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Chapter 1

Presentation of the field

In this section, we present notions related to our domains of interest. In particular, for tensors we give original definitions that are more appropriate for our study. In the neural network's section, we present the concepts necessary to understand the evolution of the state of the art research in this field. In the last section, we present graphs for their usage in deep learning.

Vector spaces considered in what follows are assumed to be finite-dimensional and over the field of real numbers \mathbb{R} .

1.1	Tensors
1.2	Neural Networks
1.3	Graphs

1.1 Tensors

Intuitively, tensors in the field of deep learning are defined as a generalization of vectors and matrices, as if vectors were tensors of rank 1 and matrices were tensors of rank 2. That is, they are objects in a vector space and their dimensions are indexed using as many indices as their rank, so that they can be represented by multidimensional arrays. In mathematics, a tensor can be defined as a special type of multilinear function (Bass, 1968; Marcus, 1975; Williamson, 2015), which image of a basis can be represented by a multidimensional array. Alternatively, Hackbush propose a mathematical construction of a tensor space as a quotient set of the span of an appropriately defined tensor product (Hackbusch, 2012), which coordinates in a basis can also be represented by a multidimensional array. In particular in the field of mathematics, tensors enjoy an intrinsic definition that neither depend on a representation nor would change the underlying object after a change of basis, whereas in our domain, tensors are confounded with their representation.

1.1.1 Definition

Our definition of tensors is such that they are a bit more than multidimensional arrays but not as much as mathematical tensors, for that they are embedded in a vector space so that deep learning objects can be later defined rigorously.

Given canonical bases, we first define a tensor space, then we relate it to the definition of the tensor product of vector spaces.

Definition 1.1.1. Tensor space

We define a tensor space \mathbb{T} of rank r as a vector space such that its canonical basis is a cartesian product of the canonical bases of r other vector spaces. Its shape is denoted $n_1 \times n_2 \times \cdots \times n_r$, where the $\{n_k\}$ are the dimensions of the vector spaces.

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Definition 1.1.2. Tensor product of vector spaces

Given r vector spaces $\mathbb{V}_1, \mathbb{V}_2, \dots, \mathbb{V}_r$, their tensor product is the tensor space \mathbb{T} spanned by the cartesian product of their canonical bases under coordinatewise sum and outer product.

We use the notation
$$\mathbb{T} = \bigotimes_{k=1}^{n} \mathbb{V}_k$$
.

Remark 1.1.1. This simpler definition is indeed equivalent with the definition of the tensor product given in (Hackbusch, 2012, p. 51). The drawback of our definition is that it depends on the canonical bases, which at first can seem limiting as being canon implies that they are bounded to a certain system of coordinates. However this is not a concern in our domain as we need not distinguish tensors from their representation.

Remark 1.1.2. For naming convenience, from now on, we will distinguish between the terms linear space and vector space i.e. we will abusively use the term vector space only to refer to a linear space that is seen as a tensor space of rank 1. If we don't know its rank, we rather use the term linear space. We also make a clear distinction between the terms dimension (that is, for a tensor space it is equal to $\prod_{k=1}^{r} n_k$) and the term rank (equal to r). Note that some authors use the term order instead of rank (e.g. Hackbusch, 2012) as the latter is affected to another notion.

Definition 1.1.3. Tensor

A tensor t is an object of a tensor space. The shape of t, which is the same as the shape of the tensor space it belongs to, is denoted $n_1^{(t)} \times n_2^{(t)} \times \cdots \times n_r^{(t)}$.

1.1.2 Manipulation

In this subsection, we describe notations and operators used to manipulate data stored in tensors.

Definition 1.1.4. Indexing

An entry of a tensor $t \in \mathbb{T}$ is one of its scalar coordinates in the canonical basis, denoted $t[i_1, i_2, \dots, i_r]$.

More precisely, if $\mathbb{T} = \bigotimes_{k=1} \mathbb{V}_k$, with bases $((e_k^i)_{i=1,\dots,n_k})_{k=1,\dots,r}$, then we have

$$t = \sum_{i_1=1}^{n_1} \cdots \sum_{i_r=1}^{n_r} t[i_1, i_2, \dots, i_r](e_1^{i_1}, \dots, e_r^{i_r})$$

The cartesian product $\mathbb{I} = \prod_{k=1}^{r} \{1, \dots, n_k\}$ is called the *index space* of \mathbb{T}

Remark 1.1.3. When using an index i_k for an entry of a tensor t, we implicitly assume that $i_k \in \{1, 2, \dots, n_k^{(t)}\}$ if nothing is specified.

Definition 1.1.5. Subtensor

A subtensor t' is a tensor of same rank composed of entries of t that are contiguous in the indexing, with at least one entry per rank. We denote $t' = t[l_1:u_1, l_2:u_2, \ldots, l_r:u_r]$, where the $\{l_k\}$ and the $\{u_k\}$ are the lower and upper bounds of the indices used by the entries that compose t'.

Remark 1.1.4. We don't necessarily write the lower bound index if it is equal to 1, neither the upper bound index if it is equal to $n_k^{(t)}$.

Definition 1.1.6. Slicing

A slice operation, along the last ranks $\{r_1, r_2, \dots, r_s\}$, and indexed by $(i_{r_1}, i_{r_2}, \dots, i_{r_s})$, is a morphism $s : \mathbb{T} = \bigotimes_{k=1}^r \mathbb{V}_k \to \bigotimes_{k=1}^{r-s} \mathbb{V}_k$, such that:

$$s(t)[i'_1, i'_2, \dots, i'_{r-s}] = t[i'_1, i'_2, \dots, i'_{r-s}, i_{r_1}, i_{r_2}, \dots, i_{r_s}]$$

$$i.e. \quad s(t) := t[:, :, \dots, :, i_{r_1}, i_{r_2}, \dots, i_{r_s}]$$

where := means that entries of the right operand are assigned to the left operand. We denote $t_{i_{r_1},i_{r_2},...i_{r_s}}$ and call it the *slice* of t. Slicing along a

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subset of ranks that are not the lasts is defined similarly. $s(\mathbb{T})$ is called a slice subspace.

Definition 1.1.7. Flattening

A flatten operation is an isomorphism $f: \mathbb{T} \to \mathbb{V}$, between a tensor space \mathbb{T} of rank r and an n-dimensional vector space \mathbb{V} , where $n = \prod_{k=1}^r n_k$. It is characterized by a bijection in the index spaces $g: \prod_{k=1}^r \{1, \dots, n_k\} \to \{1, \dots, n\}$ such that

$$\forall t \in \mathbb{T}, f(t)[g(i_1, i_2, \dots, i_r)] = f(t[i_1, i_2, \dots, i_r])$$

We call an inverse operation a de-flatten operation.

Remark 1.1.5. Row major ordering

The choice of g determines in which order the indexing is made. g is reminiscent of how data of multidimensional arrays or tensors are stored internally by programming languages. In most tensor manipulation languages, incrementing the memory address (i.e. the output of g) will first increment the last index i_r if $i_r < n_r$ (and if else $i_r = n_r$, then $i_r := 1$ and ranks are ordered in reverse lexicographic order to decide what indices are incremented). This is called row major ordering, as opposed to column major ordering. That is, in row major, g is defined as

$$g(i_1, i_2, \dots, i_r) = \sum_{p=1}^r \left(\prod_{k=p+1}^r n_k\right) i_p$$
 (1.1)

Definition 1.1.8. Reshaping

A reshape operation is an isomorphism defined on a tensor space $\mathbb{T} = \bigotimes_{k=1} \mathbb{V}_k$ such that some of its basis vector spaces $\{\mathbb{V}_k\}$ are de-flattened and some of its slice subspaces are flattened.

1.1.3 Binary operations

We define binary operations on tensors that we'll later have use for. In particular, we define tensor contraction which is sometimes called tensor multiplication, tensor product or tensor dotproduct by other sources. We also define convolution and pooling which serve as the common building blocks of convolution neural network architectures (see Section 1.2.7).

Definition 1.1.9. Contraction

A tensor contraction between two tensors, along ranks of same dimensions, is defined by natural extension of the dot product operation to tensors. More precisely, let \mathbb{T}_1 a tensor space of shape $n_1^{(1)} \times n_2^{(1)} \times \cdots \times n_{r_1}^{(1)}$, and \mathbb{T}_2 a tensor space of shape $n_1^{(2)} \times n_2^{(2)} \times \cdots \times n_{r_2}^{(2)}$, such that $\forall k \in \{1, 2, \ldots, s\}, n_{r_1-(s-k)}^{(1)} = n_k^{(2)}$, then the tensor contraction between $t_1 \in \mathbb{T}_1$ and $t_2 \in \mathbb{T}_2$ is defined as:

$$\begin{cases} t_1 \otimes t_2 = t_3 \in \mathbb{T}_3 \text{ of shape } n_1^{(1)} \times \dots \times n_{r_1-s}^{(1)} \times n_{s+1}^{(2)} \times \dots \times n_{r_2}^{(2)} \text{ where} \\ t_3[i_1^{(1)}, \dots, i_{r_1-s}^{(1)}, i_{s+1}^{(2)}, \dots, i_{r_2}^{(2)}] = \\ \sum_{k_1=1}^{n_1^{(2)}} \dots \sum_{k_s=1}^{n_s^{(2)}} t_1[i_1^{(1)}, \dots, i_{r_1-s}^{(1)}, k_1, \dots, k_s] t_2[k_1, \dots, k_s, i_{s+1}^{(2)}, \dots, i_{r_2}^{(2)}] \end{cases}$$

For the sake of simplicity, we omit the case where the contracted ranks are not the last ones for t_1 and the first ones for t_2 . But this definition still holds in the general case subject to a permutation of the indices.

Definition 1.1.10. Covariant and contravariant indices

Given a tensor contraction $t_1 \otimes t_2$, indices of the left hand operand t_1 that are not contracted are called *covariant* indices. Those that are contracted are called *contravariant* indices. For the right operand t_2 , the naming convention is the opposite. The set of covariant and contravariant indices of both operands are called the *transformation laws* of the tensor contraction.

Remark 1.1.6. Transformation law independency

Contrary to most mathematical definitions, tensors in deep learning are inde-

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pendent of any transformation law, so that they must be specified for tensor contractions.

Remark 1.1.7. Einstein summation convention

Using subscript notation for covariant indices and superscript notation for contravariant indices, the previous tensor contraction can be written using the Einstein summation convention as:

$$t_{1i_{1}^{(1)}\cdots i_{r_{1}-s}^{(1)}}{}^{k_{1}\cdots k_{s}}t_{2k_{1}\cdots k_{s}}{}^{i_{s+1}^{(2)}\cdots i_{r_{2}}^{(2)}}=t_{3i_{1}^{(1)}\cdots i_{r_{1}-s}^{(1)}}{}^{i_{s+1}^{(2)}\cdots i_{r_{2}}^{(2)}} \tag{1.2}$$

Dot product $u_k v^k = \lambda$ and matrix product $A_i^k B_k^j = C_i^j$ are common examples of tensor contractions.

Proposition 1.1.1. A contraction can be rewritten as a matrix product.

Proof. Using notation of (1.2), with the reshapings $t_1 \mapsto T_1$, $t_2 \mapsto T_2$ and $t_3 \mapsto T_3$ defined by grouping all covariant indices into a single index and all contravariant indices into another single index, we can rewrite

$$T_{1g_{i}(i_{1}^{(1)},\ldots,i_{r_{1}-s}^{(1)})}g_{k}(k_{1},\ldots,k_{s})T_{2g_{k}(k_{1},\ldots,k_{s})}g_{j}(i_{s+1}^{(2)},\ldots,i_{r_{2}}^{(2)})=T_{3g_{i}(i_{1}^{(1)},\ldots,i_{r_{1}-s}^{(1)})}g_{j}(i_{s+1}^{(2)},\ldots,i_{r_{2}}^{(2)})$$

where g_i , g_k and g_j are bijections defined similarly as in (1.1).

Definition 1.1.11. Convolution

The *n*-dimensional convolution, denoted $*^n$, between $t_1 \in \mathbb{T}_1$ and $t_2 \in \mathbb{T}_2$, where \mathbb{T}_1 and \mathbb{T}_2 are of the same rank n such that $\forall p \in \{1, 2, ..., n\}, n_p^{(1)} \geq n_p^{(2)}$, is defined as:

$$\begin{cases} t_1 *^n t_2 = t_3 \in \mathbb{T}_3 \text{ of shape } n_1^{(3)} \times \dots \times n_n^{(3)} \text{ where} \\ \forall p \in \{1, 2, \dots, n\}, n_p^{(3)} = n_p^{(1)} - n_p^{(2)} + 1 \\ t_3[i_1, \dots, i_n] = \sum_{k_1 = 1}^{n_1^{(2)}} \dots \sum_{k_n = 1}^{n_n^{(2)}} t_1[i_1 + n_1^{(2)} - k_1, \dots, i_n + n_n^{(2)} - k_n] t_2[k_1, \dots, k_n] \end{cases}$$

Proposition 1.1.2. A convolution can be rewritten as a matrix product.

Proof. Let $t_1 *^n t_2 = t_3$ defined as previously with $\mathbb{T}_1 = \bigotimes_{k=1}^r \mathbb{V}_k^{(1)}$, $\mathbb{T}_2 =$

$$\bigotimes_{k=1}^{r} \mathbb{V}_{k}^{(2)}. \text{ Let } t_{1}' \in \bigotimes_{k=1}^{r} \mathbb{V}_{k}^{(1)} \otimes \bigotimes_{k=1}^{r} \mathbb{V}_{k}^{(2)} \text{ such that } t_{1}'[i_{1}, \dots, i_{n}, k_{1}, \dots, k_{n}] = t_{1}[i_{1} + n_{1}^{(2)} - k_{1}, \dots, i_{n} + n_{n}^{(2)} - k_{n}], \text{ then}$$

$$t_3[i_1,\ldots,i_n] = \sum_{k_1=1}^{n_1^{(2)}} \cdots \sum_{k_n=1}^{n_n^{(2)}} t_1'[i_1,\ldots,i_n,k_1,\ldots,k_n] t_2[k_1,\ldots,k_n]$$

where we recognize a tensor contraction. Proposition 1.1.1 concludes. \Box

The two following operations are meant to further decrease the shape of the resulting output.

Definition 1.1.12. Strided convolution

The *n*-dimensional *strided* convolution, with strides $s = (s_1, s_2, \ldots, s_n)$, denoted $*_s^n$, between $t_1 \in \mathbb{T}_1$ and $t_2 \in \mathbb{T}_2$, where \mathbb{T}_1 and \mathbb{T}_2 are of the same rank *n* such that $\forall p \in \{1, 2, \ldots, n\}, n_p^{(1)} \geq n_p^{(2)}$, is defined as:

$$\begin{cases} t_1 *_s^n t_2 = t_4 \in \mathbb{T}_4 \text{ of shape } n_1^{(4)} \times \dots \times n_n^{(4)} \text{ where} \\ \forall p \in \{1, 2, \dots, n\}, n_p^{(4)} = \lfloor \frac{n_p^{(1)} - n_p^{(2)} + 1}{s_p} \rfloor \\ t_4[i_1, \dots, i_n] = (t_1 *^n t_2)[(i_1 - 1)s_n + 1, \dots, (i_n - 1)s_n + 1] \end{cases}$$

Remark 1.1.8. Informally, a strided convolution is defined as if it were a regular subsampling of a convolution. They match if s = (1, 1, ..., 1).

Definition 1.1.13. Pooling

Let a real-valued function f defined on all tensor spaces of any shape, e.g. the max or average function. An f-pooling operation is a mapping $t \mapsto t'$ such that each entry of t' is an image by f of a subtensor of t.

Remark 1.1.9. Usually, the set of subtensors that are reduced by f into entries of t' are defined by a regular partition of the entries of t.

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1.2 Neural Networks

A feed-forward neural network could originally be formalized as a composite function chaining linear and non-linear functions (Rumelhart et al., 1985; LeCun et al., 1989; LeCun et al., 1995), even up until the important breakthroughs that generated a surge of interest in the field (Hinton et al., 2012; Krizhevsky et al., 2012; Simonyan et al., 2014). However, in more recent advances, more complex architectures have emerged (Szegedy et al., 2015; He et al., 2016; Zoph et al., 2016; Huang et al., 2017), such that the former formalization does not suffice. We provide a definition for the first kind of neural networks (Definition 1.2.1) and use it to present its related concepts. Then we give a more generic definition (Definition 1.2.5).

Note that in this manuscript, we only consider neural networks that are feed-forward (Zell, 1994; Wikipedia, 2018).

1.2.1 Simple formalization

We denote by I_f the domain of definition of a function f ("I" stands for "input") and by $O_f = f(I_f)$ its image ("O" stands for "output"), and we represent it as $I_f \xrightarrow{f} O_f$.

Definition 1.2.1. Neural network (simply connected)

Let f be a function such that I_f and O_f are vector or tensor spaces.

f is a (simply connected) neural network function if there are a series of affine functions $(g_k)_{k=1,2,..,L}$ and a series of non-linear derivable univariate functions $(h_k)_{k=1,2,..,L}$ such that:

$$\begin{cases} \forall k \in \{1, 2, \dots, L\}, f_k = h_k \circ g_k, \\ I_f = I_{f_1} \xrightarrow{f_1} O_{f_1} \cong I_{f_2} \xrightarrow{f_2} \dots \xrightarrow{f_L} O_{f_L} = O_f, \\ f = f_L \circ \dots \circ f_2 \circ f_1 \end{cases}$$

The couple (g_k, h_k) is called the k-th layer of the neural network. L is its

depth. For $x \in I_f$, we denote by $x_k = f_k \circ ... \circ f_2 \circ f_1(x)$ the activations of the k-th layer. We denote by \mathcal{N} the set of neural network functions.

Definition 1.2.2. Activation function

An activation function h is a real-valued univariate function that is non-linear and derivable, that is also defined by extension on any linear space with the functional notation h(v)[i] = h(v[i]).

Definition 1.2.3. Layer

A couple (g, h), where g is an affine or linear function, and h is an activation function is called a *layer*. The set of layers is denoted \mathcal{L} .

Remark 1.2.1. Adoption of ReLU activations

Historically, sigmoidal and tanh activations were mostly used (Cybenko, 1989; LeCun et al., 1989). However in recent practice, the rectified linear unit (ReLU), which implements the rectifier function $h: x \mapsto max(0, x)$ with convention h'(0) = 0 (first introduced as the positive part, Jarrett et al., 2009), is the most used activation, as it was demonstrated to be faster and to obtain better results (Glorot et al., 2011). ReLU originated numerous variants e.g. leaky rectified linear unit (Maas et al., 2013), parametric rectified linear unit (PReLU, He et al., 2015), exponential linear unit (ELU, Clevert et al., 2015), scaled exponential linear unit (SELU, Klambauer et al., 2017).

Remark 1.2.2. Universal approximation

Early researches have shown that neural networks with one level of depth can approximate any real-valued function defined on a compact subset of \mathbb{R}^n . This result was first proved for sigmoidal activations (Cybenko, 1989), and then it was shown it did not depend on the sigmoidal activations (Hornik et al., 1989; Hornik, 1991).

For example, for the application of supervised learning when a neural network is trained from data (see Section 1.2.4), this result is quite important because it brings theoretical justification that the objective exists (even though it doesn't inform whether an algorithm to approach it exists or is efficient).

Remark 1.2.3. Computational difficulty

However, reaching such objective is a computationally difficult problem, which drove back interest from the field. Thanks to better hardware and to using better initialization schemes that speed up learning, researchers started to report more successes with deep neural networks (Hinton et al., 2006; Glorot et al., 2010); see (Bengio, 2009) for a review of this period. It ultimately came to a surge of interest in the field after a significant breakthrough on the imagenet dataset (Deng et al., 2009) with a deep convolutional architecture (Krizhevsky et al., 2012), see Section 1.2.7. The use of the fast ReLU activation function (Glorot et al., 2011) as well as leveraging graphical processing units with CUDA (Nickolls et al., 2008) were also key factors in overcoming this computational difficulty.

Remark 1.2.4. Expressivity and expressive efficiency

The study of the expressivity (also called representational power) of families of neural networks is the field that is interested in the range of functions that can be realized or approximated by this family (Håstad et al., 1991; Pascanu et al., 2013). In general, given a maximal error ϵ and an objective F, the more expressive is a family $N \subset \mathcal{N}$, the more likely it is to contain an approximation $f \in N$ such that $d(f, F) < \epsilon$. However, if we consider the approximation $f_{min} \in N$ that have the lowest number of neurons, it is possible that f_{min} is still too large and may be unpractical. For this reason, expressivity is often studied along the related notion of expressive efficiency (Delalleau et al., 2011; Cohen et al., 2018).

Remark 1.2.5. The class of piecewise linear functions

Of particular interest for the intuition is a result stating that a simply connected neural networks with only ReLU activations (ReNN) is a piecewise linear function (PWL) (Pascanu et al., 2013; Montufar et al., 2014), and that conversely any PWL is also a ReNN such that an upper bound of its depth is logarithmically related to the input dimension (Arora et al., 2018, th. 2.1.).

Remark 1.2.6. Benefits of depth

Expressive efficiency analysis have demonstrated the benefits of depth, *i.e.* a shallow neural network would need an unfeasible large number of neurons to approximate the function of a deep neural network (*e.g.* Delalleau et al., 2011; Bianchini et al., 2014; Poggio et al., 2015; Eldan et al., 2016; Poole et al., 2016; Raghu et al., 2016; Cohen et al., 2016; Mhaskar et al., 2016; Lin et al., 2017; Arora et al., 2018).

Remark 1.2.7. Bias

Affine functions \tilde{g} can be written as a sum between a linear function g and a constant vector b which is called the *bias*. It augments the expressivity of the neural network's family of functions. For notational convenience, we will often omit to write down the biases in the layer's equations.

1.2.2 Generic formalization

The former neural networks are said to be *simply connected* because each layer only takes as input the output of the previous one. We'll give a more general definition after first defining branching operations.

Definition 1.2.4. Branching

A binary branching operation between two tensors, $x_{k_1} \bowtie x_{k_2}$, outputs, subject to shape compatibility, either their addition, either their concatenation along a rank, or their concatenation as a list.

A branching operation between n tensors, $x_{k_1} \bowtie x_{k_2} \bowtie \cdots \bowtie x_{k_n}$, is a composition of binary branching operations, or is the identity function Id if n = 1. Branching operations are also naturally defined on tensor-valued functions through their realizations.

Definition 1.2.5. Neural network (generic definition)

The set of neural network functions \mathcal{N} is defined inductively as follows

- 1. $Id \in \mathcal{N}$
- 2. $f \in \mathcal{N} \land (g,h) \in \mathcal{L} \land O_f \subset I_g \Rightarrow h \circ g \circ f \in \mathcal{N}$

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3. for all shape compatible branching operations:

$$f_1, f_2, \dots, f_n \in \mathcal{N} \Rightarrow f_1 \bowtie f_2 \bowtie \dots \bowtie f_n \in \mathcal{N}$$

Remark 1.2.8. Examples

The neural network proposed in (Szegedy et al., 2015), called *Inception*, use depth-wise concatenation of feature maps. Residual networks (ResNets, He et al., 2016) make use of residual connections, also called skip connections, i.e. an activation that is used as input in a lower level is added to another activation at an upper level. Densely connected networks (DenseNets, Huang et al., 2017) have their activations concatenated with all lower level activations. These neural networks had demonstrated state of the art performances on the imagenet classification challenge (Deng et al., 2009), outperforming simply connected neural networks.

Remark 1.2.9. Benefits of branching operations

Recent works have provided rationales supporting benefits of using branching operations, thus giving justifications for architectures obtained with the generic formalization. In particular, (Cohen et al., 2018) have analyzed the impact of residual connections used in Wavenet-like architectures (Van Den Oord et al., 2016) in terms of expressive efficiency (see Remark 1.2.4) using tools from the field of tensor analysis; (Orhan et al., 2018) have empirically demonstrated that skip connections can resolve some inefficiency problems inherent of fully-connected networks (dead activations, activations that are always equal, linearly dependent sets of activations).

For layer indexing convenience, we still use the simple formalization in the subsequent subsections, even though the presentation would be similar with the generic formalization.

1.2.3 Interpretation

Until now, we have formally introduced a neural network as a mathematical function. As its name suggests, such function can be interpreted from a

connectivity viewpoint (LeCun, 1987).

Definition 1.2.6. Connectivity matrix

Let g a linear function. Without loss of generality subject to a flattening, let's suppose I_g and O_g are vector spaces. Then there exists a *connectivity matrix* W_g , such that:

$$\forall x \in I_g, g(x) = W_g x$$

We denote W_k the connectivity matrix of the k-th layer.

Remark 1.2.10. Biological inspiration

A (computational) neuron is a computational unit that is biologically inspired (McCulloch et al., 1943). Each neuron is capable of:

- 1. receiving modulated signals from other neurons and aggregate them,
- 2. applying to the result a derivable activation,
- 3. passing the signal to other neurons.

That is to say, each domain $\{I_{f_k}\}$ and O_f can be interpreted as a layer of neurons, with one neuron for each dimension. The connectivity matrices $\{W_k\}$ describe the connections between each successive layers. A neuron is illustrated on Figure 1.1.

placeholder

Figure 1.1: A neuron

1.2.4 Parameterization and training

Given an objective function F, training is the process of incrementally modifying a neural network f upon obtaining a better approximation of F. The most used training algorithms are based on gradient descent, as proposed in (Widrow et al., 1960). These algorithms became popular since (Rumelhart et al., 1985). Informally, f is parameterized with initial weights that characterize its linear parts. These weights are modified step by step in the opposite direction of their gradient with respect to a loss. All possible realizations of f through its weights draw a family N which expressivity is crucial for the success of the training. The common points between f and other objects of N define what is called a neural network architecture. That is

We present gradient based learning more formally in what follows.

Definition 1.2.7. Architecture Let $f \in \mathcal{N}$ with weights $(\theta_k)_k \in$.

Remark 1.2.11. Gradient descent

The most used training algorithms are based on gradient descent, as proposed in (Widrow et al., 1960). These algorithms became popular since (Rumelhart et al., 1985). In order to be trained, f is parameterized with initial weights that characterize its linear parts. These weights are modified step by step in the opposite direction of their gradient with respect to a loss.

Remark 1.2.12. Architecture

All possible realizations of f through its weights draw a family N which expressivity is crucial for the success of the training. The common points between f and other objects of N define what is called a neural network architecture.

Definition 1.2.8. Weights

Let consider the k-th layer of a neural network f. We define its weights as

coordinates of a vector θ_k , called the weight kernel, such that:

$$\forall (i, j), \begin{cases} \exists p, W_k[i, j] := \theta_k[p] \\ \text{or } W_k[i, j] = 0 \end{cases}$$

A weight p that appears multiple times in W_k is said to be *shared*. Two parameters of W_k that share a same weight p are said to be *tied*. The number of weights of the k-th layer is $n_1^{(\theta_k)}$.

Remark 1.2.13. Learning

A loss function \mathcal{L} penalizes the output $x_L = f(x)$ relatively to the approximation error |f(x)-F(x)|. Gradient w.r.t. θ_k , denoted ∇_{θ_k} , is used to update the weights via an optimization algorithm based on gradient descent and a learning rate α , that is:

$$\theta_k^{\text{(new)}} = \theta_k^{\text{(old)}} - \alpha \cdot \vec{\nabla}_{\theta_k} \left(\mathcal{L} \left(x_L, \theta_k^{\text{(old)}} \right) + R \left(\theta_k^{\text{(old)}} \right) \right)$$
(1.3)

where α can be a scalar or a vector, \cdot can denote outer or pointwise product, and R is a regularizer. They depend on the optimization algorithm.

TODO: examples of optimization

Remark 1.2.14. Linear complexity

The complexity of computing the gradients is linear with the number of weights.

Proof. Without loss of generality, we assume that the neural network is simply connected. Thanks to the chain rule, ∇_{θ_k} can be computed using gradients that are w.r.t. x_k , denoted ∇_{x_k} , which in turn can be computed using gradients w.r.t. outputs of the next layer k+1, up to the gradients given on the output layer.

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That is:

$$\vec{\nabla}_{\theta_{k}} = J_{\theta_{k}}(x_{k}) \vec{\nabla}_{x_{k}} \tag{1.4}$$

$$\vec{\nabla}_{x_{k}} = J_{x_{k}}(x_{k+1}) \vec{\nabla}_{x_{k+1}}$$

$$\vec{\nabla}_{x_{k+1}} = J_{x_{k+1}}(x_{k+2}) \vec{\nabla}_{x_{k+2}}$$

$$\vdots$$

$$\vec{\nabla}_{x_{L-1}} = J_{x_{L-1}}(x_{L}) \vec{\nabla}_{x_{L}}$$

Obtaining,

$$\vec{\nabla}_{\theta_k} = J_{\theta_k}(x_k) (\prod_{p=k}^{L-1} J_{x_p}(x_{p+1})) \vec{\nabla}_{x_L}$$
(1.6)

where $J_{\text{wrt}}(.)$ are the respective jacobians which can be determined with the layer's expressions and the $\{x_k\}$; and $\vec{\nabla}_{x_L}$ can be determined using \mathcal{L} , R and x_L .

This allows to compute the gradients with a complexity that is linear with the number of weights (only one computation of the activations), instead of being quadratic if it were done with the difference quotient expression of the derivatives (one more computation of the activations for each weight).

Remark 1.2.15. Backpropagation

We can remark that (1.5) rewrites as

$$\vec{\nabla}_{x_k} = J_{x_k}(x_{k+1}) \vec{\nabla}_{x_{k+1}}$$

$$= J_{x_k'}(h(x_k')) J_{x_k}(W_k x_k) \vec{\nabla}_{x_{k+1}}$$

$$(1.7)$$

where $x'_k = W_k x_k$, and these jacobians can be expressed as:

$$J_{x'_{k}}(h(x'_{k}))[i,j] = \delta_{i}^{j}h'(x'_{k}[i])$$

$$J_{x'_{k}}(h(x'_{k})) = I h'(x'_{k})$$
(1.8)

$$J_{x_k}(W_k x_k) = W_k^T (1.9)$$

That means that we can write $\overrightarrow{\nabla}_{x_k} = (\widetilde{h}_k \circ \widetilde{g}_k)(\overrightarrow{\nabla}_{x_{k+1}})$ such that the connectivity matrix \widetilde{W}_k is obtained by transposition. This can be interpreted as gradient calculation being a *back-propagation* on the same neural network, in opposition of the *forward-propagation* done to compute the output.

TODO: Overfitting remark

1.2.5 Examples of layer

Definition 1.2.9. Connections

The set of connections of a layer (g, h), denoted C_q , is defined as:

$$C_g = \{(i, j), \exists p, W_g[i, j] := \theta_g[p]\}$$

We have $0 \le |C_g| \le n_1^{(W_g)} n_2^{(W_g)}$.

Definition 1.2.10. Dense layer

A dense layer (g,h) is a layer such that $|C_g| = n_1^{(W_g)} n_2^{(W_g)}$, i.e. all possible connections exist. The map $(i,j) \mapsto p$ is usually a bijection, meaning that there is no weight sharing.

Definition 1.2.11. Partially connected layer

A partially connected layer (g,h) is a layer such that $|C_g| < n_1^{(W_g)} n_2^{(W_g)}$. A sparsely connected layer (g,h) is a layer such that $|C_g| \ll n_1^{(W_g)} n_2^{(W_g)}$.

Definition 1.2.12. Convolutional layer

A n-dimensional convolutional layer (g,h) is such that the weight kernel θ_g can be reshaped into a tensor w of rank n+2, and such that

$$\begin{cases} I_g \text{ and } O_g \text{ are tensor spaces of rank } n+1 \\ \forall x \in I_g, g(x) = (g(x)_q = \sum_p x_p *^n w_{p,q})_{\forall q} \end{cases}$$

where p and q index slices along the last ranks.

Definition 1.2.13. Feature maps and input channels

The slices $g(x)_q$ are typically called *feature maps*, and the slices x_p are called *input channels*. Let's denote by $n_o = n_{n+1}^{(O_g)}$ and $n_i = n_{n+1}^{(I_g)}$ the number of feature maps and input channels. In other words, Definition 1.2.12 means that for each feature maps, a convolution layer computes n_i convolutions and sums them, computing a total if $n_i \times n_o$ convolutions.

Remark 1.2.16. Note that because they are simply summed, entries of two different input channels that have the same coordinates are assumed to share some sort of relationship. For instance on images, entries of each input channel (typically corresponding to Red, Green and Blue channels) that have the same coordinates share the same pixel location.

Remark 1.2.17. Given a tensor input x, the n-dimensional convolutions between the inputs channels x_p and slices of a weight tensor $w_{p,q}$ would result in outputs y_q of shape $n_1^{(x)} - n_1^{(w)} + 1 \times \ldots \times n_n^{(x)} - n_n^{(w)} + 1$. So, in order to preserve shapes, a padding operation must pad x with $n_1^{(w)} - 1 \times \ldots \times n_n^{(w)} - 1$ zeros beforehand. For example, the padding function of the library tensor-flow (Abadi et al., 2015) pads each rank with a balanced number of zeros on the left and right indices (except if $n_t^{(w)} - 1$ is odd then there is one more zero on the left).

Definition 1.2.14. Padding

A convolutional layer with padding (g, h) is such that g can be decomposed as $g = g_{pad} \circ g'$, where g' is the linear part of a convolution layer as in Definition 1.2.12, and g_{pad} is an operation that pads zeros to its inputs such that g preserves tensor shapes.

Remark 1.2.18. One asset of padding operations is that they limit the possible loss of information on the borders of the subsequent convolutions, as well as preventing a decrease in size. Moreover, preserving shape is needed to

build some neural network architectures, especially for ones with branching operations e.g. Remark 1.2.8. On the other hand, they increase memory and computational footprints.

Proposition 1.2.1. Connectivity matrix of a convolution with padding

A convolutional layer with padding (g, h) is equivalently defined as W_g being a $n_i \times n_o$ block matrix such that its blocks are Toeplitz matrices.

Proof. Let's consider the slices indexed by p and q, and to simplify the notations, let's drop the subscripts p,q. We recall from Definition 1.1.11 that

$$y = (x *^{n} w)[j_{1}, \dots, j_{n}]$$

$$= \sum_{k_{1}=1}^{n_{1}^{(w)}} \dots \sum_{k_{n}=1}^{n_{n}^{(w)}} x[j_{1} + n_{1}^{(w)} - k_{1}, \dots, j_{n} + n_{n}^{(w)} - k_{n}] w[k_{1}, \dots, k_{n}]$$

$$= \sum_{i_{1}=j_{1}}^{j_{1}+n_{1}^{(w)}-1} \dots \sum_{i_{n}=j_{n}}^{j_{n}+n_{n}^{(w)}-1} x[i_{1}, \dots, i_{n}] w[j_{1} + n_{1}^{(w)} - i_{1}, \dots, j_{n} + n_{n}^{(w)} - i_{n}]$$

$$= \sum_{i_{1}=1}^{n_{1}^{(x)}} \dots \sum_{i_{n}=1}^{n_{n}^{(x)}} x[i_{1}, \dots, i_{n}] \widetilde{w}[i_{1}, j_{1}, \dots, i_{n}, j_{n}]$$

$$\text{where } \widetilde{w}[i_{1}, j_{1}, \dots, i_{n}, j_{n}] =$$

$$\begin{cases} w[j_{1} + n_{1}^{(w)} - i_{1}, \dots, j_{n} + n_{n}^{(w)} - i_{n}] & \text{if } \forall t, 0 \leq i_{t} - j_{t} \leq n_{t}^{(w)} - 1 \\ 0 & \text{otherwise} \end{cases}$$

Using Einstein summation convention as in (1.2) and permuting indices, we recognize the following tensor contraction

$$y_{j_1\cdots j_n} = x_{i_1\cdots i_n} \widetilde{w}^{i_1\cdots i_n}_{j_1\cdots j_n} \tag{1.10}$$

Following Proposition 1.1.1, we reshape (1.10) as a matrix product. To reshape $y \mapsto Y$, we use the row major order bijections g_j as in (1.1) defined onto $\{(j_1, \ldots, j_n), \forall t, 1 \leq j_t \leq n_t^{(y)}\}$. To reshape $x \mapsto X$, we use the same row ma-

jor order bijection g_j , however defined on the indices that support non zero-padded values, so that zero-padded values are lost after reshaping. That is, we use a bijection g_i such that $g_i(i_1, i_2, \ldots, i_n) = g_j(i_1 - o_1, i_2 - o_2, \ldots, i_n - o_n)$ defined if and only if $\forall t, 1 + o_t \leq i_t \leq n_t^{(y)}$, where the $\{o_t\}$ are the starting offsets of the non zero-padded values. $\widetilde{w} \mapsto W$ is reshaped by using g_j for its covariant indices, and g_i for its contravariant indices. The entries lost by using g_i do not matter because they would have been nullified by the resulting matrix product. We remark that W is exactly the block (p,q) of W_g (and not of $W_{g'}$). Now let's prove that it is a Toeplitz matrix.

Thanks to the linearity of the expression (1.1) of g_j , by denoting $i'_t = i_t - o_t$, we obtain

$$g_i(i_1, i_2, \dots, i_n) - g_i(j_1, j_2, \dots, j_n) = g_i(i'_1 - j_1, i'_2 - j_2, \dots, i'_n - j_n)$$
 (1.11)

To simplify the notations, let's drop the arguments of g_i and g_j . By bijectivity of g_j , (1.11) tells us that $g_i - g_j$ remains constant if and only if $i'_t - j_t$ remains constant for all t. Recall that

$$W[g_i, g_j] = \begin{cases} w[j_1 + n_1^{(w)} - i_1', \dots, j_n + n_n^{(w)} - i_n'] & \text{if } \forall t, 0 \le i_t' - j_t \le n_t^{(w)} - 1\\ 0 & \text{otherwise} \end{cases}$$

$$(1.12)$$

Hence, on a diagonal of W, $g_i - g_j$ remaining constant means that $W[g_i, g_j]$ also remains constants. So W is a Toeplitz matrix.

The converse is also true as we used invertible functions in the index spaces through the proof. \Box

Remark 1.2.19. The former proof makes clear that the result doesn't hold in case there is no padding. This is due to border effects when the index of the n^{th} rank resets in the definition of the row-major ordering function g_j that would be used. Indeed, under appropriate definitions, the matrices could be seen as almost Toeplitz.

Remark 1.2.20. Comparatively with dense layers, convolution layers enjoy a significant decrease in the number of weights. For example, an input 2 × 2 convolution on images with 3-color input channels, would breed only 12 weights per feature maps, independently of the numbers of input neurons. On image datasets, their usage also breeds a significant boost in performance compared with dense layers (Krizhevsky et al., 2012), for they allow to take advantage of the topology of the inputs while dense layers don't (LeCun et al., 1995). A more thorough comparison and explanation of their assets will be discussed in Section ??.

Definition 1.2.15. Stride

A convolutional layer with stride is a convolutional layer that computes strided convolutions (with stride > 1) instead of convolutions.

Definition 1.2.16. Pooling

A layer with pooling (g, h) is such that g can be decomposed as $g = g' \circ g_{\text{pool}}$, where g_{pool} is a pooling operation.

Remark 1.2.21. Downscaling

Layers with stride or pooling downscale the signals that passes through the layer. These types of layers allows to compute features at a coarser level, giving the intuition that the deeper a layer is in the network, the more abstract is the information captured by the weights of the layer.

TODO: below

1.2.6 Examples of regularization

Remark 1.2.22. Overfitting TODO:

A layer with dropout(g, h) is such that $h = h_1 \circ h_2$, where (g, h_2) is a layer and h_1 is a dropout operation (Srivastava et al., 2014). When dropout is used,

a certain number of neurons are randomly set to zero during the training phase, compensated at test time by scaling down the whole layer. This is done to prevent overfitting.

1.2.7 Examples of architecture

TODO: rephrase

A multilayer perceptron (MLP) (Hornik et al., 1989) is a neural network composed of only dense layers. A convolutional neural network (CNN) (LeCun et al., 1998) is a neural network composed of convolutional layers.

Neural networks are commonly used for machine learning tasks. For example, to perform supervised classification, we usually add a dense output layer $s = (g_{L+1}, h_{L+1})$ with as many neurons as classes. We measure the error between an output and its expected output with a discriminative loss function \mathcal{L} . During the training phase, the weights of the network are adapted for the classification task based on the errors that are back-propagated (Hornik et al., 1989) via the chain rule and according to a chosen optimization algorithm (e.g. Bottou, 2010).

1.3 Graphs

1.3.1 Basic definitions

Definition 1.3.1. Graph

A graph G is defined as a couple of sets $\langle V, E \rangle$ where V is the set of vertices, also called nodes, and $E \subseteq \binom{V}{2}$ is the set of edges. It is associated with a weight mapping $w: E \to \mathbb{R}^*$. The number of vertices |V| is called its order.

Figure 1.2 illustrates an example of a graph. Note that we employ interchangeably the terms *vertex* and *node*.

placeholder

Figure 1.2: Example of a graph

Definition 1.3.2. Adjacency matrix

The adjacency matrix of a graph $G = \langle V, E \rangle$ of order n, is a $n \times n$ real-valued matrix A associated to an indexing of $V = \{v_1, v_2, \dots v_n\}$, such that

$$A[i,j] = \begin{cases} w(\{v_i, v_j\}) & \text{if } \{v_i, v_j\} \in E \\ 0 & \text{otherwise} \end{cases}$$

Definition 1.3.3. Degree and degree matrix

The degree of a vertex $v \in V$ is defined as $\deg(v) = |\{u \in V, \{u, v\} \in E\}|$. The degree matrix of a graph $G = \langle V, E \rangle$ of order n, is the diagonal matrix D, associated to an indexing of $V = \{v_1, v_2, \dots v_n\}$, such that $D = \operatorname{diag}(\deg(v_1), \deg(v_2), \dots, \deg(v_n))$. 1.3. GRAPHS 33

Definition 1.3.4. Laplacian matrix

The laplacian matrix of a graph $G = \langle V, E \rangle$ of order n, associated to an indexing of $V = \{v_1, v_2, \dots v_n\}$, is defined as L = D - A, where D is the degree matrix and A is the adjacency matrix.

Definition 1.3.5. Digraph

A digraph is an oriented graph *i.e.* $E \subseteq V \times V - \{(v, v), v \in V\}$, and its weight mapping w and adjacency matrix A are not symmetric. Notions defined on graphs naturally extends to digraphs where possible.

Definition 1.3.6. Bipartite graph

A bipartite graph is a triplet of sets $\langle V^{(1)}, V^{(2)}, E \rangle$, where $V^{(1)}$ and $V^{(2)}$ are sets of vertices, $V^{(1)} \cap V^{(2)} \neq \emptyset$, and $E \subseteq V^{(1)} \times V^{(2)}$. It is associated with a weight mapping $w: E \to \mathbb{R}^*$. Its adjacency matrix A is associated to indexings of $V^{(1)} = \{v_1^{(1)}, v_2^{(1)}, \dots v_n^{(1)}\}$ and $V^{(2)} = \{v_1^{(2)}, v_2^{(2)}, \dots v_n^{(2)}\}$, such that

$$A[i,j] = \begin{cases} w\left((v_i^{(1)}, v_j^{(2)})\right) & \text{if } (v_i^{(1)}, v_j^{(2)}) \in E\\ 0 & \text{otherwise} \end{cases}$$

We come across the notion of graphs several times in deep learning:

- Connections between two layers of a deep learning model can be represented as a bipartite graph, the *connectivity graph*. It encodes how the information is propagated through a layer to another. See Section 1.3.2.1.
- Neural architectures can be represented by a graph. In particular, a computation graph is used by deep learning programming languages to keep track of the dependencies between layers of a deep learning model, in order to compute forward and back-propagation. See Section 1.3.2.2.
- A graph can represent the underlying structure of an object (often a

vector or a signal). The nodes represent its features, and the edges represent some structural property. See Section 1.3.3.1.

• Datasets can also be graph-structured. The nodes represent the objects of the dataset, and its edge represent some sort of relation between them. See Section 1.3.3.2.

1.3.2 Graphs related to models

1.3.2.1 Connectivity graph

A Connectivity graph is the bipartite graph whose adjacency matrix is the connectivity matrix of a layer of neurons. Formally, given a linear part of a layer, let \mathbf{x} and \mathbf{y} be the input and output signals, n the size of the set of input neurons $N = \{u_1, u_2, \ldots, u_n\}$, and m the size of the set of output neurons $M = \{v_1, v_2, \ldots, v_m\}$. This layer implements the equation $y = \Theta x$ where Θ is a $n \times m$ matrix.

Definition 1.3.7. The connectivity graph G = (V, E) is defined such that $V = N \cup M$ and $E = \{(u_i, v_j) \in N \times M, \Theta_{ij} \neq 0\}.$

I.e. the connectivity graph is obtained by drawing an edge between neurons for which $\Theta_{ij} \neq 0$. For instance, in the special case of a complete bipartite graph, we would obtain a dense layer. Connectivity graphs are especially useful to represent partially connected layers, for which most of the Θ_{ij} are 0. For example, in the case of layers characterized by a small local receptive field, the connectivity graph would be sparse, and output neurons would be connected to a set of input neurons that corresponds to features that are close together in the input space. Figure 1.3 depicts some examples.

TODO: Figure 1.3. It's just a placeholder right now

Connectivity graphs also allow to graphically modelize how weights are tied in a neural layer. Let's suppose the $\Theta_{i,j}$ are taking their values only into the

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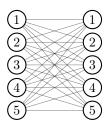


Figure 1.3: Examples

finite set $K = \{w_1, w_2, \dots, w_\kappa\}$ of size κ , which we will refer to as the *kernel* of *weights*. Then we can define a labelling of the edges $s: E \to K$. s is called the *weight sharing scheme* of the layer. This layer can then be formulated as $\forall v \in M, y_v = \sum_{u \in N, (u,v) \in E} w_{s(u,v)} x_u$. Figure 1.4 depicts the connectivity graph of a 1-d convolution layer and its weight sharing scheme.

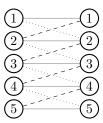


Figure 1.4: Depiction of a 1D-convolutional layer and its weight sharing scheme.

TODO: Add weight sharing scheme in Figure 1.4

- 1.3.2.2 Computation graph
- 1.3.3 Graphs related to data
- 1.3.3.1 Underlying graph structure and signals
- 1.3.3.2 Graph-structured dataset

transductive vs inductive

- 1.3.4 Special classes of graphs
- 1.3.4.1 Grids
- **1.3.4.2** Lattices
- 1.3.4.3 Spatial graphs
- 1.3.4.4 Projections of spatial graphs

Chapter 0

Trash bin

... that we may use in some section.

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TODO: Rework 1.1

0.1 Naming conventions

0.1.1 Basic notions

Let's recall the naming conventions of basic notions.

A function $f: E \to F$ maps objects $x \in E$ to objects $y \in F$, as y = f(x).

Its definition domain $\mathcal{D}_f = E$ is the set of objects onto which it is defined.

We will often just use the term domain.

We also say that f is taking values in its codomain F.

The image per f of the subset $U \subset E$, denoted f(U), is $\{y \in F, \exists x \in E, y = f(x)\}.$

The image of f is the image of its domain. We denote \mathcal{I}_f .

A vector space E, which we will always assume to be finite-dimensional in our context, is defined as \mathbb{R}^n , and is equipped with pointwise addition and scalar multiplication.

A signal s is a function taking values in a vector space. In other words, a signal can also be seen as a vector with an underlying structure, where the vector is composed from its image, and the underlying structure is defined by its domain.

For example, images are signals defined on a set of pixels. Typically, an image s in RGB representation is a mapping from pixels p to a 3d vector space, as $s_p = (r, g, b)$.

TODO?: figure

0.1.2 Graphs and graph signals

TODO: more defs on grid graphs and other graphs

A graph G = (V, E) is defined as a set of nodes V, and a set of edges $E \subseteq \binom{V}{2}$. The words node and vertex will be used equivalently, but we will rather use the first.

A graph signal, or graph-structured signal is a signal defined on the nodes of a graph, for which the underlying structure is the graph itself. A node signal is a signal defined on a node, in which case it is a node embedding in a vector space.

Although this is rarely seen, a signal can also be defined on the edges of a graph, or on an edge. We then coin it respectively dual graph signal, or edge signal / edge embedding.

Graph-structured data can refer to any of these type of signals.

0.1.3 Data and datasets

A dataset of signals is said to be *static* if all its signals share the same underlying structure, it is said to be *non-static* otherwise.

For image datasets, being non-static would mean that the dataset contains images of different sizes or different scales. For graph signal datasets, it would mean that the underlying graph structures of the signals are different.

The point in specifying that objects of a dataset of a machine learning task are signals is that we can hope to leverage their underlying structure.

TODO: figure

0.2 Disambiguation of the subject

This thesis is entitled *Deep learning models for data without a regular structure*. So either the data of interest in this manuscript do not have any structure, or either their structure is not regular.

0.2.1 Irregularly structured data

By structured data, we mean that there exists an underlying structure over which the data is defined. This kind of data are usually modelized as signals defined over a domain. These domains are then composed of objects that are related together by some sort of structural properties. For example, pixels of images can be seen as located on a grid with integer spatial coordinates (a 2d cartesian grid graph).

It then come in handy to define the notions of structure and regularity with the help of graph signals.

Definition 0.2.1. Structure

Let $s: D \to F$ be a signal defined over a finite domain.

An underlying structure of the signal s is a graph G that has the domain of s for nodes.

A dataset is said to be *structured*, if its objects can be modelized as signals with an underlying structure.

It is said to be *static* if all its objects share the same underlying structure, and *non-static* otherwise.

In other words, we chose to define "structured data" as "graph-structured data" by some graph. Hence we need to specify for which graphs this structure would be said to be regular, and for which it would not.

Definition 0.2.2. Regularity

An underlying structure is said to be *regular*, if it is a regular grid graph. It is said to be *irregular* otherwise.

A dataset is said to be *regularly structured*, if the underlying structures of its objects are regular. It is said to be *irregularly structured* otherwise.

TODO: examples

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0.2.2 Unstructured data

Data can also be unstructured. If the data is not yet embedded into a finite dimensional vector space, then we will be interested in embedding techniques used in representation learning. In the other case, it is often possible to fall back to the case of irregularly structured data. For example, vectors can be seen as signals defined over the canonical basis of the vector space, and the vectors of this basis can be related together by their covariances through the dataset. It is typical to use the graph structure that has the canonical basis for nodes, with edges obtained by covariance thresholding.

TODO: examples

What follows is a draft

0.3 Datasets

0.4 Tasks

0.5 Goals

0.6 Invariance

In order to be observed, invariances must be defined relatively to an observation. Let's give a formal definition to support our discussion.

. . .

0.7 Methods

0.8 Expressivity analysis of dense versus sparse connectivity

Let consider a tensor input x of a neural network layer l. Without loss of generality, we consider that x is a matrix of shape $n \times p$. Its rows are supposed structured by a graph $G = \langle V, E \rangle$, with |V| = n, its columns are its feature maps.

In what follows, we discuss the expressivity and efficiency of a dense layer with x as input versus a layer that would leverage G. We start with the regular case and continue onto non-regular structures.

0.8.1 Strong regular case

In the strong regular case, G is a lattice graph such that a convolution is defined naturally on it. For example, this is the case where rows of x defines ticks of a time series, or flattened pixels of an image.

Let consider a convolutional layer $c = (g_c, h_c)$ with padding, defined by q filters of width k. Define y_c its output of shape $n \times q$.

We are interested in knowing if there exists a dense layer that can efficiently replicate c.

Its connectivity matrix W_c is of shape npxnq. Obviously, the function g_c can be replicated by a dummy dense layer $d = (g_c, h_d)$ through W_c . However, whereas c has only kpq weights, d has n^2pq . If we consider the families of neural networks \mathcal{C} , \mathcal{D} spaned by their weights θ_c , θ_d , then we realize the \mathcal{C} is less expressive, but in the same time it is more efficient at representing its functions.

Let's define the notion of partial expressivity with respect to a family of functions.

Let \mathcal{F} a family of functions, \mathcal{L} a family of layer functions, and ϵ the approximation coefficient. For $f \in \mathcal{F}$, define $S_{\epsilon}(\mathcal{L}, f) = \{l \in \mathcal{L}, d(l, f) < \epsilon\}$ and

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$$S_{\epsilon}(\mathcal{L}, \mathcal{F}) = \bigcup_{f \in \mathcal{F}} S_{\epsilon}(\mathcal{L}, f).$$

TODO: reword above

By abusing and anticipating future correction of this manuscript, we consider that \mathcal{C} and \mathcal{D} are vector spaces. We are interesting in 1. proving that $S_{\epsilon}(\mathcal{C}, \mathcal{F})$ and $S_{\epsilon}(\mathcal{D}, \mathcal{F})$ are also vector spaces, and 2. analysing for which \mathcal{F} , $\frac{\dim(S_{\epsilon}(\mathcal{C}, \mathcal{F}))}{\dim(S_{\epsilon}(\mathcal{D}, \mathcal{F}))}$ is maximized.

Obviously 1. is false, so 2. is ill-posed (this draft is to be reworded afterward). Instead of using dim, we should rather use card. However they are potentially infinite families so we should rather use a notion of volume, except if we discretize. So let's discretize.

By the way, "modified" 2. is trivially maximized for $\mathcal{F} = \mathcal{C}$ (and then the ratio equals 1), so let's weaken \mathcal{F} and say it's any family with translation equivariance. We are then interested in proving that if \mathcal{F} is the family on translation equivariant function (on this domains that has to be specified when rewriting this section), then $\frac{card(S_{\epsilon}(\mathcal{C},\mathcal{F}))}{card(S_{\epsilon}(\mathcal{D},\mathcal{F}))}$ is close to 1. Equivariant in our context means commuting with translations (we should rather use the latter expression btw).

The result might be obtained without discretizing as convolutions with padding commutes with translations. Let's guess that they are close to other commuters. In fact that is even it. Proof with Fourier analysis.

0.8.2 Draft

The only dense layer that replicate g_c is obtained through the connectivity matrix W_c . \mathcal{D} is more expressive, however less efficient as we are looking for equivariant functions. It happens that equivariant functions are exactly convolutions with padding.

Index terms— Deep learning, representation learning, propagation learning, visualization, structured, unstructured regular, irregular, covariant, invariant, equivariant, tensor, scheme, weight sharing, graphs, manifold, euclidean, signal processing, graph signal processing, time series, time series database, distributed application, spatial-time series, geo time series, industrial applications, warp 10, warpscript, ...

Temptative titles

- Learning propagational representations of irregular and unstructured data
- Learning representations of unstructured or irregularly structured datasets
- Propagational learning of unstructured or irregularly structured datasets
- Learning tensorial representation of irregular and unstructured data
- Tensorial representation of propagation in deep learning for irregular and unstructured dataset
- Structural representation learning for irregular or unstructured data
- Word for both "irregularly structured" + "unstructured" = ? (maybe "unorthodox" ?)
- Unorthdox deep learning

- ...
- Deep learning of unstructured or irregularly structured datasets
- Deep learning models for data without a regular structure
- On structures in deep learning
- On deep learning for when data is lacking a regular structure
- Deep learning for non regularly structured data

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