influence of genes and evolution, nurture to the influence of experience and education

2: as the Kolmogorov (1998) complexity describes the number of bits required by the shortest algorithm that can generate the structure
[58]: Kolmogorov et al. (1976)
[59]: Willschaw et al. (1976)
[60]: Willschaw et al. (1979)

[61]: Malsburg et al. (1987)

[62]: Malsburg (2018)

[63]: Freeman III et al. (1990)

a neuron is grouped in one or multiple net fragments through network self-organization. The net fragments can be considered as filters that detect previously seen patterns in the visual input signal. An object is represented by multiple net fragments, where each fragment responds to the surface of that object and has shared neurons and connections with other net fragments representing that object. Thus, net fragments render the topological structure of the surfaces that dominate the environment. Von der Malsburg et al. [2] propose that net fragments represent shape primitives which can adapt to the shape of actual objects³. Shifter circuits are one possible implementation of networks that enable invariant responses to the position- and shape-variant representations [64, 65]. They are composed of net-fragments that can be formed by network self-organization [66]. Ref. [2] also argue that net fragments are the compositional data structure used by the brain. A hierarchy of features may be represented by nested net fragments of different size. Complex objects, such as mental constructs, can thus be seen as larger net fragments composed as mergers of pre-existing smaller net fragments.

3: adapt in spite of metric deformations, depth rotation, and position

[64]: Arathorn (2002)

[65]: Olshausen et al. (1995)

[66]: Fernandes et al. (2015)

3.2 Self-Organization

[67]: Kelso (1995)

The human brain is self-organizing [67]. Self-organization is the process by which systems consisting of many units spontaneously acquire their structure or function without interference from an external agent or system. The absence of a central control unit allow self-organizing systems to quickly adjust to new environmental conditions. Additionally, such systems have in-built redundancy with a high degree of robustness as they are made of many simpler individual units. These individual units can even fail without the overall system breaking down. Dresch [68] describes seven clearly identified properties of self-organization in the human brain: (i) modular connectivity, (ii) unsupervised learning, (iii) adaptive ability, (iv) functional resiliency, (v) functional plasticity, (vi) from-local-to-global functional organization, and (vii) dynamic system growth.

[68]: Dresch (2020)

Before summarizing the literature specific to self-organization of neural networks, the general literature on self-organization with a focus on deep learning is described in the following. Many of these fundamental algorithms for self-organization serve as inspiration for how ANNs can be designed to be self-organizing.

In nature, groups of millions units that solve complex tasks by using only local interactions can be observed. For example, ants can navigate difficult terrain with a local pheromone-based communication and thus form a collective type of intelligence. Such observations inspired researchers to build algorithms which are based on local communication and self-organization, for example ant colony optimization algorithms [69]. DeepSwarm [70] is a neural architecture search method that uses this algorithm to search for the best neural architecture. This method achieves competitive performance on rather small datasets such as MNIST, Fashion-MNIST, and CIFAR-10.

[69]: Dorigo et al. (1997)

Cellular Automata mimic developmental processes in multi-cell organisms. They contain a grid of similar cells with an internal state which

is updated periodically. The transition from a given state to a subsequent state is defined by some update rules. During an update, cells are only allowed to communicate with the neighbouring cells. Thus, self-organization is enforced by the definition of the update rules. Such automata can be used to study biological pattern formations [71] or physical systems [72]. Neural Cellular Automata [73] use neural networks to learn the update rule. The input in such a neural network is the state of a given cell and its neighbours, the output the subsequent cell state. Usually, the same network is applied to all cells. In this case, a fully connected neural network which is applied to each cell and its local neighbours can be reformulated as a CNN[74]. NCAs can be trained efficiently with gradient descent to grow given 2D patterns such as images[75, 76]. These images are grown through self-organisation (i.e. the pixels pick a color based on the color of neighboring pixels) and are surprisingly resistant to damage. For example, large parts of the images can be removed and the system is able to rebuild these pixels⁴. However, the aforementioned approaches can only grow the pattern they were trained on. A recent method called Variational Neural Cellular Automata [78] use an NCA as decoder of a Variational Autoencoder [79]. This probabilistic generative model can grow a large variety of images from a given input encoded in a vector format. However, there is still a big gap in performance compared to state-of-the-art generative models. Besides growing 2D patterns, NCAs can also create 3D patterns such as buildings in the popular video game Minecraft by utilizing 3D CNNs [80] or generate structures with specific function such as simulated robots able to locomote[81]. Moreover, self-assembling approaches based on NCAs are not restricted to grid-structures. NCAs can be generalized to graph neural networks [82]. Graph cellular automata (GCA) use graph neural networks [83] instead of CNNs to learn the state transition rules and can thus deal with more complex pattern structures than just 2D and 3D grids. NCAs can even be used to control reinforcement learning (RL) agents. Variengien et al. [84] use the observations of the environment as state of the NCA, the subsequent state predicted by the NCA are used as Q-value estimates of a deep Q-learning algorithm [85].

Self-organization can not only be used to generate structures but also to optimize the weights of a neural networks over the agents lifetime. For example, a Hebbian learning rule for meta-learning can be used to self-organize the weights of a RL agent over his lifetime[86]. This means that across multiple episodes the weights of a Hebbian based model are learned. The weights of the agents policy are reset in every episode and the Hebbian based model is used to update them. This allows the agent to adapt better to the changed conditions within the environment.

Besides optimizing the weights, self-organization has also been used to change the learning rule itself. The method “Evolve and Merge” [87] uses the so called “ABCD” Hebbian learning rule which updates the weights as follows:

$$\Delta w_{ij} = \alpha(Ao_i o_j + Bo_i + Co_j + D) \quad (3.1)$$

α is the learning rate, o_i and o_j are the activity levels of connected neurons and A , B , C , and D are learned constants. For each connection in the network is one learning rule initialized and the constants are learned. After a pre-defined number of epochs, the learning rules are

[71]: Wolfram (1984)

[73]: Wulff (1992)

[74]: Gilpin (2019)

[75]: Mordvintsev et al. (2020)

[76]: Mordvintsev et al. (2022)

4: a demo of this regeneration process is available at [77]

[78]: Kähnaman (2022)

[80]: Sudhakaran et al. (2021)

[81]: Horibe et al. (2021)

[82]: Grattarola et al. (2021)

[83]: Zhou et al. (2021)

[84]: Variengien et al. (2021)

[85]: Mnih et al. (2013)

[86]: Najarro et al. (2020)

[87]: Pedersen et al. (2021)

clustered and the ones with similar constants are merged. By repeating this process, the number of parameters can be reduced and robustness increases according to the authors.

Alternatively, it is also possible to initialize the network with shared parameters instead of starting with many rules and merging them over time. Kirsch and Schmidhuber [88] use multiple tiny recurrent neural networks (RNNs) that have the same weight parameters but different internal states⁵. By using self-organization and Hebbian learning, they show that it is possible to learn powerful learning algorithms such as backpropagation while running the network in forward-mode mode only. However, it works only for small-scale problems as it can get stuck in local optima. In general seem self-organizing systems to be hard to optimize and only to work for small datasets or simple problems so far.

Risi [89] describes why self-organizing systems are hard to train; First, the system is hard to control because there is no central entity in charge but the system must still be nudged into the right direction. Second, self-organizing systems are unpredictable (i.e. there exist no mathematical model that tells the outcome of the self-organizing process).

3.2.1 Self-Organization in Spiking Neural Networks

Spiking neural networks (SNNs) (c.f. Section Section 2.4.3) communicate through binary signals known as spikes and are very efficient on special event-based hardware[90]. There exist several methods to self-organize such architectures. For the sake of completeness, two well-known approaches are described in the following. However, since this thesis focuses on self-organization in Deep Learning systems, these approaches are only roughly described and for detailed explanations please refer to the respective literature.

Similar to deep learning, there exists a multitude of different network architectures; Shallow [91, 92] and deep networks [93, 94] structures, fully connected [95] and convolutional layers [96, 97], as well as based on different learning rules such as supervised [98, 99], unsupervised [95, 100] and reinforcement learning based [101].

A representable method for self-organization in SNNs is proposed by Raghavan et al. [102]. They introduce a stackable tool-kit to assemble multi-layer neural networks. This tool-kit is a dynamical system that encapsulates the dynamics of spiking neurons, their interactions as well as the plasticity rules that control the flow of information between layers. Based on the input, spatio-temporal waves are generated that travel across multiple layers. A dynamic learning rule tunes the connectivity between layers based on the properties of the waves tiling the layers⁶.

An alternative method proposed by Raghavan and Thomson [103] grows a neural network. They start with a single computational “cell” and use a wiring algorithm to generate a pooling architecture in a self-organizing fashion. The pooling architecture emerges through two processes; First, a layered neural network is grown. Second, self-organization of its inter-layer connections is used to form defined “pools” or receptive fields. They use the Izhikevich neuron model [46] in the first layer to generate spatio-temporal waves. The units in the second layer learn the “underlying”

[88]: Kirsch et al. (2021)

5: Intuitively, these tiny RNNs can be interpreted as more complex neurons.

[89]: Risi (2021)

[90]: Davies et al. (2018)

[91]: Masquelier et al. (2018)

[92]: Morita et al. (2019)

[93]: Carruthers (2017)

[94]: Tavanaei et al. (2016)

[95]: Diehl et al. (2015)

[96]: Zinke et al. (2018)

[97]: Meunier (2019)

[98]: Ferré et al. (2018)

[99]: Ferré et al. (2018)

[100]: Ferré et al. (2018)

[101]: Ferré et al. (2018)

[102]: Raghavan et al. (2020)

[103]: Raghavan et al. (2019)

[104]: Raghavan et al. (2019)

[105]: Raghavan et al. (2019)

[106]: Raghavan et al. (2019)

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[123]: Raghavan et al. (2019)

[124]: Raghavan et al. (2019)

[125]: Raghavan et al. (2019)

[126]: Raghavan et al. (2019)

[127]: Raghavan et al. (2019)

6: for more information please refer to [102]

[102]: Raghavan et al. (2019)

[46]: Izhikevich (2003)

pattern of activity generated in the first layer. Based on the learned patterns, the inter-layer connections are modified to generate a pooling architecture⁷.

In general, SNNs have to “convert” static input data such as images to a dynamic signal. For example, images are often converted to such signals by using Difference of Gaussian (DoG) convolution filters [50, 104]. Such filters subtract one Gaussian blurred version of an original image from another, less blurred version⁸. This subtraction results in spikes for each pixel. To encode the filter output into a temporal signal, bigger spikes are forwarded earlier in time than smaller spikes. However, such approaches lose a lot of information about the input. For example, in the process described above are all information about color and thin structures lost. To the author of this thesis, this seems to be the reason why these SNNs can’t match the performance of deep learning algorithms so far and often only work well for small gray-scale image-datasets such as MNIST.

7: for more information please refer to [103]

[104]: Vaila et al. (2019)

[50]: Kheradpisheh et al. (2018)

8: DoG filter can thus be used to reduce noise and to detect edges

3.3 Correlation within CNNs

Self-organization in neural networks can be done based on the input data. If Hebbian learning is used⁹, cells are connected based on their correlation (i.e. cells with a high correlation are wired together). One way to capture the correlation within CNNs are Gram matrices. Gram matrices are essentially the dot-product between the channels of a feature map and can capture the style of given image. They are for example used for image style transfer[105] or related fields such as texture synthesis [106]. Appendix Chapter A provides an intuitive explanation what image style transfer is and how this is related to Gram matrices.

9: “Cells that fire together wire together”

[105]: Gatys et al. (2015)

[106]: Gatys et al. (2015)

APPENDIX

Image Style Transfer

A

In the field of image style transfer^[105], such correlations are used to generate images. An image is created based on two images, one image providing the content (i.e. the “content-image”) and the other image providing the style (i.e. the “style-image”). To generate the content of the image, a random image can be fed into the model and a simple distance-based loss function such as the Euclidean norm can be used to calculate the difference between the model output and the content-image. Over time, the model would learn to output the content-image. However, only the content from the content-image and not its style is needed. A solution to this problem is using Convolutional feature maps as they capture spatial information of an image without containing the style information. Therefore, the distance loss is not calculated between the content-image and the model output but between the feature maps of the content-image and the output image. The second task is to transfer the style from the style-image to the model output. The style of an image can be described by the means and correlations across the different feature maps. This information about the image style can be calculated with a Gram matrix. A Gram matrix is the dot product¹ of the flattened feature vectors from a convolutional feature map (i.e. the dot product between all channels within a feature map). For example, a convolutional layer could have multiple channels. The first channel could have a high activation for black and white stripes in horizontal direction while the second channel could have a high activation for black and white stripes in vertical direction. If both channels activate together for the same input and thus have a high correlation, there is a high possibility that the image might contain the style of a chessboard (i.e. black and white checkered). A third channel, for example, could have a high activation for eyes. If this channel has a low correlation with the first channel but a high correlation with the second channel (i.e. black and white stripes in vertical direction), the input image might contain the face of a zebra and thus capture a “zebra-style”. Similar to the content-transfer, a distance loss such as the Euclidean distance can be used to compare the Gram matrices of the style-image and the output-image. Thus, the content is transferred by comparing entire feature maps and the style by comparing the correlations between the channels per feature map (i.e. by comparing gram matrices).

^[105]: Gatys et al. (2015)

1: the dot product between two vectors can be seen as similarity metric; it gets bigger if two vectors are more similar

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