



Deep Learning on SpiNNaker

MASTER THESIS

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Declaration

I declare that this dissertation was composed by myself, that the work contained herein is my own except where explicitly stated otherwise in the text, and that this work has not been submitted for any other degree or professional qualification except as specified.

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Abstract

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1. Introduction

Deep learning is revolutionizing the world. It has become part of our daily lives as consumers, powering major software products—from recommendation systems and translation tools to web search (LeCun et al., 2015). Major breakthroughs in fields like computer vision (Krizhevsky et al., 2012) or natural language processing (Hinton et al., 2012) were achieved through the use of deep learning. It has emerged as a driving force behind discoveries in numerous domains like particle physics (Ciodaro et al., 2012), drug discovery (Ma et al., 2015), genomics (Leung et al., 2014) and gaming (Silver et al., 2016).

Deep learning has become so ubiquitous that we are changing the way we build modern hardware to account for its computational demands. From the way edge devices like mobile phones or embedded systems are built (Deng, 2019) and modern CPUs (Perez, 2017) to specialized hardware designed only for deep learning models, such as Google’s tensor processing unit (TPU) (Jouppi et al., 2017) or NVIDIA’s EGX Edge AI platform (Boitano, 2020). Whole state-of-the-art supercomputers are built solely for deep learning. An example would be a supercomputer built by Microsoft for OpenAI, which is part of the Azure cloud (Langston, 2020).

Hardware manufacturer are faced with a major challenge in meeting the computational demands arising from inference, and more importantly, training deep learning models. OpenAI researchers have estimated that the computational costs of training increases exponentially; approximately every 3.4 months the cost doubles (Amodei et al., 2019). Amodei et al. (2019) claims the deep reinforcement learning agent AlphaGo Zero—the successor of the famous AlphaGo program, which was able to beat Go world champion Lee Sedol (Silver et al., 2017)—to be the system with the highest computational demands of approximately 1850 petaflop/s-days. AlphaGo Zero was trained for 40 days on a machine with 4 TPUs (Silver et al., 2017). With the end of Moore’s Law (Loeffler, 2018), chip makers have to get creative in scaling up computing, the same way machine learning researchers are scaling up their models (Simonite, 2016). Therefore production and research into new hardware designs for deep learning are well on the way.

Another field which has high computational demands for very specific tasks and algorithms is computational neuroscience. Computational neuroscience has long been linked to deep learning, which has its origin in research done by neuroscientists (Goodfellow et al., 2016; McCulloch and Pitts, 1943). While in the recent past deep learning research has been more focused on mathematical topics like statistics and probability theory, optimization or linear algebra, researchers are again looking to neuroscience to further improve the capabilities of deep learning models (Marblestone et al., 2016).

But the algorithms developed by computational neuroscientists are not the only aspect drawing attention from the deep learning community. Computational neuroscience has a long standing history of developing custom hardware for the efficient modeling of the human brain, so called neuromorphic computing. Neuromorphic computing—a computer architecture inspired by the biological nervous system—has been around since the 1980s (Mead, 1989). Today, neuromorphic computers are being developed to meet the demands for efficient computing needed to run large-scale spiking neural networks used for modeling brain functions (Furber, 2016). While being developed mainly for the task of modeling the human brain, deep learning has been linked to neuromorphic computing, especially in the context of commercial usability (Gomes, 2017). Both the low energy demands of neuromorphic computers—such as IBM’s True North (Cassidy et al., 2013) or The University of Manchester’s Spiking Neural Network Architecture (SpiNNaker) (Furber et al., 2006)—and their scalability and massive-parallelism are intriguing for two very important use cases of deep learning:

(i) edge computing, for example robotics and mobile devices, (ii) supercomputers and the cloud-era (Gomes, 2017).

This thesis investigates the performance of SpiNNaker machines for deep learning by training the state-of-the-art computer vision model ResNet-50 (He et al., 2015) under the closed division rules of the MLPerf training benchmark (Mattson et al., 2019). In order to benchmark ResNet-50 on SpiNNaker a prototypical implementation was developed as part of this thesis.

- here a paragraph about the results

Section 2 presents the background of this thesis. An introduction to deep learning is given in Section 2.1, as well as an overview of the benchmark in Section 2.2. Section 2.3 describes the SpiNNaker architecture and compares it to current deep learning hardware. Related work can be found in Section 3. Section 4 presents the architecture of the prototype developed for benchmarking and Section 5 presents the benchmarks and its results. In Section 6 the results of the benchmark are discussed, as well as the development process. Section 7 contains the conclusion, while Section 8 outlines the next steps for further increasing the performance of SpiNNaker by enhancing the prototype.

2. Background

This section summarizes the background knowledge needed for the following sections. First a short introduction to deep learning is given in Section 2.1. The main focus lies on the basic concepts and those concepts important for computer vision. Next, Section 2.2 outlines the context of the conducted benchmark presented in Section 5. Lastly, the SpiNNaker neuromorphic computer architecture is described in Section 2.3. SpiNNaker is also compared against the two state-of-the-art hardware solutions for deep learning that currently produce the best performance in training and inference. Namely general purpose graphical processing units (GPGPUs) and Google’s tensor processing unit (TPU).

2.1 An Introduction to Deep Learning

While it may seem that deep learning is a recent development in the field of artificial intelligence—due to all the recently announced breakthroughs (Senior et al., 2020; Vinyals et al., 2019; OpenAI, 2019; Murphy, 2019)—it has actually existed since the 1940s (Goodfellow et al., 2016). McCulloch and Pitts (1943) first described the McCulloch-Pitts neuron as a simple mathematical model of a biological neuron, which marks the origin of what is known today as deep learning.

Even though deep learning models today are still called *artificial neural networks* (due to their historical context), they are quite different from *spiking neural networks* (which SpiNNaker was designed to run efficiently). While the former has been described as “just nonlinear statistical models” (Hastie et al., 2009), the latter incorporated findings about biological neurons and is therefore more closely related to how the nervous system works (Maass, 1997). Spiking neural networks are mostly used for simulation, unlike deep learning models, which are mostly used for inference.

The history of deep learning can be broken down into three distinct phases. Only during the last of these phases was the methodology called deep learning (Goodfellow et al., 2016). Arguably the reason why deep learning seems to be a new development. The first phase, where deep learning was known as cybernetics, ranged from the 1940s to the 1960s (Goodfellow et al., 2016). As



Figure 1: Schema of a perceptron.

stated above, it was the time when the first biologically inspired representations of neurons were developed. Rosenblatt (1958) presents the first model, a single trainable artificial neuron known as the perceptron (see Figure 1).

Today’s perceptron receives a real-valued n -vector \mathbf{x} of *input* signals and builds the dot product with another real-valued n -vector known as *weights* \mathbf{w} : $\mathbf{x} \cdot \mathbf{w} = \sum_{i=1}^n x_i w_i$. The *bias* b is added to the dot product. $\mathbf{x} \cdot \mathbf{w} + b$ is then passed to the *activation function* g —some fixed transformation function appropriate for the application domain. $y = g(\mathbf{x} \cdot \mathbf{w} + b)$ is the output of the perceptron.

During *supervised learning*, we have a set of *examples*. Each example consists of an *input* vector \mathbf{x} and a associated *label* y generated by an unknown function $f^*(\mathbf{x})$. A perceptron can be trained to approximate $f^*(\mathbf{x})$. We can describe a perceptron as the mathematical function

$$y = f(\mathbf{x}; \mathbf{w}, b) = g(\mathbf{x} \cdot \mathbf{w} + b). \quad (1)$$

$f(\mathbf{x}; \mathbf{w}, b)$ is known as a (*statistical*) *model* with \mathbf{w} and b as its *trainable parameters*, which are trained/learned in order to approximate f^* with f . How a network of perceptrons—a more complex statistical model better suited for real world applications—is trained via backpropagation and gradient descent, will be explained below.

The second historical phase of deep learning is known as connectionism (1980s-1990s) (Goodfellow et al., 2016). Its main contributions to today’s knowledge were the backpropagation algorithm (Rumelhart et al., 1986a) and the approach of parallel distributed processing (Rumelhart et al., 1986b,c), which provided a mathematical framework around the idea that a large number of simple computational units (e.g. the perceptron) could achieve intelligent behavior when con-

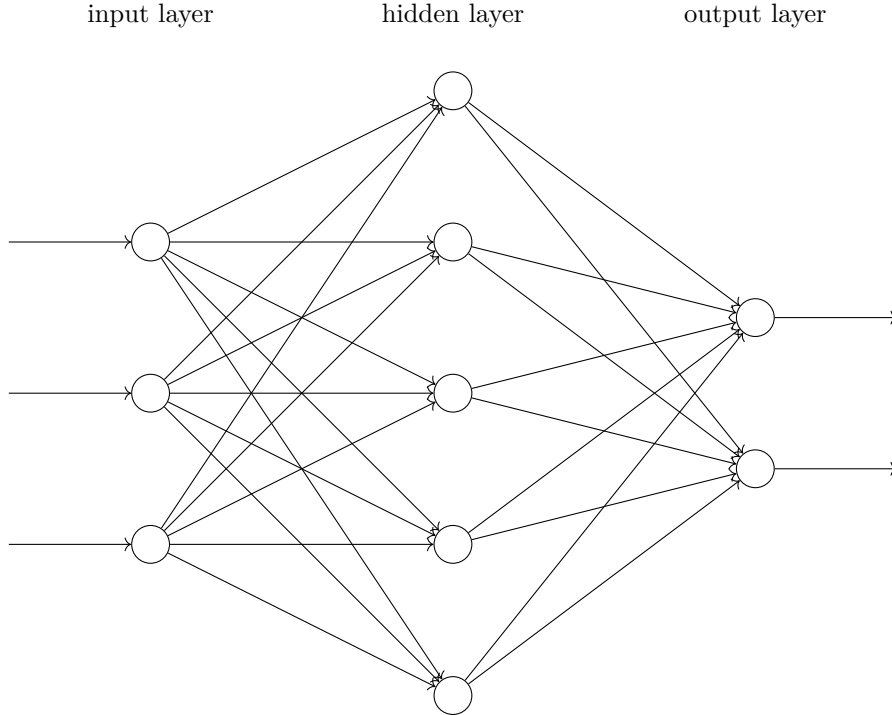


Figure 2: Schema of a MLP or feedforward neural network.

nected together (Goodfellow et al., 2016). Backpropagation enabled the training of networks of perceptrons—artificial neural networks.

The quintessential artificial neural network is the *multilayer perceptron* (MLP), also called a *feedforward neural network* (see Figure 2) (Goodfellow et al., 2016). The MLP consists of multiple perceptrons organized in *layers*. Layers are connected successively such that the output of each of its perceptrons reaches all perceptrons in the next layer. Such a layer is known to be *fully-connected* or *dense*. No cycle exists between perceptrons; the MLP is a directed acyclic graph. Unlike the single layer perceptron, the MLP has at least one *hidden layer*. A hidden layer is a layer between the input and output layers (see Figure 2).

A MLP can also be represented as a statistical model $f(\mathbf{x}; \theta)$. Computing $f(\mathbf{x})$ —also called *inference* or the *forward pass*—can be described as a layer-wise composition of functions $f^{(1)}, f^{(2)}, \dots, f^{(l)}$, each function $f^{(i)}, i < l$ being a hidden layer and $f^{(l)}$ being the output layer. The perceptron has the weight vector \mathbf{w} and the bias b as its parameters (see Equation 1). The parameters of a layer are the combination of \mathbf{w} and b for each of its perceptrons. For example, if the first hidden layer contains m perceptrons and \mathbf{x} is a n -vector, then the parameters of $f^{(1)}$ would be a matrix $\mathbf{W} : n \times m$ and a m -vector \mathbf{b} . The output of layer $f^{(1)}$ would be a m -vector computed as follows:

$$f^{(1)}(\mathbf{x}; \mathbf{W}, \mathbf{b}) = g(\mathbf{W}^\top \mathbf{x} + \mathbf{b}). \quad (2)$$

The second layer takes the output of the first and so forth. The forward pass of the MLP is computed as:

$$y = f^{(l)}(f^{(l-1)}(\dots f^{(1)}(\mathbf{x}))). \quad (3)$$

The backpropagation algorithm is a way to train the parameters of a MLP (or other deep learning models) so that it approximates the unknown function f^* which generates the labels of the examples we have in our data set. The data set used for training a model is called the *training set*. In addition to the training set there normally exists a *test set* with examples the model has not seen before (examples not in the training set). The test set is used to determine the generalization performance of the model. Backpropagation is an algorithm that allows to train a deep learning model with (*stochastic or batch*) *gradient descent*. For example, $\hat{y} = f(\mathbf{x})$ and y is the true label (y and \hat{y} are k -vectors), the error of f is computed using a *loss function* L , for example mean squared error: $1/k \sum_{i=1}^k (y_k - \hat{y}_k)^2$. In order to get the gradients of the weights of the output layer we calculate the derivative of the loss according to each weight w_{ij} in \mathbf{W} with the chain rule:

$$\frac{\delta L}{\delta w_{ij}} = \frac{\delta L}{\delta g} \frac{\delta g}{\delta h} \frac{\delta h}{\delta w_{ij}}, \quad (4)$$

h being $\mathbf{W}^\top f^{(l-1)} + \mathbf{b}$. w_{ij} is updated by performing the stochastic gradient descent¹:

$$w_{ij}^+ = w_{ij} - \mu \frac{\delta L}{\delta w_{ij}}, \quad (5)$$

with μ as the *learning rate*. The same procedure is applied to the following hidden layers. The total loss of the next hidden layer is given as:

$$L^{(l-1)} = \sum_{i=1}^n \frac{\delta L}{\delta f_i^{(l-1)}} = \sum_{i=1}^n \frac{\delta L}{\delta g} \frac{\delta g}{\delta h} \frac{\delta h}{\delta f_i^{(l-1)}}, \quad (6)$$

$f_i^{(l-1)}$ being the i -th perceptron of the hidden layer $l-1$.

Hornik et al. (1989) demonstrated that a non-linear MLP (the activation functions are non-linear transformations of $h(x) = \mathbf{W}^\top \mathbf{x} + \mathbf{b}$) can overcome the famous XOR problem of a single layer perceptron demonstrated in Minsky and Papert (1969). Another major contribution of the phase of connectionism was the neocognitron (Fukushima, 1980), the origin of today's *convolutional neural networks* (CNNs)—which are the state-of-the-art approach for building computer vision models—and the application of the backpropagation algorithm to fully automate the training of CNNs (LeCun et al., 1989).

Goodfellow et al. (2016) claims that the third and current phase of deep learning—where the name deep learning was established—starts with Hinton et al. (2006) describing a new learning algorithm called greedy layer-wise pretraining, which they applied to deep belief networks. Greedy layer-wise pretraining was soon generalized to work with other deep artificial neural network architectures (Ranzato et al., 2006; Bengio et al., 2007). While these papers may have resulted in the

1. Or (batch) gradient descent. With gradient descent the whole training set is passed through the MLP before the weights are updated with the sum over the loss of each example in the training set. Batch or Mini-batch gradient descent takes a subset of the whole training set and updates the weights after each mini-batch. Deep learning models are normally trained by passing the training set multiple times through the model. Each pass over the whole training set is called an *epoch*.

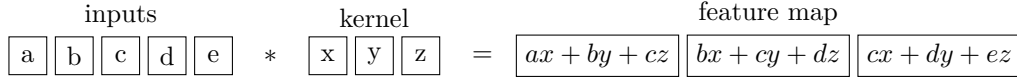


Figure 3: Example of a 1D cross-correlation operation with a kernel size of three, a single channel, a single filter, a stride of one and valid padding.

term deep learning, they were not the reason for the resurrected interest in this methodology. The two most important factors are the increase of available data and computation. The former enables better generalization (Goodfellow et al., 2016), while the latter allows training bigger models (more hidden layers—the *depth* of the neural networks increased) which can solve more complex problems (Bengio and LeCun, 2007; Goodfellow et al., 2016).

Like the perceptron, “neurons” in a *convolutional layer* are inspired by findings of neuroscientists. In this case by research done by Hubel and Wiesel about the mammalian visual cortex (Hubel and Wiesel, 1959, 1962, 1968). CNNs are just deep learning models which have at least one convolutional layer. They are applied to problems which have a grid-like topology, like time-series (1D), images (2D) or videos (3D) (Goodfellow et al., 2016).

Unlike dense layers of perceptrons, convolutional layers do not apply a full matrix multiplication $\mathbf{W}^T \mathbf{x}$ but instead a linear operation $*$ called convolution. A one dimensional discrete convolution can be described as:

$$s(i) = (x * w)(i) = \sum_n x(i + n)w(n). \quad (7)$$

Equation 7 is not really a convolution but is referred to as *cross-correlation*. Unlike true convolution, cross-correlation is not commutative (Goodfellow et al., 2016). However, commutativity is not a factor in practice, so many deep learning libraries, like Keras (Chollet et al., 2015) or the prototype developed for this thesis implement cross-correlation rather than true convolution. Convolution will refer to cross-correlation below.

In the case of deep learning, x is a nD array called the *input* and w is another nD array referred to as the *kernel*. The kernel elements are the trainable parameters (Goodfellow et al., 2016). In Equation 7, x and w are one dimensional. If we let x be a m -vector, the function $x(i)$ is defined as:

$$x(i) = \begin{cases} x_i & \text{if } 1 \leq i \leq m \\ 0 & \text{otherwise.} \end{cases} \quad (8)$$

n is the size of the kernel in the first dimension. Figure 3 shows an example of how the output of a 1D convolutional layer is computed. Figure 4 shows the schema of the convolutional layer performing the operation from Figure 3. The result of a convolution can be transformed by an activation function like the perceptron and the concept of the bias applies also.

Normally a convolutional layer does not consist of a single convolution, but applies multiple kernels to the output of the previous layer. A single convolution is called a *filter* and a layer consists of a predefined number of filters, each with its own kernel (and optionally a bias) (Brownlee, 2019). The output of a convolutional layer is often called a *feature map* (Goodfellow et al., 2016). Even though an image may seem to be a two dimensional structure of pixels, in most cases it is actually three dimensional, the third dimension being the RGB color values for each pixel. The third

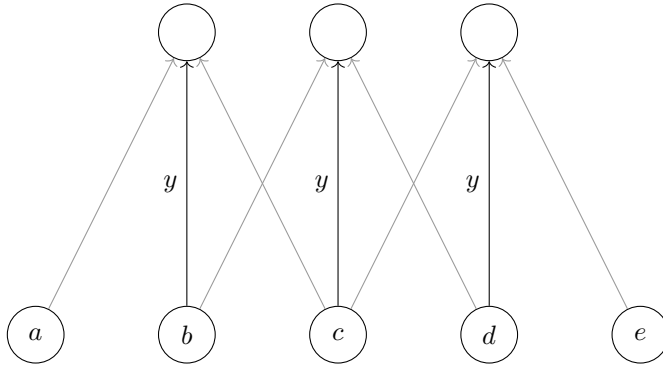


Figure 4: Schema for the convolutional layer performing the convolution shown in Figure 3. Each neuron represents one convolution. The schema shows the property of shared weights and sparse connectivity (Goodfellow et al., 2016). The black edges all have the same associated weight y , while one can see that there are much less edges compared to a dense layer shown in Figure 2.

dimension of the three RGB colors are called the *channels* (Goodfellow et al., 2016). For example, we have a data set of images with 256×256 pixels and three channels (red, green and blue). We pass the image to a convolutional layer with a 3×3 kernel shape and 64 filters. A kernel consists of 18 elements, the kernel size (for the two spatial dimensions) times the three channels of each pixel. The shape of the feature map of that layer—if we assume “same” padding (see below)—would be $256 \times 256 \times 64$, so the next layer would have 64 channels.

There are two more notable concepts of convolutional layers: *stride* and *padding*. The former refers to skipping convolutions in order to reduce the computational cost at the expense of less exact feature extraction (patterns may not be detected by the model due to the increased inaccuracy). The latter is a way of dealing with vanishing spatial dimensions of the feature map if we only perform convolutions on “valid” inputs ($1 \leq i \leq m$ in Equation 8). “Valid” padding refers to the fact that the input has no padding. The feature map of the convolutional layer will have its kernel size minus one less neurons than its input (see Figure 4). “Same” padding would be to add enough zeros evenly above and below the valid input (along each spatial dimension) so that the feature map of the convolutional layer will have the same spatial dimensions as its input (see Figure 5 (Goodfellow et al., 2016)).

Along convolutional layers, CNNs often have *pooling layers*. A pooling layer summarizes locally with the goal of making the CNN invariant to small translations of the input (Goodfellow et al., 2016), making the model less prone to *overfitting*—the state a model is in if it performs well on the training set but does not generalize well to unseen examples. *Max pooling*, for example, takes some local neighborhood of the input, exactly like a convolutional layer, and returns the maximum value of that neighborhood.

2.2 Benchmarking Deep Learning Systems for Computer Vision

In 2010 the annual (until 2017) ImageNet Large Scale Visual Recognition Challenge (ILSVRC) was launched and has become the most famous benchmark for computer vision models, producing many well-known deep learning models like AlexNet in 2012 (Krizhevsky et al., 2012), VGG16 in 2014

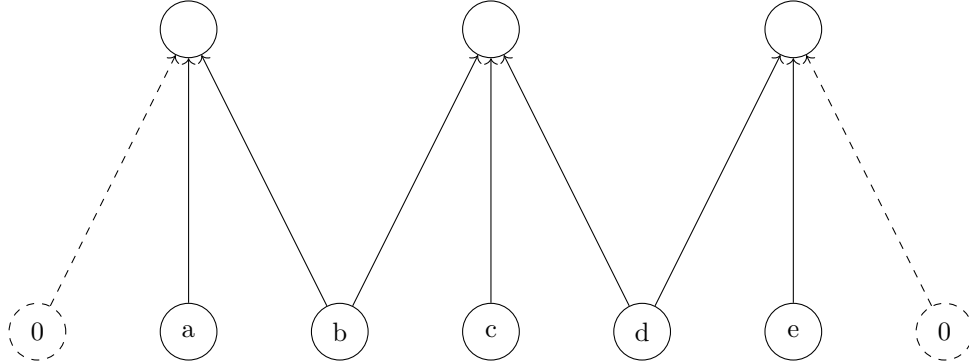


Figure 5: Example showing a layer with same padding and a stride of two.

(Simonyan and Zisserman, 2014) and the ResNet models in 2015 (He et al., 2015). The ILSVRC—like the name suggests—is based on the ImageNet data set consisting of more than 14 million images (Russakovsky et al., 2015). One task of the ILSVRC benchmark is image classification. During image classification the model is trained on 1000 categories (1.2 million images), without overlapping labels (each image has a single label, e.g. “dog”) (Russakovsky et al., 2015). The top-1 ($y = \operatorname{argmax} f(\mathbf{x})$) accuracy is measured on a test set of 150,000 images and winner is the model with the highest top-1 accuracy.

While a benchmark like the ILSVRC produces new insights into computer vision and keeps the community up to date on what is possible, deep learning has another issue on which a benchmark can shed light: training/inference speed of hardware and software systems. The MLPerf benchmark was developed to tackle this problem, so stakeholders can make informed decisions and to provide the industry—like hardware vendors, cloud providers and machine learning engineers—with a fair standard to rely on (Mattson et al., 2019). One task of the MLPerf training benchmark is training the ResNet-50 model (see below) on the image classification task from the ILSVRC 2012, until it reaches a top-1 accuracy of 74.9 percent. The wallclock time is measured and serves as the result for the benchmarked system (Mattson et al., 2019). Currently the fastest solution, from the latest MLPerf training benchmark v0.6, is Google’s cloud system based on Tensorflow and one TPUv3 pod (1024 TPUv3s) (MLPerf, 2019; Stone, 2019). Our benchmark, presented in Section 5, is based on the image classification task of the MLPerf training benchmark, making it easy to compare SpiNNaker and our prototype to other state-of-the-art deep learning systems.

The winner of the image classification task of the ILSVRC 2015 was an ensemble of residual nets (ResNets) introduced in He et al. (2015). The ensemble generated a top-5 accuracy (true label in the five highest outputs of the ensemble) of 96.4 percent. ResNets are a revolution in the sense that they are not only better classifiers than previous models, they also can be significantly deeper (He et al., 2015). He et al. (2015) presents a 152-layer deep network, eight times deeper than a “very deep convolutional network” (VGG11–VGG19) (Simonyan and Zisserman, 2014; He et al., 2015). ResNets can be so deep, without losing their ability of convergence and without degradation (saturated accuracy and higher training error with increased depth) (He et al., 2015), by introducing residual blocks with shortcut connections (see Figure 6). He et al. (2015) hypothesizes that residual blocks ease the learning of the model. These shortcut connections do not increase the complexity

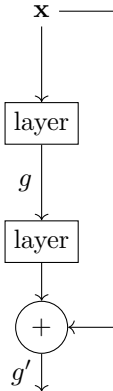


Figure 6: Schema of a residual block with two layers. \mathbf{x} is added to the output of the last layer of the residual block, before the result is passed through its activation function g' .

of the model. No additional parameters are added to the model and nothing changes during backpropagation. Only the negligible operation where \mathbf{x} is added to the output of the residual block must be performed during the forward pass. He et al. (2015) shows comprehensive tests on how residual blocks decrease degradation by comparing ResNets against their counterparts with the same architecture, but without shortcut connections. The models without shortcut connections show a higher training error than their ResNet counterpart.

As stated above, the image classification task of the MLPerf training benchmark is to train ResNet-50 (50, because it has 50 layers) until it reaches a top-1 accuracy of 74.9 percent on the test set and to measure the wallclock time it took to reach that goal. Figure 7 shows an example block from the ResNet-50 model, while Figure 8 shows its architecture. The model takes a $224 \times 224 \times 3$ image as its input and first passes it through a convolutional layer with a relatively big kernel of 7×7 and a max pooling layer. Both times the spatial dimensions are halved by applying a stride of two, so the first residual block receives a $56 \times 56 \times 64$ feature map as its input. The model consists of multiple residual blocks. Some of them have a stride of two. Each time the input is halved that way, channels are doubled. This should keep computational cost the same for each block (He et al., 2015).

2.3 SpiNNaker as a Neuromorphic Computer Architecture

Spiking Neural Network Architecture (SpiNNaker, for short) is a massively parallel neuromorphic computer system designed to run spiking neural networks with up to one billion neurons (and a trillion synapses) in real-time (Painkras et al., 2013). As stated in Section 1, neuromorphic computing is the approach of developing hardware inspired by the biological nervous system (Mead, 1989). Today, neuromorphic computer architectures range from very fast and energy efficient but inflexible direct electronic models (neurons in hardware) (Indiveri et al., 2011) to very flexible but energy demanding systems based on common consumer hardware and software (neurons in software) (Plesser et al., 2007). SpiNNaker sits somewhere in between. On the one hand, flexibility is achieved by implementing neurons in software. On the other hand, speed is achieved by massive-

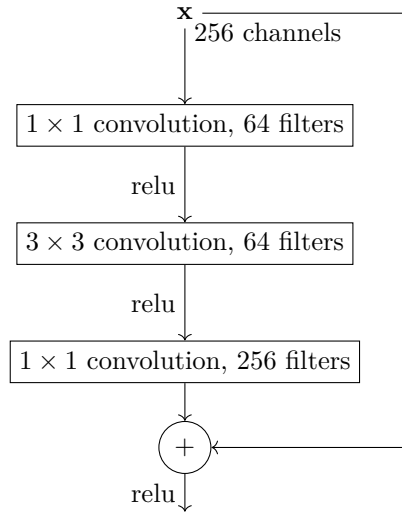


Figure 7: Example of a residual block in ResNet-50.

parallelism and energy efficiency by using energy efficient processors, rather than fast ones (Furber and Bogdan, 2020).

The SpiNNaker system’s basic building block is the SpiNNaker chip, a multiprocessor chip consisting of 18 ARM968 cores and a Network-on-Chip (NoC) system for communication between the cores (Furber and Temple, 2007; Furber and Bogdan, 2020). Each core can run up to 1000 spiking neurons, which communicate with each other over spikes—small packages with a maximum size of 72 bits which are sent over the NoC (Furber and Temple, 2007; SpiNNaker, 2020a). Each core has a 64 Kb DTCM—data tightly-coupled memory—for the application data and fast access to it. The 32 Kb ITCM stores the instructions executed by the core (Furber and Bogdan, 2020). All cores on a chip share access to 128 Mb SDRAM—synchronous dynamic random access memory—which has a higher capacity than DTCM but is also a lot slower (Furber and Bogdan, 2020; SpiNNaker, 2020b).

By today’s standards 18 cores do not qualify as a massively-parallel system. Therefore, a SpiNNaker machine consists of multiple chips connected together in a 2D triangular (six edges per router instead of four) torus (Furber and Bogdan, 2020). The biggest SpiNNaker machine is the SpiNNaker1M supercomputer in Manchester with over one million cores. The SpiNNaker1M consists of 10 cabinets, each with five card frames holding 24 SpiNN-5 boards. A SpiNN-5 board has 48 SpiNNaker chips, which means the SpiNNaker1M has 1,036,800 theoretical cores, assuming no faulty cores (Furber and Bogdan, 2020).

SpiNNaker is quite different from common deep learning accelerators, like general purpose graphics processing units (GPGPUs) or Google’s TPU (Jouppi et al., 2017). Common deep learning libraries like Tensorflow (Abadi et al., 2015), Keras (Chollet et al., 2015) or PyTorch (Paszke et al., 2019) implement deep learning on a layer basis, which means by multiplying *tensors* (n D arrays), rather than implementing deep learning on a neuron level (Goodfellow et al., 2016). Therefore, the common industry approach to building hardware accelerators for deep learning is to facilitate

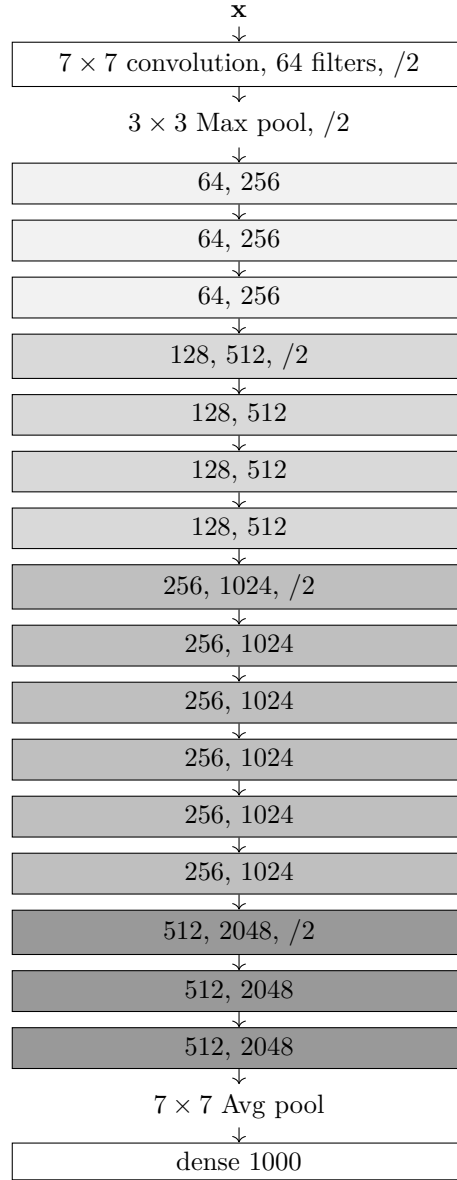


Figure 8: Schema of ResNet-50. Each block in the middle represents one residual block shown in Figure 7. The first number shows the amount of filters the first two layers of a block have, while the second number shows the filters of the last layer of the block. $/2$ indicates that a stride of two is applied (spatial dimensions are halved—each convolutional and pooling layer has “same” padding). Whenever the filters are doubled (indicated by the varying grey scales), the shortcut layer is linearly projected to match the higher channels.

fast matrix multiplication, which represents the majority of computation needed for training and inference. The leading systems, when it comes to throughput and speed according to the MLPerf training benchmark (MLPerf, 2019), are Google’s TPU (Jouppi et al., 2017) and NVIDIA’s GPU architecture Volta (Durant et al., 2017). Volta’s successor Ampere was released in 2020, which, according to NVIDIA, is much more powerful (Krashinsky and Giroux, 2020). Both architectures leverage instruction level parallelism, sacrificing significant chip space for specialized units performing a fused multiply-add-accumulate matrix operation (MAC). NVIDIA calls these units tensor cores. The Tesla V100 has 640 tensor cores, each performing one 4×4 MAC per clock cycle, a theoretical peak performance of 125 terraflop/s (Markidis et al., 2018). The TPUv1 comes with 256×256 MACs performing 8 bit (unsigned) integer operations (Jouppi et al., 2017). The TPUv2 added support for mixed precision floating point operations (16 bit multiply with 32 bit add and accumulate, same as the tensor core of the Volta architecture) (Kennedy, 2017; Markidis et al., 2018). The SpiNNaker cores do not have a MAC unit, being designed to run spiking neurons efficiently, rather than lots of matrix multiplications. That means the prototype presented in Section 4 must put more focus on leveraging SpiNNaker’s massive parallelism, rather than relying on fast instruction level parallelism. For example with optimized domain decomposition and smarter algorithms than matrix multiplication.

3. Related Work

Like Gomes (2017) states, implementing deep learning on neuromorphic chips has been a desire for some time. This section will outline two approaches of implementing deep learning models on neuromorphic hardware. One being the SNN toolbox (Rueckauer et al., 2017), the other being an implementation of CNNs on IBM’s TrueNorth system (Esser et al., 2016).

The SNN toolbox takes pre-trained deep learning models and translates them into spiking neural networks. Its front-end supports a wide range of different input formats from various deep learning libraries, including Keras, Tensorflow, PyTorch or Caffe (Jia et al., 2014), while its back-end supports different spiking neural network simulators like Brian2 (Stimberg et al., 2019), the simulator independent language PyNN (Davison et al., 2009) or direct mappings to neuromorphic computers like SpiNNaker or Intel’s Loihi (Davies et al., 2018; SNN toolbox, 2020). It supports complex CNNs like VGG16 or Inception-v3 (Szegedy et al., 2015; Rueckauer et al., 2017). Rueckauer et al. (2017) shows that using the converted version of LeNet (LeCun et al., 1989) on the MNIST data set (LeCun et al., 2020) and BinaryNet (Courbariaux and Bengio, 2016) on CIFAR-10 (Krizhevsky, 2009) requires two times less operations than the original CNNs without considerable loss in accuracy. Unfortunately for bigger problems, namely VGG16 and Inception-v3 on the ImageNet data set, the converted models have a much lower accuracy than their original counterpart (63.9 percent accuracy for the original VGG16 and only an accuracy of 49.6 for the converted model) (Rueckauer et al., 2017). Another caveat of the SNN toolbox is the fact that it only supports inference and not training, which is the far more complex task computationally.

IBM’s TrueNorth neuromorphic architecture has the goal of achieving energy efficiency and performance through scalability, same as SpiNNaker. Esser et al. (2016) presents Eedn (energy-efficient deep neuromorphic networks). Eedn is an approach of generating CNNs on the TrueNorth system, enabling both inference and training. TrueNorth—unlike SpiNNaker—uses one bit spikes (Esser et al., 2016), which means its substantially different from contemporary consumer hardware and deep learning accelerators and Eedn is needed to translate the CNN in order to make it run on TrueNorth. SpiNNaker on the other hand, like stated above, is just a collection of low power ARM

cores connected over a NoC. Spikes on SpiNNaker are communicated with small multicast packets (up to 72 bits, see above) (Furber and Bogdan, 2020). These packets can be used to transfer any information from one core to another. This makes it much easier to implement deep learning on SpiNNaker, because focus lies more on how to deconstruct the model and map it onto the cores, rather than having to translate the model into a SpiNNaker-specific format. Nonetheless, Eedn models show promising results. Esser et al. (2016) presents tests on 6 well known, industry-strength data sets with the Eedn models having approximately the same accuracy as the original models. The throughput of the TrueNorth system is promising as well. Esser et al. (2016) shows that TrueNorth is able to process between 1,200 and 2,600 $32 \times 32 \times 3$ images per second.

4. Deep Learning on SpiNNaker

This section will describe the developed deep learning prototype in detail. While Section 2.3 gave a short overview over the SpiNNaker hardware, Section 4.1 will present a short introduction to the SpiNNaker software toolchain and programming model. Section 4.2 will present the architecture of the prototype. Lastly, Section 4.3 will shine light onto the hardships and problems encountered and mistakes made during the development process.

4.1 The SpiNNaker Programming Model

Parallel programming is hard (Lee, 2011), especially on novel hardware architectures like SpiNNaker (Brown et al., 2015). SpiNNaker provides layers of software abstractions over the hardware to make programming and exploiting the capabilities of it as easy as possible (Furber and Bogdan, 2020).

The SpiNNaker hardware is designed to tackle problems which can be decomposed into many small, autonomous units without a central computational overseer (Brown et al., 2015). These problems are commonly known as embarrassingly parallel problems (Foster, 1995). The software toolchain lets the user describe their program as a graph. Each vertex of the graph represents one unit of computation (e.g. a bunch of spiking neurons or a perceptron in our case) and directed edges represent the communication between the units (Furber and Bogdan, 2020).

There are two type of graphs, application graphs and machine graphs. A vertex in the machine graph—a machine vertex—is directly mapped to a single SpiNNaker core. An application graph is an abstraction over a machine graph. Application vertices have atoms. Atoms are atomic units of computation. The atoms of an application vertex are distributed onto machine vertices. This makes programming and scaling easier and also facilitates proper resource exploitation, for example by mapping 1,000 small spiking neurons—atoms of a neuron population represented as a application vertex—onto a single core, instead of having them distributed across 1,000 cores, which would be the case if they were to be implemented directly as machine vertices (Furber and Bogdan, 2020). For the prototype we used a machine graph, since it is easier to implement. The development process was guided by the UNIX rule of optimization: prototype before polishing (Raymond, 2003), or in Donald Knuth’s words: “premature optimization is the root of all evil” (Knuth, 1974).

The toolchain is written in the Python programming language. The SpiNNaker machine is connected to a host device via Ethernet (Rowley et al., 2019). The toolchain—running on the host—is mainly responsible for the generation of the graph and its execution on the connected SpiNNaker machine. It goes through a stage of mapping the graph onto the available cores, before data generation (where e.g. parameters of the vertex are loaded into SDRAM) and finally running the application (Furber and Bogdan, 2020).



Figure 9: Example of a machine vertex “source” connected to four other machine vertices over two outgoing edge partitions. If the source vertex sends a MC packet with zero as key, the two upper vertices will receive the packet, whereas with one as key the two lower vertices would receive the packet.

Each vertex is represented as a Python object—instantiated from the appropriate classes—and has an associated binary, the program to be executed by the machine (Furber and Bogdan, 2020). The source code of the binary to be executed on the machine is written in the C programming language and compiled with the gcc compiler from the GNU ARM embedded toolchain (ARM, 2020; Rowley et al., 2019). Machine vertices are not common C programs. They do not own the control flow but instead are event-based, like the ECMAScript programming language (ECMA, 2020). The SpiNNaker1 API provides the operating system executing the vertex and serves the mechanism for registering software callback functions, triggered when a certain event occurs (Furber and Bogdan, 2020). The two events used by the vertices of this prototype were: (i) receiving a packet and (ii) a periodical update event, called every x cycles.

Machine vertices communicate with each other via MC (multicast) packets. A MC packet has two or three segments: (i) one control byte, (ii) 4 byte key and (iii) optionally 4 byte payload. This makes a MC packet either 40 or 72 bits long (Furber and Bogdan, 2020). A MC packet is sent via the directed edges between vertices. Each edge has an associated outgoing edge partition. An outgoing edge partition has one source vertex and n destination vertices and—in the case of the machine graph—one unique routing key (allocated by the toolchain). The routing key—a 32 bit unsigned integer—is unique in the sense that no other outgoing edge partition will have the same key. Otherwise routing would fail in the sense that packets will be sent to the wrong destinations. If the source vertex sends a MC packet it uses the key of the outgoing edge partition. The packet will

reach all destination vertices of the partition. A vertex can have multiple outgoing edge partitions (see Figure 9) (Furber and Bogdan, 2020).

The toolchain offers support for live IO, enabling external devices (like robots, or in our case the host) to interact with the application running on SpiNNaker. Interaction happens, again, via MC packets and by adding extra vertices for input and output to the graph. Live input is enabled by the `ReverseIPTagMulticastSource` (RIPTMCS) machine vertex and live output by the `LivePacketGatherer` (LPG) (Furber and Bogdan, 2020). The toolchain provides a `LiveEventConnection` for the external device, which supplies the appropriate abstractions over the networking. Like the SpiNNaker1 API, the `LiveEventConnection` provides an event-based interface with callbacks.

4.2 The Prototype

The underlying assumption made for developing the prototype was, that because SpiNNaker was designed to run spiking neurons, it would be a natural fit to implement deep learning on a neuron level as well. Besides that, neurons are an easy-to-understand abstraction over the mechanisms of deep learning and rather straight-forward to implement. This design decision was made against the trend of both deep learning research and state-of-the-art deep learning libraries. The former more commonly abstracts over layers while the latter implements deep learning as a computational graph (Goodfellow et al., 2016). Problems with this assumption are discussed in Section 4.3. The API of the prototype was designed to resemble the API of the Keras deep learning library (Chollet et al., 2015). An example comparison can be seen in Listing 1.

The prototype exposes the `Model` class to the user. The model is the main interface to SpiNNaker and—as the name clearly suggests—functions as the representation of a deep learning model. It is designed to be used the same way as Keras’s `Sequential` model (see Listing 1). The second user interface are the layers—the building blocks of a model. Layers are added sequentially to a model instance by calling the model’s `add` method. A layer has at most one preceding and one succeeding layer (see Figure 10). While this suits most common needs, modern deep learning models, like the inception nets (Szegedy et al., 2014), have layers connected to multiple preceding and succeeding layers. Also the shortcut connections of the ResNets (see Section 2.2) are not straight forward to implement with the sequential API. Keras therefore exposes another API, its functional API. The prototype was not developed so far and only supports sequential models.

The model stores the trainable parameters (the weights of the layers), which can be accessed via the `set_weights` and `get_weights` methods of the model class. The parameters are generated during the `add` call by the added layer, since weights are different for different layer types and can depend on the preceding layer. For example, a dense layer generates the weight matrix $\mathbf{W} : n \times m$ and the bias m -vector. n is the amount of neurons in the preceding layer, m the amount of neurons of the dense layer. Convolutional layers do not depend on the previous layer for their weights. For example a 1D convolutional layer generates a 3D array $\mathbf{W} : kernel\ size \times channels \times filters$ and a bias *filters*-vector. \mathbf{W} are the kernels for each filter (see Section 2.1). The kernel of a 2D convolutional layer simply has one more spatial dimension, so the 2D layer would generate a 4D array as weights. The input layer does not generate any weights. The formats of the weights is again inspired by Keras and enables seamless interoperability between a SpiNNaker deep learning model and the subset of the Keras API supported by the prototype (see Listing 1, line 25). Having our prototype so closely resemble Keras had the great advantage of enabling precise integration testing. We could simply build two equal models, one in Keras and one with our prototype, and

```

1 import tensorflow as tf
2 import numpy as np
3
4 # prototype library
5 from spiDNN import Model
6 from spiDNN.layers import Input, Dense
7
8 # random test set
9 X = np.random.rand(500, 64)
10
11 # the keras model
12 keras_model = tf.keras.Sequential()
13 keras_model.add(tf.keras.layers.Dense(128, input_shape=(64,)))
14 keras_model.add(tf.keras.layers.Dense(128, activation="relu"))
15 keras_model.add(tf.keras.layers.Dense(10, activation="softmax"))
16
17 # the equivalent model for SpiNNaker
18 spinn_model = Model()
19 spinn_model.add(Input(64))
20 spinn_model.add(Dense(128))
21 spinn_model.add(Dense(128, activation="relu"))
22 spinn_model.add(Dense(10, activation="softmax"))
23
24 # this call ensures both models have the same parameters
25 model.set_weights(keras_model.get_weights())
26
27 # predict the results for the random test set
28 # (with random weights)
29 p_ = kmodel.predict(X)
30 p = model.predict(X)
31
32 error = np.absolute(p - p_)
33
34 # difference in prediction can happen,
35 # due to floating point errors
36 assert np.amax(error) < 1e-4

```

Listing 1: Example code comparing Keras to the prototype. The code would result in a model akin to the one shown in Figure 10.

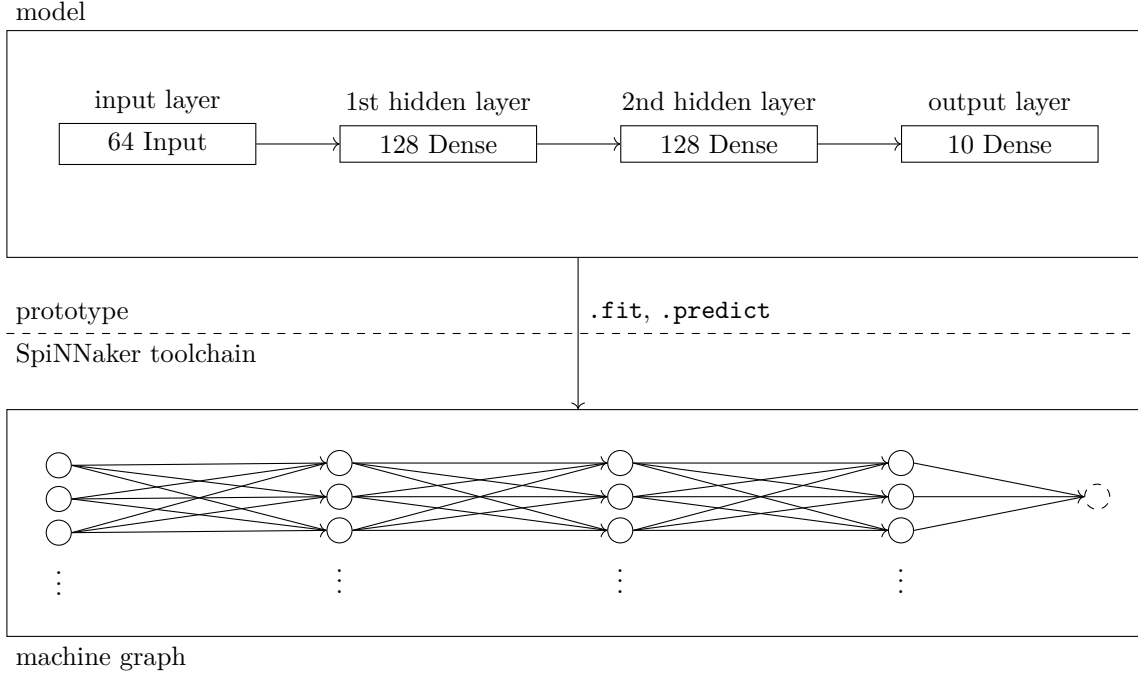


Figure 10: Illustration of how a machine graph is generated by the prototype. The dashed circle represents an auxiliary machine vertex, in this case the LPG. This machine graph would be generated when the `predict` method of the model is called.

compare their outputs against each other (see Listing 1, lines 29ff). The same goes for testing backpropagation, where we could simply train both models and compare the updated weights.

Supported layers are: (i) input, (ii) dense and (iii) 1D convolutional layers. A flatten layer (used for flattening a feature map into a one dimensional vector so it can be processed by a dense layer) is implicitly implemented (see Listing 2). Besides layers, the prototype supports the following activation functions:

- identity (default when activation unspecified): $f(x) = x$.
- tanh: $f(x) = \tanh(x)$
- sigmoid: $f(x) = \frac{1}{1 + \exp(-x)}$
- relu (rectified linear unit): $f(x) = \max(0, x)$
- softmax: $f(x_i) = \frac{\exp(x_i)}{\sum_{j=1}^N \exp(x_j)}$

The only supported optimizer (algorithm for training the weights) is gradient descent with a constant learning rate. Gradient descend is probably the most common optimization algorithm for

```

1 import tensorflow as tf
2 import numpy as np
3
4 # prototype library
5 from spiDNN import Model
6 from spiDNN.layers import Input, Dense, Conv1D
7
8 # 64 features, each with 4 channels
9 input_shape = (64, 4)
10
11 # random test set
12 X = np.random.rand(500, *input_shape)
13
14 # the keras model
15 keras_model = tf.keras.Sequential()
16 # this layer has 20 filters and a kernel size of 3
17 keras_model.add(tf.keras.layers.Conv1D(20, 3, input_shape=input_shape))
18 keras_model.add(tf.keras.layers.Conv1D(5, 3, activation="relu"))
19 keras_model.add(tf.keras.layers.Flatten())
20 keras_model.add(tf.keras.layers.Dense(10, activation="softmax"))
21
22 # the equivalent model for SpiNNaker
23 spinn_model = Model()
24 spinn_model.add(Input(*input_shape))
25 spinn_model.add(Conv1D(20, (3,)))
26 # The feature map of this layer is implicitly flattened,
27 # because the next layer is a dense layer
28 spinn_model.add(Conv1D(5, (3,), activation="relu"))
29 spinn_model.add(Dense(10, activation="softmax"))
30
31 # this call ensures both models have the same parameters
32 model.set_weights(keras_model.get_weights())
33
34 # predict the results for the random test set
35 # (with random weights)
36 p_ = kmodel.predict(X)
37 p = model.predict(X)
38
39 error = np.absolute(p - p_)
40
41 # difference in prediction can happen,
42 # due to floating point errors
43 assert np.amax(error) < 1e-4

```

Listing 2: Example code comparing 1D CNNs in Keras to the prototype.

deep learning (Goodfellow et al., 2016). 1D convolutional layers have support for same and valid (default) padding and they can be strided (see Section 2.1).

Like stated above, a deep learning model is decomposed into neurons in order to execute it on SpiNNaker. Neurons are part of the internal APIs of the prototype and not exposed to the user. The neuron of a dense layer can be seen in Figure 1. Each neuron of a convolutional layer performs one single convolution operation with all filters of the layer. For example, when we look at Figure 3, $ax + by + cz$ would be one convolution with one filter. If the layer has a second filter with a kernel w, v, u , the neuron would produce a second value $aw + bv + cu$. The input a, b, c stays the same. A convolutional layer is schematized in Figure 4.

Figure 10 shows how to operate the prototype. The user defines the sequential deep learning model, like described above. Only when the inference or training operations are called is the model translated into a SpiNNaker machine graph and executed on the connected machine. For inference one has to call the `predict` method of the model and for training the `fit` method.

Algorithm 1 shows what happens during the `predict` method. First, an auxiliary layer for the live IO is created, the extractor layer. The extractor layer has a single machine vertex, a LPG instance (see Section 4.1) for streaming the predictions off of SpiNNaker, back to the host. The first layer—which must always be an input layer—handles streaming the data onto the machine. Neurons of a input layer are simple wrappers around the RIPTMCS machine vertices provided by the SpiNNaker toolchain (see Section 4.1).

The layer interface specifies a `reset` method, which is called for each layer in Algorithm 1, line 2. This method resets the neurons of a layer (simply deletes them). If, for example, the model was trained before the call to the `predict` method—probably the most common workflow in deep learning—the neurons from the training graph would still exist. Beyond the training these neurons are meaningless, because they were part of a different run on SpiNNaker and must be deleted (when the toolchain is resetted it deletes its machine graph instance, but the vertex objects remain). They could be deleted right before the `fit` or `predict` method exits, but for unit testing it is convenient to still have the neurons beyond a run on SpiNNaker.

After the machine and the SpiNNaker toolchain are set up, the machine vertices (neurons) are initialized (see Algorithm 1, line 4). The layer interface always sits between the model and the neurons. The model only ever calls methods of the layers. This is a design pattern in software engineering called the layered pattern (not to be confused with the layers provided by our prototype) (Morlion, 2018). This design pattern is a common way to abstract—e.g. TCP/IP or the OSI model (Tanenbaum and Wetherall, 2013)—and holds code complexity down, makes ownership clear and helps writing well-defined interfaces. For the initialization of the neurons, the layer interface provides a method `init_neurons`. This method generates the neurons, which are stored in the `neurons` property of each layer (a Python list). The `init_neurons` method furthermore adds the neurons to the machine graph of the SpiNNaker toolchain.

After the neurons are created and initialized, the connections for the forward pass are generated (see Algorithm 1, lines 5 and 6). The layer interface offers the `connect_incoming` and `connect_outgoing` functions for this. Incoming and outgoing refer to the called layer, respectively. For example, the forward pass is built with calling `connect_incoming` for each hidden layer, the output layer and the auxiliary layer for extracting the predictions. Each of these layers is connected with the preceding layer. All layers except convolutional layers connect every of their neurons with every neuron of the layer connected with. Connecting convolutional layers is more complicated, because they also need to take care of stride and padding.

Last thing to be set up, before execution can start, is the live IO connection for streaming the observations onto the board and the predictions off it.

Algorithm 1 : predict method

```

1: create extractor layer
2: reset layers
3: setup SpiNNaker
4: initialize machine vertices
5: establish forward pass connections
6: establish connection between output layer and extractor
7: setup live IO
8: execute model on SpiNNaker (run forever) {the live IO connection stops the execution}
9: stop the SpiNNaker machine
10: close live IO
11: return predictions {predictions collected by live IO}

```

- communication structure (partitions and global partition manager)
- ping-pong
- forward pass
- graph structure (especially focused on edge and host-SpiNN communication)
- backward pass
- Conv1D

4.3 Problems

- what is all missing (2D, Pooling and Shortcut connections)
- Too much time spend in receive callback
- interpreting neurons as domain decomposition over linear algebra compute graph
- backward pass: gradients computed two times so comm fabric is not overly used by unique partitions/lots of unused packages
- How I crushed *nd*-kernels into a single blob of weights
- backprop design change (shared weights \rightarrow redundant packets (easier and nicer to implement)). Really stress this point and present all three things tried (e-mail).
- resource invariance (first prototype, deemed too difficult to put resources into it as well (just three months time))
- bottlenecks (one layer much higher computational effort but even less resources (10–1024–20))
- conv neurons of resnet way too big and are horrific scalers (small spatial input but big channels computationally much more effort but again, less resources)

5. Benchmark

6. Discussion

- present possible solutions for problems encountered
- space used inefficiently (cores and memory) → better domain decomposition
- with this streaming model I have no easy way to do batch normalization—implications for vanishing/exploding gradients during training

7. Conclusion

8. Next Steps

- profiling
- cost model
- multiple copies of the same network on the same machine → use all resources available
- better domain decomposition (SpiNNaker application graph or custom solution (application graph not helpful for neurons which become too big))
- smart algorithms vs. integrating with state-of-the-art libraries (investing time in stuff like SLIDE and the one paper by the Austrian guys about sparse connections explicitly mentioning SpiNNaker and neuromorphic chips or rather work on a trans-/compiler that efficiently translates linear algebra operations (like TF, PyTorch,...) onto SpiNNaker)
- integrate into compiler projects like Apache-TVM, XLA, Glow, nGraph, etc.
- implementing ONNX spec to make it easy for developers to use SpiNNaker (develop in PyTorch → run on SpiNNaker)

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