Artificial Neural Networks and Deep Learning

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This lecture follows the first parts of these books fairly closely, which treat artificial neural networks.

Textbook, 2nd ed.  
Springer-Verlag  
Heidelberg, DE 2015  
(in German)

Textbook, 2nd ed.  
Springer-Verlag  
Heidelberg, DE 2016  
(in English)
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  Motivation, Biological Background

- **Threshold Logic Units**
  Definition, Geometric Interpretation, Limitations, Networks of TLUs, Training

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- **Deep Learning**
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- **Hopfield Networks and Boltzmann Machines**
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- **Recurrent Neural Networks**
  Differential Equations, Vector Networks, Backpropagation through Time
Motivation: Why (Artificial) Neural Networks?

- **(Neuro-)Biology / (Neuro-)Physiology / Psychology:**
  - Exploit similarity to real (biological) neural networks.
  - Build models to understand nerve and brain operation by simulation.

- **Computer Science / Engineering / Economics**
  - Mimic certain cognitive capabilities of human beings.
  - Solve learning/adaptation, prediction, and optimization problems.

- **Physics / Chemistry**
  - Use neural network models to describe physical phenomena.
  - Special case: spin glasses (alloys of magnetic and non-magnetic metals).
Motivation: Why Neural Networks in AI?

Physical-Symbol System Hypothesis [Newell and Simon 1976]
A physical-symbol system has the necessary and sufficient means for general intelligent action.

Neural networks process simple signals, not symbols.

So why study neural networks in Artificial Intelligence?

- Symbol-based representations work well for inference tasks, but are fairly bad for perception tasks.
- Symbol-based expert systems tend to get slower with growing knowledge, human experts tend to get faster.
- Neural networks allow for highly parallel information processing.
- There are several successful applications in industry and finance.
Diagram of a typical myelinated vertebrate motoneuron (source: Wikipedia, Ruiz-Villarreal 2007), showing the main parts involved in its signaling activity like the *dendrites*, the *axon*, and the *synapses*.
Structure of a prototypical biological neuron (simplified)
(Very) simplified description of neural information processing

- Axon terminal releases chemicals, called neurotransmitters.
- These act on the membrane of the receptor dendrite to change its polarization. (The inside is usually 70mV more negative than the outside.)
- Decrease in potential difference: excitatory synapse
  Increase in potential difference: inhibitory synapse
- If there is enough net excitatory input, the axon is depolarized.
- The resulting action potential travels along the axon. (Speed depends on the degree to which the axon is covered with myelin.)
- When the action potential reaches the terminal buttons, it triggers the release of neurotransmitters.
Recording the Electrical Impulses (Spikes)

pictures not available in online version
An actual recording of the electrical potential also contains the so-called **local field potential (LFP)**, which is dominated by the electrical current flowing from all nearby dendritic synaptic activity within a volume of tissue. The LFP is removed in a preprocessing step (high-pass filtering, $\sim 300\text{Hz}$).

Spikes are detected in the filtered signal with a simple threshold approach. Aligning all detected spikes allows us to distinguishing multiple neurons based on the shape of their spikes. This process is called **spike sorting**.
## (Personal) Computers versus the Human Brain

<table>
<thead>
<tr>
<th></th>
<th><strong>Personal Computer</strong></th>
<th><strong>Human Brain</strong></th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>processing units</strong></td>
<td>1 CPU, 2–10 cores, $10^{10}$ transistors</td>
<td>$10^{11}$ neurons</td>
</tr>
<tr>
<td></td>
<td>1–2 graphics cards/GPUs, $10^3$ cores/shaders</td>
<td></td>
</tr>
<tr>
<td></td>
<td>$10^{10}$ transistors</td>
<td></td>
</tr>
<tr>
<td><strong>storage capacity</strong></td>
<td>$10^{10}$ bytes main memory (RAM)</td>
<td>$10^{11}$ neurons</td>
</tr>
<tr>
<td></td>
<td>$10^{12}$ bytes external memory</td>
<td>$10^{14}$ synapses</td>
</tr>
<tr>
<td><strong>processing speed</strong></td>
<td>$10^{-9}$ seconds, $10^9$ operations per second</td>
<td>&gt; $10^{-3}$ seconds</td>
</tr>
<tr>
<td></td>
<td>&lt; 1000 per second</td>
<td>&lt; 1000 per second</td>
</tr>
<tr>
<td><strong>bandwidth</strong></td>
<td>$10^{12}$ bits/second</td>
<td>$10^{14}$ bits/second</td>
</tr>
<tr>
<td><strong>neural updates</strong></td>
<td>$10^6$ per second</td>
<td>$10^{14}$ per second</td>
</tr>
</tbody>
</table>
• The processing/switching time of a neuron is relatively large (> $10^{-3}$ seconds), but updates are computed in parallel.

• A serial simulation on a computer takes several hundred clock cycles per update.

Advantages of Neural Networks:

• High processing speed due to massive parallelism.

• Fault Tolerance:
  Remain functional even if (larger) parts of a network get damaged.

• “Graceful Degradation”:
  gradual degradation of performance if an increasing number of neurons fail.

• Well suited for inductive learning
  (learning from examples, generalization from instances).

It appears to be reasonable to try to mimic or to recreate these advantages by constructing artificial neural networks.
Threshold Logic Units
A **Threshold Logic Unit (TLU)** is a processing unit for numbers with \( n \) inputs \( x_1, \ldots, x_n \) and one output \( y \). The unit has a **threshold** \( \theta \) and each input \( x_i \) is associated with a **weight** \( w_i \). A threshold logic unit computes the function

\[
y = \begin{cases} 
1, & \text{if } \sum_{i=1}^{n} w_i x_i \geq \theta, \\
0, & \text{otherwise}.
\end{cases}
\]

TLUs mimic the thresholding behavior of biological neurons in a (very) simple fashion.
Threshold Logic Units: Examples

Threshold logic unit for the conjunction $x_1 \land x_2$.

\[
\begin{array}{c}
\text{Threshold logic unit for the conjunction } x_1 \land x_2. \\
\begin{array}{ccc}
\times_1 & \rightarrow & 3 \\
\times_2 & \rightarrow & 2 \\
\end{array}
\end{array}
\]

\[
\begin{array}{|c|c|c|c|}
\hline
x_1 & x_2 & 3x_1 + 2x_2 & y \\
\hline
0 & 0 & 0 & 0 \\
1 & 0 & 3 & 0 \\
0 & 1 & 2 & 0 \\
1 & 1 & 5 & 1 \\
\hline
\end{array}
\]

Threshold logic unit for the implication $x_2 \rightarrow x_1$.

\[
\begin{array}{c}
\text{Threshold logic unit for the implication } x_2 \rightarrow x_1. \\
\begin{array}{ccc}
\times_1 & \rightarrow & 2 \\
\times_2 & \rightarrow & -2 \\
\end{array}
\end{array}
\]

\[
\begin{array}{|c|c|c|c|}
\hline
x_1 & x_2 & 2x_1 - 2x_2 & y \\
\hline
0 & 0 & 0 & 1 \\
1 & 0 & 2 & 1 \\
0 & 1 & -2 & 0 \\
1 & 1 & 0 & 1 \\
\hline
\end{array}
\]
Threshold Logic Units: Examples

Threshold logic unit for \((x_1 \land \overline{x_2}) \lor (x_1 \land x_3) \lor (\overline{x_2} \land x_3)\).

Rough Intuition:

- Positive weights are analogous to excitatory synapses.
- Negative weights are analogous to inhibitory synapses.
Review of line representations

Straight lines are usually represented in one of the following forms:

Explicit Form: \[ g \equiv x_2 = bx_1 + c \]
Implicit Form: \[ g \equiv a_1x_1 + a_2x_2 + d = 0 \]
Point-Direction Form: \[ g \equiv \vec{x} = \vec{p} + k\vec{r} \]
Normal Form: \[ g \equiv (\vec{x} - \vec{p})^\top \vec{n} = 0 \]

with the parameters:

- \( b \): Gradient of the line
- \( c \): Section of the \( x_2 \) axis (intercept)
- \( \vec{p} \): Vector of a point of the line (base vector)
- \( \vec{r} \): Direction vector of the line
- \( \vec{n} \): Normal vector of the line
A straight line and its defining parameters:

\[ b = \frac{r_2}{r_1} \]

\[ \vec{q} = \frac{\vec{p}^\top \vec{n}}{\lVert \vec{n} \rVert} \frac{\vec{n}}{\lVert \vec{n} \rVert} \]

\[ \vec{n} = (a_1, a_2) \]

\[ d = -\vec{p}^\top \vec{n} \]
How to determine the side on which a point $\vec{y}$ lies:

$$\vec{z} = \frac{\vec{y}^\top \vec{n}}{|\vec{n}|} \frac{\vec{n}}{|\vec{n}|}$$

$$\vec{q} = \frac{\vec{p}^\top \vec{n}}{|\vec{n}|} \frac{\vec{n}}{|\vec{n}|}$$
Threshold Logic Units: Geometric Interpretation

Threshold logic unit for $x_1 \land x_2$.

Threshold logic unit for $x_2 \rightarrow x_1$. 

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Artificial Neural Networks and Deep Learning
Threshold Logic Units: Geometric Interpretation

Visualization of 3-dimensional Boolean functions:

Threshold logic unit for \((x_1 \land \overline{x_2}) \lor (x_1 \land x_3) \lor \overline{x_2} \land x_3)\).
The biimplication problem $x_1 \leftrightarrow x_2$: There is no separating line.

<table>
<thead>
<tr>
<th>$x_1$</th>
<th>$x_2$</th>
<th>$y$</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>0</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
</tbody>
</table>

**Formal proof** by *reductio ad absurdum*:

since $(0, 0) \mapsto 1$: $0 \geq \theta$, \hspace{10mm} (1)

since $(1, 0) \mapsto 0$: $w_1 < \theta$, \hspace{10mm} (2)

since $(0, 1) \mapsto 0$: $w_2 < \theta$, \hspace{10mm} (3)

since $(1, 1) \mapsto 1$: $w_1 + w_2 \geq \theta$. \hspace{10mm} (4)

(2) and (3): $w_1 + w_2 < 2\theta$. With (4): $2\theta > \theta$, or $\theta > 0$. Contradiction to (1).
Linear Separability

**Definition:** Two sets of points in a Euclidean space are called **linearly separable**, iff there exists at least one point, line, plane or hyperplane (depending on the dimension of the Euclidean space), such that all points of the one set lie on one side and all points of the other set lie on the other side of this point, line, plane or hyperplane (or on it). That is, the point sets can be separated by a **linear decision function**. Formally: Two sets $X, Y \subset \mathbb{R}^m$ are linearly separable iff $\vec{w} \in \mathbb{R}^m$ and $\theta \in \mathbb{R}$ exist such that

$$\forall \vec{x} \in X : \quad \vec{w}^\top \vec{x} < \theta \quad \text{and} \quad \forall \vec{y} \in Y : \quad \vec{w}^\top \vec{y} \geq \theta.$$ 

- **Boolean functions** define two points sets, namely the set of points that are mapped to the function value 0 and the set of points that are mapped to 1. ⇒ The term “linearly separable” can be transferred to Boolean functions.

- As we have seen, **conjunction** and **implication** are **linearly separable** (as are **disjunction**, NAND, NOR etc.).

- The **biimplication** is not **linearly separable** (and neither is the **exclusive or (XOR)**).
Definition: A set of points in a Euclidean space is called **convex** if it is non-empty and connected (that is, if it is a region) and for every pair of points in it every point on the straight line segment connecting the points of the pair is also in the set.

Definition: The **convex hull** of a set of points $X$ in a Euclidean space is the smallest convex set of points that contains $X$. Alternatively, the convex hull of a set of points $X$ is the intersection of all convex sets that contain $X$.

Theorem: Two sets of points in a Euclidean space are **linearly separable** if and only if their convex hulls are disjoint (that is, have no point in common).

- For the biimplication problem, the convex hulls are the diagonal line segments.
- They share their intersection point and are thus not disjoint.
- Therefore the biimplication is not linearly separable.
Threshold Logic Units: Limitations

Total number and number of linearly separable Boolean functions
(On-Line Encyclopedia of Integer Sequences, oeis.org, A001146 and A000609):

<table>
<thead>
<tr>
<th>inputs</th>
<th>Boolean functions</th>
<th>linearly separable functions</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>4</td>
<td>4</td>
</tr>
<tr>
<td>2</td>
<td>16</td>
<td>14</td>
</tr>
<tr>
<td>3</td>
<td>256</td>
<td>104</td>
</tr>
<tr>
<td>4</td>
<td>65,536</td>
<td>1,882</td>
</tr>
<tr>
<td>5</td>
<td>4,294,967,296</td>
<td>94,572</td>
</tr>
<tr>
<td>6</td>
<td>18,446,744,073,709,551,616</td>
<td>15,028,134</td>
</tr>
<tr>
<td>n</td>
<td>$2^{(2^n)}$</td>
<td>no general formula known</td>
</tr>
</tbody>
</table>

- For many inputs a threshold logic unit can compute almost no functions.
- Networks of threshold logic units are needed to overcome the limitations.
Solving the biimplication problem with a network.

Idea: logical decomposition

\[ x_1 \leftrightarrow x_2 \equiv (x_1 \rightarrow x_2) \land (x_2 \rightarrow x_1) \]

![Diagram](attachment:network_diagram.png)

- Computes \( y_1 = x_1 \rightarrow x_2 \)
- Computes \( y_2 = x_2 \rightarrow x_1 \)
- Computes \( y = y_1 \land y_2 \)
- \( y = x_1 \leftrightarrow x_2 \)
Solving the biimplication problem: Geometric interpretation

- The first layer computes new Boolean coordinates for the points.
- After the coordinate transformation the problem is linearly separable.
Representing Arbitrary Boolean Functions

**Algorithm:** Let \( y = f(x_1, \ldots, x_n) \) be a Boolean function of \( n \) variables.

(i) Represent the given function \( f(x_1, \ldots, x_n) \) in disjunctive normal form. That is, determine \( D_f = C_1 \lor \ldots \lor C_m \), where all \( C_j \) are conjunctions of \( n \) literals, that is, \( C_j = l_{j1} \land \ldots \land l_{jn} \) with \( l_{ji} = x_i \) (positive literal) or \( l_{ji} = \neg x_i \) (negative literal).

(ii) Create a neuron for each conjunction \( C_j \) of the disjunctive normal form (having \( n \) inputs — one input for each variable), where

\[
 w_{ji} = \begin{cases} 
 2, & \text{if } l_{ji} = x_i, \\
 -2, & \text{if } l_{ji} = \neg x_i,
\end{cases} \quad \text{and} \quad \theta_j = n - 1 + \frac{1}{2} \sum_{i=1}^{n} w_{ji}.
\]

(iii) Create an output neuron (having \( m \) inputs — one input for each neuron that was created in step (ii)), where

\[
 w_{(n+1)k} = 2, \quad k = 1, \ldots, m, \quad \text{and} \quad \theta_{n+1} = 1.
\]

Remark: weights are set to \( \pm 2 \) instead of \( \pm 1 \) in order to ensure integer thresholds.
Example: ternary Boolean function:

<table>
<thead>
<tr>
<th>$x_1$</th>
<th>$x_2$</th>
<th>$x_3$</th>
<th>$y$</th>
<th>$C_j$</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>1</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>$x_1 \land \overline{x_2} \land \overline{x_3}$</td>
</tr>
<tr>
<td>0</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td></td>
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<tr>
<td>1</td>
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<td>1</td>
<td>0</td>
<td>1</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>0</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>$\overline{x_1} \land x_2 \land x_3$</td>
</tr>
<tr>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>$x_1 \land x_2 \land x_3$</td>
</tr>
</tbody>
</table>

$D_f = C_1 \lor C_2 \lor C_3$

One conjunction for each row where the output $y$ is 1 with literals according to input values.

First layer (conjunctions):

$C_1 = x_1 \land \overline{x_2} \land \overline{x_3}$

$C_2 = \overline{x_1} \land x_2 \land x_3$

$C_3 = x_1 \land x_2 \land x_3$

Second layer (disjunction):

$D_f = C_1 \lor C_2 \lor C_3$
Example:

Ternary Boolean function:

<table>
<thead>
<tr>
<th>$x_1$</th>
<th>$x_2$</th>
<th>$x_3$</th>
<th>$y$</th>
<th>$C_j$</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>$x_1 \land \overline{x_2} \land \overline{x_3}$</td>
</tr>
<tr>
<td>1</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td></td>
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<tr>
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<td></td>
</tr>
<tr>
<td>0</td>
<td>0</td>
<td>1</td>
<td>1</td>
<td>$\overline{x_1} \land x_2 \land x_3$</td>
</tr>
<tr>
<td>1</td>
<td>0</td>
<td>1</td>
<td>0</td>
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<tr>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>$x_1 \land x_2 \land x_3$</td>
</tr>
</tbody>
</table>

$D_f = C_1 \lor C_2 \lor C_3$

One conjunction for each row where the output $y$ is 1 with literals according to input value.

Resulting network of threshold logic units:

$C_1 = x_1 \land \overline{x_2} \land \overline{x_3}$

$D_f = C_1 \lor C_2 \lor C_3$

$C_2 = \overline{x_1} \land x_2 \land x_3$

$C_3 = x_1 \land x_2 \land x_3$
Theorem: Two sets of points in a Euclidean space are **linearly separable** if and only if their convex hulls are disjoint (that is, have no point in common).

Example function on the preceding slide:

\[
y = f(x_1, x_2, x_3) = (x_1 \land x_2 \land x_3) \lor (\overline{x_1} \land x_2 \land x_3) \lor (x_1 \land x_2 \land x_3)
\]

- The convex hulls of the two point sets are not disjoint (red: intersection).
- Therefore the function \( y = f(x_1, x_2, x_3) \) is not linearly separable.
Training Threshold Logic Units
Training Threshold Logic Units

- Geometric interpretation provides a way to construct threshold logic units with 2 and 3 inputs, but:
  - Not an automatic method (human visualization needed).
  - Not feasible for more than 3 inputs.

- **General idea of automatic training:**
  - Start with random values for weights and threshold.
  - Determine the error of the output for a set of training patterns.
  - Error is a function of the weights and the threshold: \( e = e(w_1, \ldots, w_n, \theta) \).
  - Adapt weights and threshold so that the error becomes smaller.
  - Iterate adaptation until the error vanishes.
Training Threshold Logic Units

Single input threshold logic unit for the negation \( \neg x \).

\[
\begin{array}{ccc}
  x & w & \theta \\
  \downarrow & \downarrow & \downarrow \\
  y & & \\
\end{array}
\]

Output error as a function of weight and threshold.

\[
\begin{array}{c|c|c}
  x & y \\
  0 & 1 \\
  1 & 0 \\
\end{array}
\]

- error for \( x = 0 \)
- error for \( x = 1 \)
- sum of errors
Training Threshold Logic Units

- The error function cannot be used directly, because it consists of plateaus.

- Solution: If the computed output is wrong, take into account how far the weighted sum is from the threshold (that is, consider “how wrong” the relation of weighted sum and threshold is).

Modified output error as a function of weight and threshold.

![Graphs showing modified output error for different values of x and the sum of errors.](image-url)
Training Threshold Logic Units

Schemata of resulting directions of parameter changes.

- Start at a random point.
- Iteratively adapt parameters according to the direction corresponding to the current point.
- Stop if the error vanishes.
Formal Training Rule: Let \( \vec{x} = (x_1, \ldots, x_n)^\top \) be an input vector of a threshold logic unit, \( o \) the desired output for this input vector and \( y \) the actual output of the threshold logic unit. If \( y \neq o \), then the threshold \( \theta \) and the weight vector \( \vec{w} = (w_1, \ldots, w_n)^\top \) are adapted as follows in order to reduce the error:

\[
\theta^{\text{(new)}} = \theta^{\text{(old)}} + \Delta \theta \quad \text{with} \quad \Delta \theta = -\eta (o - y),
\]

\[
\forall i \in \{1, \ldots, n\} : \quad w_i^{\text{(new)}} = w_i^{\text{(old)}} + \Delta w_i \quad \text{with} \quad \Delta w_i = \eta (o - y)x_i,
\]

where \( \eta \) is a parameter that is called learning rate. It determines the severity of the weight changes. This procedure is called Delta Rule or Widrow–Hoff Procedure [Widrow and Hoff 1960].

- **Online Training:** Adapt parameters after each training pattern.
- **Batch Training:** Adapt parameters only at the end of each epoch, that is, after a traversal of all training patterns.
procedure online_training (var $\vec{w}$, var $\theta$, $L$, $\eta$);
var $y$, $e$; (* output, sum of errors *)
begin
    repeat (* training loop *)
        $e := 0$; (* initialize the error sum *)
        for all $(\vec{x}, o) \in L$ do begin (* traverse the patterns *)
            if ($\vec{w}^\top \vec{x} \geq \theta$) then $y := 1$; (* compute the output *)
                else $y := 0$; (* of the threshold logic unit *)
            if ($y \neq o$) then begin (* if the output is wrong *)
                $\theta := \theta - \eta(o - y)$; (* adapt the threshold *)
                $\vec{w} := \vec{w} + \eta(o - y)\vec{x}$; (* and the weights *)
                $e := e + |o - y|$; (* sum the errors *)
            end; (* repeat the computations *)
        end; (* until the error vanishes *)
    until ($e \leq 0$);
end;
Training Threshold Logic Units: Delta Rule

procedure batch_training (var \( \vec{w} \), var \( \theta, L, \eta \));
var \( y, e, \theta_c, \vec{w}_c \); (* output, sum of errors, sums of changes *)
begin
  repeat
    \( e := 0; \theta_c := 0; \vec{w}_c := \vec{0}; \) (* training loop *)
    for all \((\vec{x}, o) \in L\) do begin
      \( \text{if } (\vec{w}^\top \vec{x} \geq \theta) \text{ then } y := 1; \) (* initializations *)
      \( \text{else } y := 0; \) (* traverse the patterns *)
      \( \text{if } (y \neq o) \text{ then begin } \) (* compute the output *)
        \( \theta_c := \theta_c - \eta(o - y); \) (* of the threshold logic unit *)
        \( \vec{w}_c := \vec{w}_c + \eta(o - y)\vec{x}; \) (* if the output is wrong *)
        \( e := e + |o - y|; \) (* sum the changes of the *)
      \( \text{end; } \) (* threshold and the weights *)
    \end;
    \( \theta := \theta + \theta_c; \) (* sum the errors *)
    \( \vec{w} := \vec{w} + \vec{w}_c; \) (* adapt the threshold *)
    \( \text{until } (e \leq 0); \) (* and the weights *)
  end;
(* repeat the computations *)
(* until the error vanishes *)
## Training Threshold Logic Units: Online

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Training Threshold Logic Units

Example training procedure: Online and batch training.

Online Training

Batch Training

Batch Training

\[ x \xrightarrow{-1} \frac{1}{2} \rightarrow y \]
Training Threshold Logic Units: Conjunction

Threshold logic unit with two inputs for the conjunction.

\[ y = \begin{cases} 1 & \text{if } x_1 w_1 + x_2 w_2 \geq \theta \\ 0 & \text{otherwise} \end{cases} \]

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Christian Borgelt
Artificial Neural Networks and Deep Learning
Training Threshold Logic Units: Conjunction

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</table>
Training Threshold Logic Units: Convergence

**Convergence Theorem:** Let \( L = \{(\vec{x}_1, o_1), \ldots, (\vec{x}_m, o_m)\} \) be a set of training patterns, each consisting of an input vector \( \vec{x}_i \in \mathbb{R}^n \) and a desired output \( o_i \in \{0, 1\} \). Furthermore, let \( L_0 = \{(\vec{x}, o) \in L \mid o = 0\} \) and \( L_1 = \{(\vec{x}, o) \in L \mid o = 1\} \). If \( L_0 \) and \( L_1 \) are linearly separable, that is, if \( \vec{w} \in \mathbb{R}^n \) and \( \theta \in \mathbb{R} \) exist such that

\[
\forall (\vec{x}, 0) \in L_0 : \quad \vec{w}^\top \vec{x} < \theta \quad \text{and} \quad
\forall (\vec{x}, 1) \in L_1 : \quad \vec{w}^\top \vec{x} \geq \theta,
\]

then online as well as batch training terminate.

- The algorithms terminate only when the error vanishes.
- Therefore the resulting threshold and weights must solve the problem.
- For not linearly separable problems the algorithms do not terminate (oscillation, repeated computation of same non-solving \( \vec{w} \) and \( \theta \)).
Turning the threshold value into a weight:

\[ \sum_{i=1}^{n} w_i x_i \geq \theta \]

\[ \sum_{i=1}^{n} w_i x_i - \theta \geq 0 \]
**Formal Training Rule** (with threshold turned into a weight):
Let \( \vec{x} = (x_0 = 1, x_1, \ldots, x_n)^\top \) be an (extended) input vector of a threshold logic unit, \( o \) the desired output for this input vector and \( y \) the actual output of the threshold logic unit. If \( y \neq o \), then the (extended) weight vector \( \vec{w} = (w_0 = -\theta, w_1, \ldots, w_n)^\top \) is adapted as follows in order to reduce the error:

\[
\forall i \in \{0, \ldots, n\} : \quad w_i^{\text{(new)}} = w_i^{\text{(old)}} + \Delta w_i \quad \text{with} \quad \Delta w_i = \eta (o - y) x_i,
\]

where \( \eta \) is a parameter that is called **learning rate**. It determines the severity of the weight changes. This procedure is called **Delta Rule** or **Widrow–Hoff Procedure** [Widrow and Hoff 1960].

- Note that with extended input and weight vectors, there is only one update rule (no distinction of threshold and weights).
- Note also that the (extended) input vector may be \( \vec{x} = (x_0 = -1, x_1, \ldots, x_n)^\top \) and the corresponding (extended) weight vector \( \vec{w} = (w_0 = +\theta, w_1, \ldots, w_n)^\top \).
• Single threshold logic units have strong limitations: They can only compute linearly separable functions.

• Networks of threshold logic units can compute arbitrary Boolean functions.

• Training single threshold logic units with the delta rule is easy and fast and guaranteed to find a solution if one exists.

• Networks of threshold logic units cannot be trained, because
  ○ there are no desired values for the neurons of the first layer(s),
  ○ the problem can usually be solved with several different functions computed by the neurons of the first layer(s) (non-unique solution).

• When this situation became clear, neural networks were first seen as a “research dead end”.
General (Artificial) Neural Networks
Basic graph theoretic notions

A (directed) graph is a pair $G = (V, E)$ consisting of a (finite) set $V$ of vertices or nodes and a (finite) set $E \subseteq V \times V$ of edges.

We call an edge $e = (u, v) \in E$ directed from vertex $u$ to vertex $v$.

Let $G = (V, E)$ be a (directed) graph and $u \in V$ a vertex. Then the vertices of the set

$$\text{pred}(u) = \{v \in V \mid (v, u) \in E\}$$

are called the predecessors of the vertex $u$ and the vertices of the set

$$\text{succ}(u) = \{v \in V \mid (u, v) \in E\}$$

are called the successors of the vertex $u$. 
General definition of a neural network

An (artificial) **neural network** is a (directed) graph \( G = (U, C) \), whose vertices \( u \in U \) are called **neurons** or **units** and whose edges \( c \in C \) are called **connections**.

The set \( U \) of vertices is partitioned into

- the set \( U_{\text{in}} \) of **input neurons**,
- the set \( U_{\text{out}} \) of **output neurons**, and
- the set \( U_{\text{hidden}} \) of **hidden neurons**.

It is

\[
U = U_{\text{in}} \cup U_{\text{out}} \cup U_{\text{hidden}},
\]

\[
U_{\text{in}} \neq \emptyset, \quad U_{\text{out}} \neq \emptyset, \quad U_{\text{hidden}} \cap (U_{\text{in}} \cup U_{\text{out}}) = \emptyset.
\]
Each connection \((v, u) \in C\) possesses a weight \(w_{uv}\) and each neuron \(u \in U\) possesses three (real-valued) state variables:

- the **network input** \(\text{net}_u\),
- the **activation** \(\text{act}_u\), and
- the **output** \(\text{out}_u\).

Each input neuron \(u \in U_{\text{in}}\) also possesses a fourth (real-valued) state variable,

- the **external input** \(\text{ext}_u\).

Furthermore, each neuron \(u \in U\) possesses three functions:

- the **network input function** \(f_{\text{net}}^{(u)} : \mathbb{R}^2 | \text{pred}(u)| + \kappa_1(u) \rightarrow \mathbb{R}\),
- the **activation function** \(f_{\text{act}}^{(u)} : \mathbb{R}^{\kappa_2(u)} \rightarrow \mathbb{R}\), and
- the **output function** \(f_{\text{out}}^{(u)} : \mathbb{R} \rightarrow \mathbb{R}\),

which are used to compute the values of the state variables.
Types of (artificial) neural networks:

- If the graph of a neural network is acyclic, it is called a feed-forward network.
- If the graph of a neural network contains cycles (backward connections), it is called a recurrent network.

Representation of the connection weights as a matrix:

\[
\begin{pmatrix}
 u_1 & u_2 & \ldots & u_r \\
 w_{u_1u_1} & w_{u_1u_2} & \ldots & w_{u_1u_r} \\
 w_{u_2u_1} & w_{u_2u_2} & \ldots & w_{u_2u_r} \\
 \vdots & \vdots & \ddots & \vdots \\
 w_{u_ru_1} & w_{u_ru_2} & \ldots & w_{u_ru_r}
\end{pmatrix}
\begin{pmatrix}
 u_1 \\
 u_2 \\
 \vdots \\
 u_r
\end{pmatrix}
\]
A simple recurrent neural network

Weight matrix of this network

\[
\begin{pmatrix}
0 & 0 & 4 \\
1 & 0 & 0 \\
-2 & 3 & 0 \\
\end{pmatrix}
\]
A generalized neuron is a simple numeric processor

\[ \text{out}_{v_1} = \text{in}_{uv_1} \]

\[ w_{uv_1} \]

\[ \vdots \]

\[ \vdots \]

\[ \vdots \]

\[ \text{out}_{v_n} = \text{in}_{uv_n} \]

\[ w_{uv_n} \]

\[ \sigma_1, \ldots, \sigma_l \]

\[ \theta_1, \ldots, \theta_k \]

\[ f_{\text{net}}(u) \]

\[ \text{net}_u \]

\[ f_{\text{act}}(u) \]

\[ \text{act}_u \]

\[ f_{\text{out}}(u) \]

\[ \text{ext}_u \]
General Neural Networks: Example

\[ f^{(u)}(\bar{w}_u, \bar{n}_u) = \sum_{v \in \text{pred}(u)} w_{uv} \text{in}_{uv} = \sum_{v \in \text{pred}(u)} w_{uv} \text{out}_{v} \]

\[ f^{(u)}(\text{net}_u, \theta) = \begin{cases} 1, & \text{if } \text{net}_u \geq \theta, \\ 0, & \text{otherwise}. \end{cases} \]

\[ f^{(u)}(\text{act}_u) = \text{act}_u \]
Updating the activations of the neurons

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- Order in which the neurons are updated:
  $u_3, u_1, u_2, u_3, u_1, u_2, u_3, \ldots$

- Input phase: activations/outputs in the initial state.

- Work phase: activations/outputs of the next neuron to update (bold) are computed from the outputs of the other neurons and the weights/threshold.

- A stable state with a unique output is reached.
Updating the activations of the neurons

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- Order in which the neurons are updated:
  - $u_3, u_2, u_1, u_3, u_2, u_1, u_3, \ldots$
- No stable state is reached (oscillation of output).
Definition of learning tasks for a neural network

A fixed learning task $L_{\text{fixed}}$ for a neural network with

- $n$ input neurons $U_{\text{in}} = \{u_1, \ldots, u_n\}$ and
- $m$ output neurons $U_{\text{out}} = \{v_1, \ldots, v_m\}$,

is a set of training patterns $l = (\vec{\mathbf{i}}^{(l)}, \vec{\mathbf{o}}^{(l)})$, each consisting of

- an input vector $\vec{\mathbf{i}}^{(l)} = (\mathbf{ext}_u^{(l)} u_1, \ldots, \mathbf{ext}_u^{(l)} u_n)$ and
- an output vector $\vec{\mathbf{o}}^{(l)} = (\mathbf{o}_v^{(l)} v_1, \ldots, \mathbf{o}_v^{(l)} v_m)$.

A fixed learning task is solved, if for all training patterns $l \in L_{\text{fixed}}$ the neural network computes from the external inputs contained in the input vector $\vec{\mathbf{i}}^{(l)}$ of a training pattern $l$ the outputs contained in the corresponding output vector $\vec{\mathbf{o}}^{(l)}$. 
Solving a fixed learning task: Error definition

- Measure how well a neural network solves a given fixed learning task.
- Compute differences between desired and actual outputs.
- Do not sum differences directly in order to avoid errors canceling each other.
- Square has favorable properties for deriving the adaptation rules.

\[
e = \sum_{l \in L_{\text{fixed}}} e^{(l)} = \sum_{v \in U_{\text{out}}} e_{v} = \sum_{l \in L_{\text{fixed}}} \sum_{v \in U_{\text{out}}} e_{v}^{(l)},
\]

where \( e_{v}^{(l)} = \left( o_{v}^{(l)} - \text{out}_{v}^{(l)} \right)^{2} \)
Definition of learning tasks for a neural network

A free learning task $L_{\text{free}}$ for a neural network with

- $n$ input neurons $U_{\text{in}} = \{u_1, \ldots, u_n\}$,

is a set of training patterns $l = (\vec{v}(l))$, each consisting of

- an input vector $\vec{v}(l) = (\text{ext}^{(l)}_{u_1}, \ldots, \text{ext}^{(l)}_{u_n})$.

Properties:

- There is no desired output for the training patterns.
- Outputs can be chosen freely by the training method.
- Solution idea: Similar inputs should lead to similar outputs. (clustering of input vectors)
Normalization of the input vectors

- Compute expected value and (corrected) standard deviation for each input:

\[ \mu_k = \frac{1}{|L|} \sum_{l \in L} \text{ext}_{u_k}^{(l)} \quad \text{and} \quad \sigma_k = \sqrt{\frac{1}{|L| - 1} \sum_{l \in L} \left( \text{ext}_{u_k}^{(l)} - \mu_k \right)^2} , \]

- Normalize the input vectors to expected value 0 and standard deviation 1:

\[ \text{ext}_{u_k}^{(l)(new)} = \frac{\text{ext}_{u_k}^{(l)(old)} - \mu_k}{\sigma_k} \]

- Such a normalization avoids unit and scaling problems. It is also known as z-scaling or z-score standardization.
Multi-layer Perceptrons (MLPs)
An \textbf{r-layer perceptron} is a neural network with a graph \( G = (U, C) \) that satisfies the following conditions:

\begin{enumerate}
  \item \( U_{\text{in}} \cap U_{\text{out}} = \emptyset \),
  \item \( U_{\text{hidden}} = U_{\text{hidden}}^{(1)} \cup \cdots \cup U_{\text{hidden}}^{(r-2)} \)
    \quad \forall 1 \leq i < j \leq r - 2 : \quad U_{\text{hidden}}^{(i)} \cap U_{\text{hidden}}^{(j)} = \emptyset,
  \item \( C \subseteq \left( U_{\text{in}} \times U_{\text{hidden}}^{(1)} \right) \cup \left( \bigcup_{i=1}^{r-3} U_{\text{hidden}}^{(i)} \times U_{\text{hidden}}^{(i+1)} \right) \cup \left( U_{\text{hidden}}^{(r-2)} \times U_{\text{out}} \right)
    \quad \text{or, if there are no hidden neurons} \quad (r = 2, U_{\text{hidden}} = \emptyset),
    \quad C \subseteq U_{\text{in}} \times U_{\text{out}}.
\end{enumerate}

- Feed-forward network with strictly layered structure.
General structure of a multi-layer perceptron
Multi-layer Perceptrons

- The network input function of each hidden neuron and of each output neuron is the **weighted sum** of its inputs, that is,

\[ \forall u \in U_{\text{hidden}} \cup U_{\text{out}} : \quad f_{\text{net}}^{(u)}(\vec{w}_u, \vec{in}_u) = \vec{w}_u \vec{in}_u = \sum_{v \in \text{pred}(u)} w_{uv} \text{out}_v. \]

- The activation function of each hidden neuron is a so-called **sigmoid function**, that is, a monotonically increasing function

\[ f : \mathbb{R} \rightarrow [0, 1] \quad \text{with} \quad \lim_{x \to -\infty} f(x) = 0 \quad \text{and} \quad \lim_{x \to \infty} f(x) = 1. \]

- The activation function of each output neuron is either also a sigmoid function or a **linear function**, that is,

\[ f_{\text{act}}(\text{net}, \theta) = \alpha \text{net} - \theta. \]

Only the step function is a neurobiologically plausible activation function.
Sigmoid Activation Functions

step function:
\[ f_{\text{act}}(\text{net}, \theta) = \begin{cases} 
1, & \text{if net} \geq \theta, \\
0, & \text{otherwise}. 
\end{cases} \]

semi-linear function:
\[ f_{\text{act}}(\text{net}, \theta) = \begin{cases} 
1, & \text{if net} > \theta + \frac{1}{2}, \\
0, & \text{if net} < \theta - \frac{1}{2}, \\
(\text{net} - \theta) + \frac{1}{2}, & \text{otherwise}. 
\end{cases} \]

sine until saturation:
\[ f_{\text{act}}(\text{net}, \theta) = \begin{cases} 
1, & \text{if net} > \theta + \frac{\pi}{2}, \\
0, & \text{if net} < \theta - \frac{\pi}{2}, \\
\sin(\text{net} - \theta) + \frac{1}{2}, & \text{otherwise}. 
\end{cases} \]

logistic function:
\[ f_{\text{act}}(\text{net}, \theta) = \frac{1}{1 + e^{-(\text{net} - \theta)}} \]
• All sigmoid functions on the previous slide are **unipolar**, that is, they range from 0 to 1.

• Sometimes **bipolar** sigmoid functions are used (ranging from $-1$ to $+1$), like the hyperbolic tangent (**tangens hyperbolicus**).

**hyperbolic tangent:**

$$f_{\text{act}}(\text{net}, \theta) = \tanh(\text{net} - \theta)$$

$$= \frac{e^{(\text{net} - \theta)} - e^{-(\text{net} - \theta)}}{e^{(\text{net} - \theta)} + e^{-(\text{net} - \theta)}}$$

$$= \frac{1 - e^{-2(\text{net} - \theta)}}{1 + e^{-2(\text{net} - \theta)}}$$

$$= \frac{2}{1 + e^{-2(\text{net} - \theta)}} - 1$$
Let \( U_1 = \{v_1, \ldots, v_m\} \) and \( U_2 = \{u_1, \ldots, u_n\} \) be the neurons of two consecutive layers of a multi-layer perceptron.

Their connection weights are represented by an \( n \times m \) matrix

\[
W = \begin{pmatrix}
  w_{u_1v_1} & w_{u_1v_2} & \cdots & w_{u_1v_m} \\
  w_{u_2v_1} & w_{u_2v_2} & \cdots & w_{u_2v_m} \\
  \vdots & \vdots & & \vdots \\
  w_{u_nv_1} & w_{u_nv_2} & \cdots & w_{u_nv_m}
\end{pmatrix},
\]

where \( w_{u_iv_j} = 0 \) if there is no connection from neuron \( v_j \) to neuron \( u_i \).

Advantage: The computation of the network input can be written as

\[
\vec{\text{net}}_{U_2} = W \cdot \vec{\text{in}}_{U_2} = W \cdot \vec{\text{out}}_{U_1}
\]

where \( \vec{\text{net}}_{U_2} = (\text{net}_{u_1}, \ldots, \text{net}_{u_n})^\top \) and \( \vec{\text{in}}_{U_2} = \vec{\text{out}}_{U_1} = (\text{out}_{v_1}, \ldots, \text{out}_{v_m})^\top \).
Solving the biimplication problem with a multi-layer perceptron.

![Diagram of a multi-layer perceptron solving the biimplication problem](image)

Note the additional input neurons compared to the TLU solution.

\[
\mathbf{W}_1 = \begin{pmatrix} -2 & 2 \\ 2 & -2 \end{pmatrix} \quad \text{and} \quad \mathbf{W}_2 = \begin{pmatrix} 2 & 2 \end{pmatrix}
\]
Multi-layer Perceptrons: Fredkin Gate

\[ \begin{align*}
  s &= 0 0 0 0 1 1 1 1 1 \\
  x_1 &= 0 0 1 1 0 0 1 1 \\
  x_2 &= 0 1 0 1 0 1 0 1 \\
  y_1 &= 0 0 1 1 0 1 0 1 \\
  y_2 &= 0 1 0 1 0 0 1 1
\end{align*} \]
The Fredkin gate (after Edward Fredkin *1934) or controlled swap gate (CSWAP) is a computational circuit that is used in conservative logic and reversible computing.

Conservative logic is a model of computation that explicitly reflects the physical properties of computation, like the reversibility of the dynamical laws and the conservation of certain quantities (e.g. energy) [Fredkin and Toffoli 1982].

The Fredkin gate is reversible in the sense that the inputs can be computed as functions of the outputs in the same way in which the outputs can be computed as functions of the inputs (no information loss, no entropy gain).

The Fredkin gate is universal in the sense that all Boolean functions can be computed using only Fredkin gates.

Note that both outputs, $y_1$ and $y_2$ are not linearly separable, because the convex hull of the points mapped to 0 and the convex hull of the points mapped to 1 share the point in the center of the cube.
Theorem: Two sets of points in a Euclidean space are **linearly separable** if and only if their convex hulls are disjoint (that is, have no point in common).

Both outputs $y_1$ and $y_2$ of a Fredkin gate are not linearly separable:

- Convex hull of points with $y_1 = 0$
- Convex hull of points with $y_1 = 1$
- Convex hull of points with $y_2 = 0$
- Convex hull of points with $y_2 = 1$
• A billiard-ball computer (a.k.a. conservative logic circuit) is an idealized model of a reversible mechanical computer that simulates the computations of circuits by the movements of spherical billiard balls in a frictionless environment [Fredkin and Toffoli 1982].

**Switch** (control output location of an input ball by another ball; equivalent to an AND gate in its second/lower output):
Excursion: The (Negated) Fredkin Gate

- Only reflectors (black) and inputs/outputs (color) shown.
- Meaning of switch variable is negated compared to value table on previous slide: 0 switches inputs, 1 passes them through.
Excursion: The (Negated) Fredkin Gate

• All possible paths of billiard-balls through this gate are shown. Inputs/outputs are in color; red: switch, green: $x_1/y_1$, blue: $x_2/y_2$.

• Billiard-balls are shown in places where their trajectories may change direction.
This Fredkin gate implementation contains four switches (shown in light blue).

The other parts serve the purpose to equate the travel times of the balls through the gate.
Excursion: The (Negated) Fredkin Gate

- Reminder: in this implementation the meaning of the switch variable is negated.
- Switch is zero (no ball enters at red arrow):
  blue and green balls switch to the other output.
Excursion: The (Negated) Fredkin Gate

- If a ball is entered at the switch input, it is passed through.
- Since there are no other inputs, the other outputs remain 0 (no ball in, no ball out).
Excursion: The (Negated) Fredkin Gate

- A ball is entered at the switch input and passed through.
- A ball entered at the first input $x_1$ is also passed through to output $y_1$.
  (Note the symmetry of the trajectories.)
Excursion: The (Negated) Fredkin Gate

- A ball is entered at the switch input and passed through.
- A ball entered at the second input $x_2$ is also passed through to output $y_2$. (Note the symmetry of the trajectories.)
Excursion: The (Negated) Fredkin Gate

- A ball is entered at the switch input and passed through.
- Ball entered at both inputs $x_1$ and $x_2$ and are also passed through to the outputs $y_1$ and $y_2$. (Note the symmetry of the trajectories.)
Multi-layer Perceptrons: Fredkin Gate

\[ W_1 = \begin{pmatrix} 2 & -2 & 0 \\ 2 & 2 & 0 \\ 0 & 2 & 2 \\ 0 & -2 & 2 \end{pmatrix} \]

\[ W_2 = \begin{pmatrix} 2 & 0 & 2 & 0 \\ 0 & 2 & 0 & 2 \end{pmatrix} \]
Why Non-linear Activation Functions?

With weight matrices we have for two consecutive layers $U_1$ and $U_2$

$$\text{net}_{U_2} = \mathbf{W} \cdot \text{in}_{U_2} = \mathbf{W} \cdot \text{out}_{U_1}.$$ 

If the activation functions are linear, that is,

$$f_{\text{act}}(\text{net}, \theta) = \alpha \text{net} - \theta.$$

the activations of the neurons in the layer $U_2$ can be computed as

$$\mathbf{a}_{ctU_2} = \mathbf{D}_{\text{act}} \cdot \text{net}_{U_2} - \mathbf{\theta},$$

where

- $\mathbf{a}_{ctU_2} = (a_{ctu_1}, \ldots, a_{ctu_n})^\top$ is the activation vector,
- $\mathbf{D}_{\text{act}}$ is an $n \times n$ diagonal matrix of the factors $\alpha_{u_i}$, $i = 1, \ldots, n$, and
- $\mathbf{\theta} = (\theta_{u_1}, \ldots, \theta_{u_n})^\top$ is a bias vector.
Why Non-linear Activation Functions?

If the output function is also linear, it is analogously

\[ \text{out}_{U_2} = D_{out} \cdot \text{act}_{U_2} - \xi, \]

where

- \( \text{out}_{U_2} = (\text{out}_{u_1}, \ldots, \text{out}_{u_n})^\top \) is the output vector,
- \( D_{out} \) is again an \( n \times n \) diagonal matrix of factors, and
- \( \xi = (\xi_{u_1}, \ldots, \xi_{u_n})^\top \) a bias vector.

Combining these computations we get

\[ \text{out}_{U_2} = D_{out} \cdot (D_{act} \cdot (W \cdot \text{out}_{U_1}) - \theta) - \xi \]

and thus

\[ \text{out}_{U_2} = A_{12} \cdot \text{out}_{U_1} + \tilde{b}_{12} \]

with an \( n \times m \) matrix \( A_{12} \) and an \( n \)-dimensional vector \( \tilde{b}_{12} \).
Therefore we have

\[ \text{out}_{U_2} = A_{12} \cdot \text{out}_{U_1} + \vec{b}_{12} \]

and

\[ \text{out}_{U_3} = A_{23} \cdot \text{out}_{U_2} + \vec{b}_{23} \]

for the computations of two consecutive layers \( U_2 \) and \( U_3 \).

These two computations can be combined into

\[ \text{out}_{U_3} = A_{13} \cdot \text{out}_{U_1} + \vec{b}_{13}, \]

where \( A_{13} = A_{23} \cdot A_{12} \) and \( \vec{b}_{13} = A_{23} \cdot \vec{b}_{12} + \vec{b}_{23} \).

**Result:** With linear activation and output functions any multi-layer perceptron can be reduced to a two-layer perceptron.
Multi-layer Perceptrons: Function Approximation

- Up to now: representing and learning Boolean functions $f : \{0, 1\}^n \rightarrow \{0, 1\}$.
- Now: representing and learning real-valued functions $f : \mathbb{R}^n \rightarrow \mathbb{R}$.

General idea of function approximation:

- Approximate a given function by a step function.
- Construct a neural network that computes the step function.
A neural network that computes the step function shown on the preceding slide. According to the input value only one step is active at any time. The output neuron has the identity as its activation and output functions.
Theorem: Any Riemann-integrable function can be approximated with arbitrary accuracy by a four-layer perceptron.

• But: Error is measured as the area between the functions.

• More sophisticated mathematical examination allows a stronger assertion: With a three-layer perceptron any continuous function can be approximated with arbitrary accuracy (error: maximum function value difference).
Universal Approximation Theorem [Hornik 1991]:

Let $\varphi(\cdot)$ be a continuous, bounded and nonconstant function, let $X$ denote an arbitrary compact subset of $\mathbb{R}^m$, and let $C(X)$ denote the space of continuous functions on $X$.

Given any function $f \in C(X)$ and $\varepsilon > 0$, there exists an integer $N$, real constants $v_i, \theta_i \in \mathbb{R}$ and real vectors $\vec{w}_i \in \mathbb{R}^m$, $i = 1, \ldots, N$, such that we may define

$$F(\vec{x}) = \sum_{i=1}^{N} v_i \varphi(\vec{w}_i^\top \vec{x} - \theta_i)$$

as an approximate realization of the function $f$ where $f$ is independent of $\varphi$. That is,

$$|F(\vec{x}) - f(\vec{x})| < \varepsilon$$

for all $\vec{x} \in X$. In other words, functions of the form $F(\vec{x})$ are dense in $C(X)$.

Note that it is not the shape of the activation function, but the layered structure of the feedforward network that renders multi-layer perceptrons universal approximators.
Multi-layer Perceptrons: Function Approximation

By using relative step heights one layer can be saved.
A neural network that computes the step function shown on the preceding slide. The output neuron has the identity as its activation and output functions.
By using semi-linear functions the approximation can be improved.
A neural network that computes the step function shown on the preceding slide. The output neuron has the identity as its activation and output functions.

\[ \theta_i = \frac{x_i}{\Delta x} \]

\[ \Delta x = x_{i+1} - x_i \]
Mathematical Background: Regression
Training neural networks is closely related to regression.

Given:
- A dataset \( ((x_1, y_1), \ldots, (x_n, y_n)) \) of \( n \) data tuples and
- a hypothesis about the functional relationship, e.g. \( y = g(x) = a + bx \).

Approach: Minimize the sum of squared errors, that is,

\[
F(a, b) = \sum_{i=1}^{n} (g(x_i) - y_i)^2 = \sum_{i=1}^{n} (a + bx_i - y_i)^2.
\]

Necessary conditions for a minimum
(a.k.a. Fermat’s theorem, after Pierre de Fermat, 1601–1665):

\[
\frac{\partial F}{\partial a} = \sum_{i=1}^{n} 2(a + bx_i - y_i) = 0 \quad \text{and} \\
\frac{\partial F}{\partial b} = \sum_{i=1}^{n} 2(a + bx_i - y_i)x_i = 0
\]
Result of necessary conditions: System of so-called **normal equations**, that is,

\[ na + \left( \sum_{i=1}^{n} x_i \right) b = \sum_{i=1}^{n} y_i, \]

\[ \left( \sum_{i=1}^{n} x_i \right) a + \left( \sum_{i=1}^{n} x_i^2 \right) b = \sum_{i=1}^{n} x_i y_i. \]

- Two linear equations for two unknowns \( a \) and \( b \).
- System can be solved with standard methods from linear algebra.
- Solution is unique unless all \( x \)-values are identical.
- The resulting line is called a **regression line**.
Linear Regression: Example

\[ y = \frac{3}{4} + \frac{7}{12}x. \]

<table>
<thead>
<tr>
<th>$x$</th>
<th>$1$</th>
<th>$2$</th>
<th>$3$</th>
<th>$4$</th>
<th>$5$</th>
<th>$6$</th>
<th>$7$</th>
<th>$8$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$y$</td>
<td>$1$</td>
<td>$3$</td>
<td>$2$</td>
<td>$3$</td>
<td>$4$</td>
<td>$3$</td>
<td>$5$</td>
<td>$6$</td>
</tr>
</tbody>
</table>
Generalization to polynomials

\[ y = p(x) = a_0 + a_1 x + \ldots + a_m x^m \]

Approach: Minimize the sum of squared errors, that is,

\[ F(a_0, a_1, \ldots, a_m) = \sum_{i=1}^{n} (p(x_i) - y_i)^2 = \sum_{i=1}^{n} (a_0 + a_1 x_i + \ldots + a_m x_i^m - y_i)^2 \]

Necessary conditions for a minimum: All partial derivatives vanish, that is,

\[ \frac{\partial F}{\partial a_0} = 0, \quad \frac{\partial F}{\partial a_1} = 0, \quad \ldots, \quad \frac{\partial F}{\partial a_m} = 0. \]
System of normal equations for polynomials

\[
n a_0 + \left( \sum_{i=1}^{n} x_i \right) a_1 + \ldots + \left( \sum_{i=1}^{n} x_i^m \right) a_m = \sum_{i=1}^{n} y_i
\]

\[
\left( \sum_{i=1}^{n} x_i \right) a_0 + \left( \sum_{i=1}^{n} x_i^2 \right) a_1 + \ldots + \left( \sum_{i=1}^{n} x_i^{m+1} \right) a_m = \sum_{i=1}^{n} x_i y_i
\]

\[
\vdots
\]

\[
\left( \sum_{i=1}^{n} x_i^m \right) a_0 + \left( \sum_{i=1}^{n} x_i^{m+1} \right) a_1 + \ldots + \left( \sum_{i=1}^{n} x_i^{2m} \right) a_m = \sum_{i=1}^{n} x_i^m y_i
\]

- \( m + 1 \) linear equations for \( m + 1 \) unknowns \( a_0, \ldots, a_m \).
- System can be solved with standard methods from linear algebra.
- Solution is unique unless the points lie exactly on a polynomial of lower degree.
Mathematical Background: Multilinear Regression

Generalization to more than one argument

$$z = f(x, y) = a + bx + cy$$

Approach: Minimize the sum of squared errors, that is,

$$F(a, b, c) = \sum_{i=1}^{n} (f(x_i, y_i) - z_i)^2 = \sum_{i=1}^{n} (a + bx_i + cy_i - z_i)^2$$

Necessary conditions for a minimum: All partial derivatives vanish, that is,

$$\frac{\partial F}{\partial a} = \sum_{i=1}^{n} 2(a + bx_i + cy_i - z_i) = 0,$$

$$\frac{\partial F}{\partial b} = \sum_{i=1}^{n} 2(a + bx_i + cy_i - z_i)x_i = 0,$$

$$\frac{\partial F}{\partial c} = \sum_{i=1}^{n} 2(a + bx_i + cy_i - z_i)y_i = 0.$$
System of normal equations for several arguments

\[na + \left(\sum_{i=1}^{n} x_i\right)b + \left(\sum_{i=1}^{n} y_i\right)c = \sum_{i=1}^{n} z_i\]

\[\left(\sum_{i=1}^{n} x_i\right)a + \left(\sum_{i=1}^{n} x_i^2\right)b + \left(\sum_{i=1}^{n} x_i y_i\right)c = \sum_{i=1}^{n} z_i x_i\]

\[\left(\sum_{i=1}^{n} y_i\right)a + \left(\sum_{i=1}^{n} x_i y_i\right)b + \left(\sum_{i=1}^{n} y_i^2\right)c = \sum_{i=1}^{n} z_i y_i\]

- 3 linear equations for 3 unknowns \(a\), \(b\), and \(c\).
- System can be solved with standard methods from linear algebra.
- Solution is unique unless all data points lie on a straight line.
General multilinear case:

\[ y = f(x_1, \ldots, x_m) = a_0 + \sum_{k=1}^{m} a_k x_k \]

Approach: Minimize the sum of squared errors, that is,

\[ F(\vec{a}) = (X\vec{a} - \vec{y})^\top (X\vec{a} - \vec{y}) , \]

where

\[ X = \begin{pmatrix} 1 & x_{11} & \cdots & x_{1m} \\ \vdots & \vdots & \ddots & \vdots \\ 1 & x_{1n} & \cdots & x_{mn} \end{pmatrix}, \quad \vec{y} = \begin{pmatrix} y_1 \\ \vdots \\ y_n \end{pmatrix}, \quad \text{and} \quad \vec{a} = \begin{pmatrix} a_0 \\ a_1 \\ \vdots \\ a_m \end{pmatrix} \]

Necessary conditions for a minimum:

\[ \nabla_{\vec{a}} F(\vec{a}) = \nabla_{\vec{a}} (X\vec{a} - \vec{y})^\top (X\vec{a} - \vec{y}) = 0 \]
Multilinear Regression

- $\tilde{\nabla}_a F(\tilde{a})$ may easily be computed by remembering that the differential operator

$$\tilde{\nabla}_a = \left( \frac{\partial}{\partial a_0}, \ldots, \frac{\partial}{\partial a_m} \right)$$

behaves formally like a vector that is “multiplied” to the sum of squared errors.

- Alternatively, one may write out the differentiation componentwise.

With the former method we obtain for the derivative:

$$\tilde{\nabla}_a F(\tilde{a}) = \tilde{\nabla}_a ((X\tilde{a} - \tilde{y})^\top (X\tilde{a} - \tilde{y}))$$

$$= \left( \tilde{\nabla}_a (X\tilde{a} - \tilde{y}) \right)^\top (X\tilde{a} - \tilde{y}) + ((X\tilde{a} - \tilde{y})^\top \left( \tilde{\nabla}_a (X\tilde{a} - \tilde{y}) \right))^\top$$

$$= \left( \tilde{\nabla}_a (X\tilde{a} - \tilde{y}) \right)^\top (X\tilde{a} - \tilde{y}) + \left( \tilde{\nabla}_a (X\tilde{a} - \tilde{y}) \right)^\top (X\tilde{a} - \tilde{y})$$

$$= 2X^\top (X\tilde{a} - \tilde{y})$$

$$= 2X^\top X\tilde{a} - 2X^\top \tilde{y} = \tilde{0}$$
Multilinear Regression

Necessary condition for a minimum therefore:

$$\nabla_{\vec{a}} F(\vec{a}) = \nabla_{\vec{a}} (X\vec{a} - \vec{y})^\top (X\vec{a} - \vec{y})$$

$$= 2X^\top X\vec{a} - 2X^\top \vec{y} \triangleq 0$$

As a consequence we obtain the system of normal equations:

$$X^\top X\vec{a} = X^\top \vec{y}$$

This system has a solution unless $X^\top X$ is singular. If it is regular, we have

$$\vec{a} = (X^\top X)^{-1}X^\top \vec{y}.$$  

$(X^\top X)^{-1}X^\top$ is called the (Moore-Penrose-)Pseudoinverse of the matrix $X$.

With the matrix-vector representation of the regression problem an extension to multipolynomial regression is straightforward: Simply add the desired products of powers to the matrix $X$. 
Generalization to non-polynomial functions

Simple example: \( y = ax^b \)

Idea: Find transformation to linear/polynomial case.

Transformation for the above example: \( \ln y = \ln a + b \cdot \ln x \).

Special case: **logistic function**

\[
y = \frac{Y}{1 + e^{a+bx}} \quad \Leftrightarrow \quad \frac{1}{y} = \frac{1 + e^{a+bx}}{Y} \quad \Leftrightarrow \quad \frac{Y - y}{y} = e^{a+bx}.
\]

Result: Apply so-called **Logit-Transformation**

\[
\ln \left( \frac{Y - y}{y} \right) = a + bx.
\]
Logistic Regression: Example

<table>
<thead>
<tr>
<th>$x$</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
</tr>
</thead>
<tbody>
<tr>
<td>$y$</td>
<td>0.4</td>
<td>1.0</td>
<td>3.0</td>
<td>5.0</td>
<td>5.6</td>
</tr>
</tbody>
</table>

Transform the data with

$$z = \ln \left( \frac{Y - y}{y} \right), \quad Y = 6.$$ 

The transformed data points are

<table>
<thead>
<tr>
<th>$x$</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
</tr>
</thead>
<tbody>
<tr>
<td>$z$</td>
<td>2.64</td>
<td>1.61</td>
<td>0.00</td>
<td>−1.61</td>
<td>−2.64</td>
</tr>
</tbody>
</table>

The resulting regression line and therefore the desired function are

$$z \approx -1.3775x + 4.133 \quad \text{and} \quad y \approx \frac{6}{1 + e^{-1.3775x+4.133}}.$$ 

**Attention:** Note that the error is minimized only in the transformed space! Therefore the function in the original space may not be optimal!
The logistic regression function can be computed by a single neuron with

- network input function $f_{\text{net}}(x) \equiv wx$ with $w \approx -1.3775$,
- activation function $f_{\text{act}}(\text{net}, \theta) \equiv \left(1 + e^{-(\text{net} - \theta)}\right)^{-1}$ with $\theta \approx 4.133$ and
- output function $f_{\text{out}}(\text{act}) \equiv 6 \text{act}$. 
Logistic Function: Two-dimensional Example

Example logistic function for two arguments $x_1$ and $x_2$:

$$ y = \frac{1}{1 + \exp(4 - x_1 - x_2)} = \frac{1}{1 + \exp\left(4 - (1, 1)^\top (x_1, x_2)\right)} $$

The blue lines have show where the logistic function has a certain value in $\{0.1, \ldots, 0.9\}$. 
Logistic Regression: Two Class Problems

- Let $C$ be a class attrib., $\text{dom}(C) = \{c_1, c_2\}$, and $\vec{X}$ an $m$-dim. random vector.
  Let $P(C = c_1 \mid \vec{X} = \vec{x}) = p(\vec{x})$ and $P(C = c_2 \mid \vec{X} = \vec{x}) = 1 - p(\vec{x})$.

- **Given:** A set of data points $X = \{\vec{x}_1, \ldots, \vec{x}_n\}$ (realizations of $\vec{X}$), each of which belongs to one of the two classes $c_1$ and $c_2$.

- **Desired:** A simple description of the function $p(\vec{x})$.

- **Approach:** Describe $p$ by a logistic function:

  $$ p(\vec{x}) = \frac{1}{1 + e^{a_0 + \vec{a} \cdot \vec{x}}} = \frac{1}{1 + \exp(a_0 + \sum_{i=1}^{m} a_i x_i)} $$

  Apply logit transformation to $p(x)$:

  $$ \ln \left( \frac{1 - p(\vec{x})}{p(\vec{x})} \right) = a_0 + \vec{a} \cdot \vec{x} = a_0 + \sum_{i=1}^{m} a_i x_i $$

  The values $p(\vec{x}_i)$ may be obtained by kernel estimation (see next slide).
Logistic Regression: Kernel Estimation

- **Idea:** Define an “influence function” (kernel), which describes how strongly a data point influences the probability estimate for neighboring points.

- **Common choice for the kernel function:** Gaussian function

\[
K(\vec{x}, \vec{y}) = \frac{1}{(2\pi \sigma^2)^{\frac{m}{2}}} \exp \left(-\frac{(\vec{x} - \vec{y})^\top (\vec{x} - \vec{y})}{2\sigma^2}\right)
\]

- Kernel estimate of probability density given a data set \( X = \{\vec{x}_1, \ldots, \vec{x}_n\} \):

\[
\hat{f}(\vec{x}) = \frac{1}{n} \sum_{i=1}^{n} K(\vec{x}, \vec{x}_i).
\]

- Kernel estimation applied to a two class problem:

\[
\hat{p}(\vec{x}) = \frac{\sum_{i=1}^{n} c(\vec{x}_i) K(\vec{x}, \vec{x}_i)}{\sum_{i=1}^{n} K(\vec{x}, \vec{x}_i)}.
\]

It is \( c(\vec{x}_i) = 1 \) if \( x_i \) belongs to class \( c_1 \) and \( c(\vec{x}_i) = 0 \) otherwise.
Training Multi-layer Perceptrons
Training Multi-layer Perceptrons: Gradient Descent

- Problem of logistic regression: Works only for two-layer perceptrons.
- More general approach: gradient descent.
- Necessary condition: differentiable activation and output functions.

Illustration of the gradient of a real-valued function $z = f(x, y)$ at a point $(x_0, y_0)$.

It is $\nabla z|_{(x_0, y_0)} = \left( \frac{\partial z}{\partial x}|_{x_0}, \frac{\partial z}{\partial y}|_{y_0} \right)$. 

($\nabla$ is a differential operator called “nabla” or “del”.)
Gradient Descent: Formal Approach

**General Idea:** Approach the minimum of the error function in small steps.

Error function:

\[
e = \sum_{l \in L_{\text{fixed}}} e^{(l)} = \sum_{v \in U_{\text{out}}} e_v = \sum_{l \in L_{\text{fixed}}} \sum_{v \in U_{\text{out}}} e_v^{(l)},
\]

Form gradient to determine the direction of the step (here and in the following: extended weight vector \( \vec{w}_u = (-\theta_u, w_{up1}, \ldots, w_{upn}) \)):

\[
\nabla_{\vec{w}_u} e = \frac{\partial e}{\partial \vec{w}_u} = \left( -\frac{\partial e}{\partial \theta_u}, \frac{\partial e}{\partial w_{up1}}, \ldots, \frac{\partial e}{\partial w_{upn}} \right).
\]

Exploit the sum over the training patterns:

\[
\nabla_{\vec{w}_u} e = \frac{\partial e}{\partial \vec{w}_u} = \frac{\partial}{\partial \vec{w}_u} \sum_{l \in L_{\text{fixed}}} e^{(l)} = \sum_{l \in L_{\text{fixed}}} \frac{\partial e^{(l)}}{\partial \vec{w}_u}.
\]
Gradient Descent: Formal Approach

Single pattern error depends on weights only through the network input:

\[
\nabla_{\mathbf{w}_u} e^{(l)} = \frac{\partial e^{(l)}}{\partial \mathbf{w}_u} = \frac{\partial e^{(l)}}{\partial \text{net}^{(l)}_u} \frac{\partial \text{net}^{(l)}_u}{\partial \mathbf{w}_u}.
\]

Since \(\text{net}^{(l)}_u = \mathbf{w}_u^\top \mathbf{in}^{(l)}_u\) (note: extended input vector \(\mathbf{in}^{(l)}_u = (1, \text{in}^{(l)}_{p_1u}, \ldots, \text{in}^{(l)}_{p_nu})\)), we have for the second factor

\[
\frac{\partial \text{net}^{(l)}_u}{\partial \mathbf{w}_u} = \mathbf{in}^{(l)}_u.
\]

For the first factor we consider the error \(e^{(l)}\) for the training pattern \(l = (\mathbf{in}^{(l)}_u, \mathbf{o}^{(l)}_u)\):

\[
e^{(l)} = \sum_{v \in U_{\text{out}}} e^{(l)}_v = \sum_{v \in U_{\text{out}}} \left( o^{(l)}_v - \text{out}^{(l)}_v \right)^2,
\]

that is, the sum of the errors over all output neurons.
Therefore we have

$$\frac{\partial e^{(l)}}{\partial \text{net}^{(l)}_u} = \frac{\partial \sum_{v \in U_{\text{out}}} \left( o_v^{(l)} - \text{out}_v^{(l)} \right)^2}{\partial \text{net}^{(l)}_u} = \sum_{v \in U_{\text{out}}} \frac{\partial \left( o_v^{(l)} - \text{out}_v^{(l)} \right)^2}{\partial \text{net}^{(l)}_u}. $$

Since only the actual output $\text{out}_v^{(l)}$ of an output neuron $v$ depends on the network input $\text{net}_u^{(l)}$ of the neuron $u$ we are considering, it is

$$\frac{\partial e^{(l)}}{\partial \text{net}^{(l)}_u} = -2 \sum_{v \in U_{\text{out}}} \left( o_v^{(l)} - \text{out}_v^{(l)} \right) \frac{\partial \text{out}_v^{(l)}}{\partial \text{net}_u^{(l)}} \delta_u^{(l)},$$

which also introduces the abbreviation $\delta_u^{(l)}$ for the important sum appearing here.
Gradient Descent: Formal Approach

Distinguish two cases:

- The neuron \( u \) is an **output neuron**.
- The neuron \( u \) is a **hidden neuron**.

In the first case we have

\[
\forall u \in U_{\text{out}} : \quad \delta^{(l)}_u = \left( o^{(l)}_u - \text{out}^{(l)}_u \right) \frac{\partial \text{out}^{(l)}_u}{\partial \text{net}^{(l)}_u}
\]

Therefore we have for the gradient

\[
\forall u \in U_{\text{out}} : \quad \vec{\nabla}_{\vec{w}_u} e^{(l)}_u = \frac{\partial e^{(l)}_u}{\partial \vec{w}_u} = -2 \left( o^{(l)}_u - \text{out}^{(l)}_u \right) \frac{\partial \text{out}^{(l)}_u}{\partial \text{net}^{(l)}_u} \vec{\text{in}}^{(l)}_u
\]

and thus for the weight change

\[
\forall u \in U_{\text{out}} : \quad \Delta \vec{w}^{(l)}_u = -\frac{\eta}{2} \vec{\nabla}_{\vec{w}_u} e^{(l)}_u = \eta \left( o^{(l)}_u - \text{out}^{(l)}_u \right) \frac{\partial \text{out}^{(l)}_u}{\partial \text{net}^{(l)}_u} \vec{\text{in}}^{(l)}_u.
\]
Gradient Descent: Formal Approach

Exact formulae depend on the choice of the activation and the output function, since it is
\[ \text{out}^{(l)}_u = f_{\text{out}}(\text{act}^{(l)}_u) = f_{\text{out}}(f_{\text{act}}(\text{net}^{(l)}_u)). \]

Consider the special case with

- output function is the identity,
- activation function is logistic, that is, \( f_{\text{act}}(x) = \frac{1}{1+e^{-x}}. \)

The first assumption yields
\[
\frac{\partial \text{out}^{(l)}_u}{\partial \text{net}^{(l)}_u} = \frac{\partial \text{act}^{(l)}_u}{\partial \text{net}^{(l)}_u} = f'_{\text{act}}(\text{net}^{(l)}_u).
\]
For a logistic activation function we have

\[
f'_{\text{act}}(x) = \frac{d}{dx} \left(1 + e^{-x}\right)^{-1} = -\left(1 + e^{-x}\right)^{-2} (-e^{-x})
\]

\[
= \frac{1 + e^{-x} - 1}{(1 + e^{-x})^2} = \frac{1}{1 + e^{-x}} \left(1 - \frac{1}{1 + e^{-x}}\right)
\]

\[
= f_{\text{act}}(x) \cdot (1 - f_{\text{act}}(x)),
\]

and therefore

\[
f'_{\text{act}}(\text{net}_u^{(l)}) = f_{\text{act}}(\text{net}_u^{(l)}) \cdot \left(1 - f_{\text{act}}(\text{net}_u^{(l)})\right) = \text{out}_u^{(l)} \left(1 - \text{out}_u^{(l)}\right).
\]

The resulting weight change is therefore

\[
\Delta \vec{w}_u^{(l)} = \eta \left(o_u^{(l)} - \text{out}_u^{(l)}\right) \text{out}_u^{(l)} \left(1 - \text{out}_u^{(l)}\right) \vec{m}_u^{(l)},
\]

which makes the computations very simple.
logistic activation function:

\[ f_{\text{act}}(\text{net}^{(l)}_u) = \frac{1}{1 + e^{-\text{net}^{(l)}_u}} \]

derivative of logistic function:

\[ f'_{\text{act}}(\text{net}^{(l)}_u) = f_{\text{act}}(\text{net}^{(l)}_u) \cdot (1 - f_{\text{act}}(\text{net}^{(l)}_u)) \]

- If a logistic activation function is used (shown on left), the weight changes are proportional to \( \lambda^{(l)}_u = \text{out}^{(l)}_u \left(1 - \text{out}^{(l)}_u\right) \) (shown on right; see preceding slide).

- Weight changes are largest, and thus the training speed highest, in the vicinity of \( \text{net}^{(l)}_u = 0 \). Far away from \( \text{net}^{(l)}_u = 0 \), the gradient becomes (very) small (“saturation regions”) and thus training (very) slow.
Example logistic function for two arguments $x_1$ and $x_2$:

$$y = \frac{1}{1 + \exp(4 - x_1 - x_2)} = \frac{1}{1 + \exp\left(4 - (1,1)^\top (x_1, x_2)\right)}$$

The blue lines have show where the logistic function has a certain value in \{0.1, \ldots, 0.9\}. 

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Consider now: The neuron $u$ is a hidden neuron, that is, $u \in U_k$, $0 < k < r - 1$.

The output $\text{out}_v^{(l)}$ of an output neuron $v$ depends on the network input $\text{net}_u^{(l)}$ only indirectly through the successor neurons $\text{succ}(u) = \{s \in U \mid (u, s) \in C\} = \{s_1, \ldots, s_m\} \subseteq U_{k+1}$, namely through their network inputs $\text{net}_s^{(l)}$.

We apply the chain rule to obtain

$$
\delta_u^{(l)} = \sum_{v \in U_{\text{out}}} \sum_{s \in \text{succ}(u)} (o_v^{(l)} - \text{out}_v^{(l)}) \frac{\partial \text{out}_v^{(l)}}{\partial \text{net}_s^{(l)}} \frac{\partial \text{net}_s^{(l)}}{\partial \text{net}_u^{(l)}}.
$$

Exchanging the sums yields

$$
\delta_u^{(l)} = \sum_{s \in \text{succ}(u)} \left( \sum_{v \in U_{\text{out}}} (o_v^{(l)} - \text{out}_v^{(l)}) \frac{\partial \text{out}_v^{(l)}}{\partial \text{net}_u^{(l)}} \right) \frac{\partial \text{net}_s^{(l)}}{\partial \text{net}_u^{(l)}} = \sum_{s \in \text{succ}(u)} \delta_s^{(l)} \frac{\partial \text{net}_s^{(l)}}{\partial \text{net}_u^{(l)}}.
$$
Consider the network input

\[
net_s^{(l)} = \overrightarrow{w_s} \overrightarrow{in}_s^{(l)} = \left( \sum_{p \in \text{pred}(s)} w_{sp} \text{out}_p^{(l)} \right) - \theta_s,
\]

where one element of \( \overrightarrow{in}_s^{(l)} \) is the output \( \text{out}_u^{(l)} \) of the neuron \( u \). Therefore it is

\[
\frac{\partial net_s^{(l)}}{\partial net_u^{(l)}} = \left( \sum_{p \in \text{pred}(s)} w_{sp} \frac{\partial \text{out}_p^{(l)}}{\partial net_u^{(l)}} \right) - \frac{\partial \theta_s}{\partial net_u^{(l)}} = w_{su} \frac{\partial \text{out}_u^{(l)}}{\partial net_u^{(l)}}.
\]

The result is the recursive equation (error backpropagation)

\[
\delta_u^{(l)} = \left( \sum_{s \in \text{succ}(u)} \delta_s^{(l)} w_{su} \right) \frac{\partial \text{out}_u^{(l)}}{\partial net_u^{(l)}}.
\]
The resulting formula for the weight change is

\[
\Delta \vec{w}^{(l)}_u = -\frac{\eta}{2} \vec{w}_u e^{(l)} = \eta \delta_u^{(l)} \vec{m}_u^{(l)} = \eta \left( \sum_{s \in \text{succ}(u)} \delta_s^{(l)} w_{su} \right) \frac{\partial \text{out}^{(l)}_u}{\partial \text{net}^{(l)}_u} \vec{m}_u^{(l)}.
\]

Consider again the special case with

- output function is the identity,
- activation function is logistic.

The resulting formula for the weight change is then

\[
\Delta \vec{w}^{(l)}_u = \eta \left( \sum_{s \in \text{succ}(u)} \delta_s^{(l)} w_{su} \right) \text{out}^{(l)}_u (1 - \text{out}^{(l)}_u) \vec{m}_u^{(l)}.
\]
Error Backpropagation: Cookbook Recipe

∀u ∈ U_{in}:
out_u^{(l)} = ext_u^{(l)}

forward propagation:

∀u ∈ U_{hidden} ∪ U_{out}:
out_u^{(l)} = \left(1 + \exp\left(-\sum_{p \in \text{pred}(u)} w_{up} out_p^{(l)}\right)\right)^{-1}

backward propagation:

∀u ∈ U_{hidden}:
\delta_u^{(l)} = \left(\sum_{s \in \text{succ}(u)} \delta_s^{(l)} w_{su}\right) \lambda_u^{(l)}

activation derivative:

\lambda_u^{(l)} = out_u^{(l)} \left(1 - out_u^{(l)}\right)

error factor:

∀u ∈ U_{out}:
\delta_u^{(l)} = (o_u^{(l)} - out_u^{(l)}) \lambda_u^{(l)}

weight change:

\Delta w_{up}^{(l)} = \eta \delta_u^{(l)} out_p^{(l)}
Gradient descent training for the negation $\neg x$

\[ x \xrightarrow{w} \theta \xrightarrow{} y \]

<table>
<thead>
<tr>
<th>$x$</th>
<th>$y$</th>
</tr>
</thead>
<tbody>
<tr>
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<tr>
<td>1</td>
<td>0</td>
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</tbody>
</table>

error for $x = 0$   error for $x = 1$   sum of errors

Note: error for $x = 0$ and $x = 1$ is effectively the squared logistic activation function!
Gradient Descent: Examples

<table>
<thead>
<tr>
<th>epoch</th>
<th>$\theta$</th>
<th>$w$</th>
<th>error</th>
</tr>
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<tr>
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Online Training

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<th>error</th>
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<tr>
<td>220</td>
<td>-1.63</td>
<td>-3.53</td>
<td>0.044</td>
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</tbody>
</table>

Batch Training
Gradient Descent: Examples

Visualization of gradient descent for the negation $\neg x$

- Training is obviously successful.
- Error cannot vanish completely due to the properties of the logistic function.
Gradient Descent: Examples

Example function:

\[ f(x) = \frac{5}{6}x^4 - 7x^3 + \frac{115}{6}x^2 - 18x + 6, \]

<table>
<thead>
<tr>
<th>(i)</th>
<th>(x_i)</th>
<th>(f(x_i))</th>
<th>(f'(x_i))</th>
<th>(\Delta x_i)</th>
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Gradient descent with initial value 0.2 and learning rate 0.001.
Gradient Descent: Examples

Example function:

\[ f(x) = \frac{5}{6}x^4 - 7x^3 + \frac{115}{6}x^2 - 18x + 6, \]

\[
\begin{array}{|c|c|c|c|c|}
\hline
i & x_i & f(x_i) & f'(x_i) & \Delta x_i \\
\hline
0 & 1.500 & 2.719 & 3.500 & -0.875 \\
1 & 0.625 & 0.655 & -1.431 & 0.358 \\
2 & 0.983 & 0.955 & 2.554 & -0.639 \\
3 & 0.344 & 1.801 & -7.157 & 1.789 \\
4 & 2.134 & 4.127 & 0.567 & -0.142 \\
5 & 1.992 & 3.989 & 1.380 & -0.345 \\
6 & 1.647 & 3.203 & 3.063 & -0.766 \\
7 & 0.881 & 0.734 & 1.753 & -0.438 \\
8 & 0.443 & 1.211 & -4.851 & 1.213 \\
9 & 1.656 & 3.231 & 3.029 & -0.757 \\
10 & 0.898 & 0.766 & & \\
\hline
\end{array}
\]

Gradient descent with initial value 1.5 and learning rate 0.25.
Gradient Descent: Examples

Example function:

\[ f(x) = \frac{5}{6}x^4 - 7x^3 + \frac{115}{6}x^2 - 18x + 6, \]

<table>
<thead>
<tr>
<th>( i )</th>
<th>( x_i )</th>
<th>( f(x_i) )</th>
<th>( f'(x_i) )</th>
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<tr>
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<td>3.327</td>
<td>2.585</td>
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<td></td>
</tr>
</tbody>
</table>

Gradient descent with initial value 2.6 and learning rate 0.05.
Gradient Descent: Variants

Weight update rule:

\[ w(t + 1) = w(t) + \Delta w(t) \]

**Standard backpropagation:**

\[ \Delta w(t) = -\frac{\eta}{2} \nabla w e(t) \]

**Manhattan training:**

\[ \Delta w(t) = -\eta \text{sgn}(\nabla w e(t)). \]

Fixed step width (grid), only sign of gradient (direction) is evaluated.

**Momentum term:**

\[ \Delta w(t) = -\frac{\eta}{2} \nabla w e(t) + \beta \Delta w(t - 1), \]

Part of previous change is added, may lead to accelerated training \((\beta \in [0.5, 0.95])\).
Self-adaptive error backpropagation:

\[ \eta_w(t) = \begin{cases} 
  c^- \cdot \eta_w(t-1), & \text{if } \nabla_w e(t) \cdot \nabla_w e(t-1) < 0, \\
  c^+ \cdot \eta_w(t-1), & \text{if } \nabla_w e(t) \cdot \nabla_w e(t-1) > 0 \\
  \eta_w(t-1), & \text{otherwise.} 
\end{cases} \]

Resilient error backpropagation:

\[ \Delta w(t) = \begin{cases} 
  c^- \cdot \Delta w(t-1), & \text{if } \nabla_w e(t) \cdot \nabla_w e(t-1) < 0, \\
  c^+ \cdot \Delta w(t-1), & \text{if } \nabla_w e(t) \cdot \nabla_w e(t-1) > 0 \\
  \Delta w(t-1), & \text{otherwise.} 
\end{cases} \]

Typical values: \( c^- \in [0.5, 0.7] \) and \( c^+ \in [1.05, 1.2] \).
Quickpropagation

The weight update rule can be derived from the triangles:

$$
\Delta w(t) = \frac{\nabla_{we}(t)}{\nabla_{we}(t-1) - \nabla_{we}(t)} \cdot \Delta w(t-1).
$$
Gradient Descent: Examples

<table>
<thead>
<tr>
<th>epoch</th>
<th>$\theta$</th>
<th>$w$</th>
<th>error</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
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<tr>
<td>220</td>
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<td>-3.53</td>
<td>0.044</td>
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</table>

without momentum term

<table>
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<tr>
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<th>$\theta$</th>
<th>$w$</th>
<th>error</th>
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with momentum term ($\beta = 0.9$)
Gradient Descent: Examples

- Dots show position every 20 (without momentum term) or every 10 epochs (with momentum term).
- Learning with a momentum term ($\beta = 0.9$) is about twice as fast.
Gradient Descent: Examples

Example function:

\[ f(x) = \frac{5}{6} x^4 - 7x^3 + \frac{115}{6} x^2 - 18x + 6, \]

<table>
<thead>
<tr>
<th>$i$</th>
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<th>$f'(x_i)$</th>
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Gradient descent with initial value 0.2, learning rate 0.001 and momentum term $\beta = 0.9$. 
Gradient Descent: Examples

Example function: \[ f(x) = \frac{5}{6}x^4 - 7x^3 + \frac{115}{6}x^2 - 18x + 6, \]

<table>
<thead>
<tr>
<th>( i )</th>
<th>( x_i )</th>
<th>( f(x_i) )</th>
<th>( f'(x_i) )</th>
<th>( \Delta x_i )</th>
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</table>

Gradient descent with initial value 1.5, initial learning rate 0.25, and self-adapting learning rate \((c^+ = 1.2, c^- = 0.5)\).
Other Extensions of Error Backpropagation

Flat Spot Elimination:

\[ \Delta w(t) = -\frac{\eta}{2} \nabla_w e(t) + \zeta \]

- Eliminates slow learning in saturation region of logistic function (\(\zeta \approx 0.1\)).
- Counteracts the decay of the error signals over the layers.

Weight Decay:

\[ \Delta w(t) = -\frac{\eta}{2} \nabla_w e(t) - \xi w(t), \]

- Helps to improve the robustness of the training results (\(\xi \leq 10^{-3}\)).
- Can be derived from an extended error function penalizing large weights:

\[ e^* = e + \frac{\xi}{2} \sum_{u \in U_{out} \cup U_{hidden}} \left( \theta_u^2 + \sum_{p \in pred(u)} w_{up}^2 \right). \]
Number of Hidden Neurons

- Note that the approximation theorem only states that there exists a number of hidden neurons and weight vectors $\vec{v}$ and $\vec{w}_i$ and thresholds $\theta_i$, but not how they are to be chosen for a given $\varepsilon$ of approximation accuracy.

- For a single hidden layer the following rule of thumb is popular:
  number of hidden neurons = (number of inputs + number of outputs) / 2

- Better, though computationally expensive approach:
  - Randomly split the given data into two subsets of (about) equal size, the training data and the validation data.
  - Train multi-layer perceptrons with different numbers of hidden neurons on the training data and evaluate them on the validation data.
  - Repeat the random split of the data and training/evaluation many times and average the results over the same number of hidden neurons. Choose the number of hidden neurons with the best average error.
  - Train a final multi-layer perceptron on the whole data set.
Number of Hidden Neurons

Principle of training data/validation data approach:

- **Underfitting:** If the number of neurons in the hidden layer is too small, the multi-layer perceptron may not be able to capture the structure of the relationship between inputs and outputs precisely enough due to a lack of parameters.

- **Overfitting:** With a larger number of hidden neurons a multi-layer perceptron may adapt not only to the regular dependence between inputs and outputs, but also to the accidental specifics (errors and deviations) of the training data set.

- Overfitting will usually lead to the effect that the error a multi-layer perceptron yields on the validation data will be (possibly considerably) greater than the error it yields on the training data.

  The reason is that the validation data set is likely distorted in a different fashion than the training data, since the errors and deviations are random.

- Minimizing the error on the validation data by properly choosing the number of hidden neurons prevents both under- and overfitting.
Objective: select the model that best fits the data, taking the model complexity into account.

The more complex the model, the better it usually fits the data.

- The blue curve fits the data points perfectly, but it is not a good model.
The described method of iteratively splitting the data into training and validation data may be referred to as **cross validation**.

However, this term is more often used for the following specific procedure:

- The given data set is split into *n* parts or subsets (also called folds) of about equal size (so-called **n-fold cross validation**).
- If the output is nominal (also sometimes called symbolic or categorical), this split is done in such a way that the relative frequency of the output values in the subsets/folds represent as well as possible the relative frequencies of these values in the data set as a whole. This is also called **stratification** (derived from the Latin *stratum*: layer, level, tier).
- Out of these *n* data subsets (or folds) *n* pairs of training and validation data set are formed by using one fold as a validation data set while the remaining *n* − 1 folds are combined into a training data set.
The advantage of the cross validation method is that one random split of the data yields \( n \) different pairs of training and validation data set.

An obvious disadvantage is that (except for \( n = 2 \)) the size of the training and the test data set are considerably different, which makes the results on the validation data statistically less reliable.

It is therefore only recommended for sufficiently large data sets or sufficiently small \( n \), so that the validation data sets are of sufficient size.

Repeating the split (either with \( n = 2 \) or greater \( n \)) has the advantage that one obtains many more training and validation data sets, leading to more reliable statistics (here: for the number of hidden neurons).

The described approaches fall into the category of resampling methods.

Other well-known statistical resampling methods are bootstrap, jackknife, subsampling and permutation test.
Avoiding Overfitting: Alternatives

- An alternative way to prevent overfitting is the following approach:
  - During training the performance of the multi-layer perceptron is evaluated after each epoch (or every few epochs) on a **validation data set**.
  - While the error on the training data set should always decrease with each epoch, the error on the validation data set should, after decreasing initially as well, increase again as soon as overfitting sets in.
  - At this moment training is terminated and either the current state or (if available) the state of the multi-layer perceptron, for which the error on the validation data reached a minimum, is reported as the training result.

- Furthermore a stopping criterion may be derived from the shape of the error curve on the training data over the training epochs, or the network is trained only for a fixed, relatively small number of epochs (also known as **early stopping**).

- Disadvantage: these methods stop the training of a complex network early enough, rather than adjust the complexity of the network to the “correct” level.
Sensitivity Analysis
Sensitivity Analysis

Problem of Multi-Layer Perceptrons:

- The knowledge that is learned by a neural network is encoded in matrices/vectors of real-valued numbers and is therefore often difficult to understand or to extract.

- Geometric interpretation or other forms of intuitive understanding are possible only for very simple networks, but fail for complex practical problems.

- Thus a neural network is often effectively a *black box*, that computes its output, though in a mathematically precise way, in a way that is very difficult to understand and to interpret.

Idea of Sensitivity Analysis:

- Try to find out to which inputs the output(s) react(s) most sensitively.

- This may also give hints which inputs are not needed and may be discarded.
Question: How important are different inputs to the network?

Idea: Determine change of output relative to change of input.

\[
\forall u \in U_{in} : \quad s(u) = \frac{1}{|L_{fixed}|} \sum_{l \in L_{fixed}} \sum_{v \in U_{out}} \frac{\partial out_v^{(l)}}{\partial ext_u^{(l)}}.
\]

Formal derivation: Apply chain rule.

\[
\frac{\partial out_v}{\partial ext_u} = \frac{\partial out_v}{\partial out_u} \frac{\partial out_u}{\partial ext_u} = \frac{\partial out_v}{\partial net_v} \frac{\partial net_v}{\partial out_u} \frac{\partial out_u}{\partial ext_u}.
\]

Simplification: Assume that the output function is the identity.

\[
\frac{\partial out_u}{\partial ext_u} = 1.
\]
Sensitivity Analysis

For the second factor we get the general result:

\[
\frac{\partial \text{net}_v}{\partial \text{out}_u} = \frac{\partial}{\partial \text{out}_u} \sum_{p \in \text{pred}(v)} w_{vp} \text{out}_p = \sum_{p \in \text{pred}(v)} w_{vp} \frac{\partial \text{out}_p}{\partial \text{out}_u}.
\]

This leads to the recursion formula

\[
\frac{\partial \text{out}_v}{\partial \text{out}_u} = \frac{\partial \text{out}_v}{\partial \text{out}_u} \frac{\partial \text{net}_v}{\partial \text{out}_u} = \frac{\partial \text{out}_v}{\partial \text{net}_v} \sum_{p \in \text{pred}(v)} w_{vp} \frac{\partial \text{out}_p}{\partial \text{out}_u}.
\]

However, for the first hidden layer we get

\[
\frac{\partial \text{net}_v}{\partial \text{out}_u} = w_{vu}, \quad \text{therefore} \quad \frac{\partial \text{out}_v}{\partial \text{out}_u} = \frac{\partial \text{out}_v}{\partial \text{net}_v} w_{vu}.
\]

This formula marks the start of the recursion.
Consider as usual the special case with

- output function is the identity,
- activation function is logistic.

The recursion formula is in this case

\[
\frac{\partial \text{out}_v}{\partial \text{out}_u} = \text{out}_v(1 - \text{out}_v) \sum_{p \in \text{pred}(v)} w_{vp} \frac{\partial \text{out}_p}{\partial \text{out}_u}
\]

and the anchor of the recursion is

\[
\frac{\partial \text{out}_v}{\partial \text{out}_u} = \text{out}_v(1 - \text{out}_v)w_{vu}.
\]
Attention: Use weight decay to stabilize the training results!

\[ \Delta w(t) = -\frac{\eta}{2} \nabla w e(t) - \xi w(t), \]

- Without weight decay the results of a sensitivity analysis can depend (strongly) on the (random) initialization of the network.

Example: Iris data, 1 hidden layer, 3 hidden neurons, 1000 epochs

<table>
<thead>
<tr>
<th>attribute</th>
<th>$\xi = 0$</th>
<th>$\xi = 0.0001$</th>
</tr>
</thead>
<tbody>
<tr>
<td>sepal length</td>
<td>0.0216 0.0399 0.0232 0.0515</td>
<td>0.0367 0.0325 0.0351 0.0395</td>
</tr>
<tr>
<td>sepal width</td>
<td>0.0423 0.0341 0.0460 0.0447</td>
<td>0.0385 0.0376 0.0421 0.0425</td>
</tr>
<tr>
<td>petal length</td>
<td>0.1789 0.2569 0.1974 0.2805</td>
<td>0.2048 0.1928 0.1838 0.1861</td>
</tr>
<tr>
<td>petal width</td>
<td>0.2017 0.1356 0.2198 0.1325</td>
<td>0.2020 0.1962 0.1750 0.1743</td>
</tr>
</tbody>
</table>
Reminder: The Iris Data

- Collected by Edgar Anderson on the Gaspé Peninsula (Canada).
- First analyzed by Ronald Aylmer Fisher (famous statistician).
- 150 cases in total, 50 cases per Iris flower type.
- Measurements of sepal length and width and petal length and width (in cm).
- Most famous data set in pattern recognition and data analysis.
Application: Recognition of Handwritten Digits

- 9298 segmented and digitized digits of handwritten ZIP codes.
- Appeared on US Mail envelopes passing through the post office in Buffalo, NY.
- Written by different people, great variety of sizes, writing styles and instruments, with widely varying levels of care.
- In addition: 3349 printed digits from 35 different fonts.
Application: Recognition of Handwritten Digits

- Acquisition, binarization, location of the ZIP code on the envelope and segmentation into individual digits were performed by Postal Service contractors.
- Further segmentation were performed by the authors of [LeCun et al. 1990].
- Segmentation of totally unconstrained digit strings is a difficult problem.
- Several ambiguous characters are the result of missegmentation (especially broken 5s, see picture above).
• Segmented digits vary in size, but are typically around 40 to 60 pixels.
• The size of the digits is normalized to $16 \times 16$ pixels with a simple linear transformation (preserve aspect ratio).
• As a result of the transformation, the resulting image is not binary, but has multiple gray levels, which are scaled to fall into $[-1, +1]$.
• Remainder of the recognition is performed entirely by an MLP.
Objective: train an MLP to recognize the digits.

- 7291 handwritten and 2549 printed digits in training set; 2007 handwritten and 700 printed digits (different fonts) in test set.
- Both training and test set contain multiple examples that are ambiguous, unclassifiable, or even misclassified.
- Picture above: normalized digits from the test set.
• **Challenge:** all connections are adaptive, although heavily constrained (in contrast to earlier work which used manually chosen first layers).

• **Training:** error backpropagation / gradient descent.

• **Input:** 16 × 16 pixel images of the normalized digits (256 input neurons).

• **Output:** 10 neurons, that is, one neuron per class.

  If an input image belongs to class $i$ (shows digit $i$), the output neuron $u_i$ should produce a value of $+1$, all other output neurons should produce a value of $-1$.

• **Problem:** for a fully connected network with multiple hidden layers the number of parameters is excessive (risk of overfitting).

• **Solution:** instead of a fully connected network, use a locally connected one.
Application: Recognition of Handwritten Digits

- Used network has four hidden layers plus input and output layer.
- Local connections in all but the last layer:
  - Two-dimensional layout of the input and hidden layers (neuron grids).
  - Connections to a neuron in the next layer from a group of neighboring neurons in the preceding layer (small sub-grid, local receptive field).
Each neuron of the (first) hidden layer is connected to a small number of input neurons that refer to a contiguous region of the input image (left).

Connection weights are shared, same network is evaluated at different locations. The input field is “moved” step by step over the whole image (right).

Equivalent to a convolution with a small size kernel ("receptive field").
• Idea: local connections can implement feature detectors.

• In addition: analogous connections are forced to have identical weights to allow for features appearing in different parts of the input (weight sharing).

• Equivalent to a convolution with a small size kernel, followed by an application of the activation function.
Application: Recognition of Handwritten Digits

- Note: Weight sharing considerably reduces the number of free parameters.
- The outputs of a set of neurons with shared weights constitute a feature map.
- In practice, multiple feature maps, extracting different features, are needed.
- The idea of local, convolutional feature maps can be applied to subsequent hidden layers as well to obtain more complex and abstract features.
Higher level features require less precise coding of their location; therefore each additional layer performs local averaging and subsampling.

The loss of spatial resolution is partially compensated by an increase in the number of different features.

Layers H1 and H3 are shared-weight feature extractors, Layers H2 and H4 are averaging/subsampling layers.
Application: Recognition of Handwritten Digits

- Input image is enlarged from $16 \times 16$ to $28 \times 28$ to avoid boundary problems.
- In the first hidden layer, four feature maps are represented. All neurons in the same feature map share the same weight vectors.
- The second hidden layer averages over $2 \times 2$ blocks in the first hidden layer.
- 12 feature maps in H3, averaging over $2 \times 2$ blocks in H4.
- Feature maps in hidden layers H2 and H3 are connected as follows:

<table>
<thead>
<tr>
<th>H2/H3</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
<th>6</th>
<th>7</th>
<th>8</th>
<th>9</th>
<th>10</th>
<th>11</th>
<th>12</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>×</td>
<td>×</td>
<td>×</td>
<td>×</td>
<td>×</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>2</td>
<td></td>
<td>×</td>
<td>×</td>
<td>×</td>
<td>×</td>
<td>×</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>3</td>
<td></td>
<td></td>
<td>×</td>
<td>×</td>
<td>×</td>
<td>×</td>
<td>×</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>4</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

- The output layer is fully connected to the fourth hidden layer H4.
- In total: 4635 neurons, 98442 connections, 2578 independent parameters.
Application: Recognition of Handwritten Digits

- After 30 training epochs, the error rate on the training set was 1.1% (MSE: 0.017), and 3.4% on the test set (MSE: 0.024). Compared to human error: 2.5%

- All classification errors occurred on handwritten digits.

- Substitutions (misclassified) versus Rejections (unclassified):
  - Reject an input if the difference in output between the two neurons with the highest output is less than a threshold.
  - Best result on the test set: 5.7% rejections for 1% substitutions.
  - On handwritten data alone: 9% rejections for 1% substitutions or 6% rejections for 2% substitutions.

- Atypical input (see below) is also correctly classified.

picture not available in online version
Demonstration of multi-layer perceptron training:

- Visualization of the training process
- Biimplication and Exclusive Or, two continuous functions
- http://www.borgelt.net/mlpd.html
Multi-Layer Perceptron Software: mlp/mlpgui

Software for training general multi-layer perceptrons:

- Command line version written in C, fast training
- Graphical user interface in Java, easy to use

http://www.borgelt.net/mlp.html
http://www.borgelt.net/mlpgui.html
Deep Learning
(Multi-Layer Perceptrons with Many Layers)
Deep Learning: Motivation

- **Reminder:** *Universal Approximation Theorem*
  Any continuous function on an arbitrary compact subspace of $\mathbb{R}^n$ can be approximated arbitrarily well with a three-layer perceptron.

- This theorem is often cited as (allegedly!) meaning that
  - we can confine ourselves to multi-layer perceptrons with only one hidden layer,
  - there is no real need to look at multi-layer perceptrons with more hidden layers.

- **However:** The theorem says nothing about the number of hidden neurons that may be needed to achieve a desired approximation accuracy.

- Depending on the function to approximate, a very large number of neurons may be necessary.

- Allowing for more hidden layers may enable us to achieve the same approximation quality with a significantly lower number of neurons.
• Very simple and commonly used example is the **n-bit parity function**: Output is 1 if an even number of inputs is 1; output is 0 otherwise.

• Can easily be represented by a multi-layer perceptron with only one hidden layer. (See reminder of TLU algorithm on next slide; adapt to MLPs.)

• However, the solution has $2^{n-1}$ **hidden neurons**: disjunctive normal form is a disjunction of $2^{n-1}$ conjunctions, which represent the $2^{n-1}$ input combinations with an even number of set bits.

• Number of hidden neurons **grows exponentially** with the number of inputs.

• However, if more hidden layers are admissible, **linear growth** is possible:
  - Start with a *biimplication* of two inputs.
  - Continue with a chain of *exclusive ors*, each of which adds another input.
  - Such a network needs $n + 3(n - 1) = 4n - 3$ neurons in total
    ($n$ input neurons, $3(n - 1) - 1$ **hidden neurons**, 1 output neuron)
Algorithm: Let $y = f(x_1, \ldots, x_n)$ be a Boolean function of $n$ variables.

(i) Represent the given function $f(x_1, \ldots, x_n)$ in disjunctive normal form. That is, determine where all $K_j$ are conjunctions of $n$ literals, that is, $K_j = l_{j1} \land \ldots \land l_{jn}$ with $l_{ji} = x_i$ (positive literal) or $l_{ji} = \neg x_i$ (negative literal).

(ii) Create a neuron for each conjunction $K_j$ of the disjunctive normal form (having $n$ inputs — one input for each variable), where

$$w_{ji} = \begin{cases} 2, & \text{if } l_{ji} = x_i, \\ -2, & \text{if } l_{ji} = \neg x_i, \end{cases}$$

and

$$\theta_j = n - 1 + \frac{1}{2} \sum_{i=1}^{n} w_{ji}.$$

(iii) Create an output neuron (having $m$ inputs — one input for each neuron that was created in step (ii)), where

$$w_{(n+1)k} = 2, \quad k = 1, \ldots, m,$$

and

$$\theta_{n+1} = 1.$$

Remark: weights are set to $\pm 2$ instead of $\pm 1$ in order to ensure integer thresholds.
Deep Learning: $n$-bit Parity Function

- Implementation of the $n$-bit parity function with a chain of one biimplication and $n - 2$ exclusive or sub-networks.

- Note that the structure is not strictly layered, but could be completed. If this is done, the number of neurons increases to $n(n + 1) - 1$. 
Deep Learning: Motivation

- Similar to the situation w.r.t. linearly separable functions:
  - Only few $n$-ary Boolean functions are linearly separable (see next slide).
  - Only few $n$-ary Boolean functions need few hidden neurons in a single layer.

- An $n$-ary Boolean function has $k_0$ input combinations with an output of 0 and $k_1$ input combinations with an output of 1 (clearly $k_0 + k_1 = 2^n$).

- We need at most $2^{\min\{k_0, k_1\}}$ hidden neurons if we choose disjunctive normal form for $k_1 \leq k_0$ and conjunctive normal form for $k_1 > k_0$.

- As there are $\binom{2^n}{k_0}$ possible functions with $k_0$ input combinations mapped to 0 and $k_1$ mapped to 1, many functions require a substantial number of hidden neurons.

- Although the number of neurons may be reduced with minimization methods like the Quine–McCluskey algorithm [Quine 1952, 1955; McCluskey 1956], the fundamental problem remains.
Reminder: Limitations of Threshold Logic Units

Total number and number of linearly separable Boolean functions
(On-Line Encyclopedia of Integer Sequences, oeis.org, A001146 and A000609):

<table>
<thead>
<tr>
<th>inputs</th>
<th>Boolean functions</th>
<th>linearly separable functions</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>4</td>
<td>4</td>
</tr>
<tr>
<td>2</td>
<td>16</td>
<td>14</td>
</tr>
<tr>
<td>3</td>
<td>256</td>
<td>104</td>
</tr>
<tr>
<td>4</td>
<td>65,536</td>
<td>1,882</td>
</tr>
<tr>
<td>5</td>
<td>4,294,967,296</td>
<td>94,572</td>
</tr>
<tr>
<td>6</td>
<td>18,446,744,073,709,551,616</td>
<td>15,028,134</td>
</tr>
<tr>
<td>$n$</td>
<td>$2^{(2^n)}$</td>
<td>no general formula known</td>
</tr>
</tbody>
</table>

- For many inputs a threshold logic unit can compute almost no functions.
- Networks of threshold logic units are needed to overcome the limitations.
Deep Learning: Motivation

- In practice the problem is mitigated considerably by the simple fact that training data sets are necessarily limited in size.

- Complete training data for an $n$-ary Boolean function has $2^n$ training examples. Data sets for practical problems usually contain much fewer sample cases. This leads to many input configurations for which no desired output is given; freedom to assign outputs to them allows for a simpler representation.

- Nevertheless, using more than one hidden layer promises in many cases to reduce the number of needed neurons.

- This is the focus of the area of deep learning. (Depth means the length of the longest path in the network graph.)

- For multilayer perceptrons (longest path: number of hidden layers plus one), deep learning starts with more than one hidden layer.

- For 10 or more hidden layers, one sometimes speaks of very deep learning.
Deep Learning: Main Problems

• Deep learning multilayer perceptrons suffer from two main problems: overfitting and vanishing gradient.

• Overfitting results mainly from the increased number of adaptable parameters.

• **Weight decay** prevents large weights and thus an overly precise adaptation.

• **Sparsity constraints** help to avoid overfitting:
  ○ there is only a restricted number of neurons in the hidden layers or
  ○ only few of the neurons in the hidden layers should be active (on average).
  May be achieved by adding a regularization term to the error function (compares the observed number of active neurons with desired number and pushes adaptations into a direction that tries to match these numbers).

• Furthermore, a training method called **dropout training** may be applied: some units are randomly omitted from the input/hidden layers during training.
Deep Learning: Dropout

- Desired characteristic: Robustness against neuron failures.

- Approach during training:
  - Use only $p\%$ of the neurons (e.g. $p = 50$: half of the neurons).
  - Choose dropout neurons randomly.

- Approach during execution:
  - Use all of the neurons.
  - Multiply all weights by $p\%$.

- Result of this approach:
  - More robust representation.
  - Better generalization.
Reminder: Cookbook Recipe for Error Backpropagation

Forward propagation:
\[ \forall u \in U_{in}: \quad \text{out}_{u}^{(l)} = \text{ext}_{u}^{(l)} \]

\[ \forall u \in U_{in}: \quad \text{out}_{u}^{(l)} = \text{ext}_{u}^{(l)} \]

Backward propagation:
\[ \forall u \in U_{hidden} \cup U_{out}: \quad \text{out}_{u}^{(l)} = \left(1 + \exp \left(-\sum_{p \in \text{pred}(u)} w_{up} \text{out}_{p}^{(l)}\right)\right)^{-1} \]

Errors:
- logistic activation function
- implicit bias value
- error factor:

Activation derivative:
\[ \lambda_{u}^{(l)} = \text{out}_{u}^{(l)} (1 - \text{out}_{u}^{(l)}) \]

Weight change:
\[ \Delta w_{up}^{(l)} = \eta \delta_{u}^{(l)} \text{out}_{p}^{(l)} \]
logistic activation function:
\[ f_{\text{act}}(\text{net}_u^{(l)}) = \frac{1}{1 + e^{-\text{net}_u^{(l)}}} \]

derivative of logistic function:
\[ f'_{\text{act}}(\text{net}_u^{(l)}) = f_{\text{act}}(\text{net}_u^{(l)}) \cdot (1 - f_{\text{act}}(\text{net}_u^{(l)})) \]

- If a logistic activation function is used (shown on left), the weight changes are proportional to \( \lambda_u^{(l)} = \text{out}_u^{(l)} \left( 1 - \text{out}_u^{(l)} \right) \) (shown on right; see recipe).

- This factor is also propagated back, but cannot be larger than \( \frac{1}{4} \) (see right).

⇒ The gradient tends to vanish if many layers are backpropagated through. Learning in the early hidden layers can become very slow [Hochreiter 1991].
• In principle, a small gradient may be counteracted by a large weight.

\[
\delta_u^{(l)} = \left( \sum_{s \in \text{succ}(u)} \delta_s^{(l)} w_{su} \right) \lambda_u^{(l)}.
\]

• However, usually a large weight, since it also enters the activation function, drives the activation function to its saturation regions.

• Thus, (the absolute value of) the derivative factor is usually the smaller, the larger (the absolute value of) the weights.

• Furthermore, the connection weights are commonly initialized to a random value in the range from $-1$ to $+1$.

⇒ **Initial training steps are particularly affected:** both the gradient as well as the weights provide a factor less than 1.

• Theoretically, there can be exceptions to this description. However, they are rare in practice and one usually observes a vanishing gradient.
Alternative way to understand the vanishing gradient effect:

- The logistic activation function is a **contracting function**: for any two arguments $x$ and $y$, $x \neq y$, we have $|f_{\text{act}}(x) - f_{\text{act}}(y)| < |x - y|$. (Obvious from the fact that its derivative is always $< 1$; actually $\leq \frac{1}{4}$.)

- If several logistic functions are chained, these contractions combine and yield an even stronger contraction of the input range.

- As a consequence, a rather large change of the input values will produce only a rather small change in the output values, and the more so, the more logistic functions are chained together.

- Therefore the function that maps the inputs of a multilayer perceptron to its outputs usually becomes the flatter the more layers the multilayer perceptron has.

- Consequently the gradient in the first hidden layer (were the inputs are processed) becomes the smaller.
Deep Learning: Different Activation Functions

rectified maximum/ramp function:
\[ f_{\text{act}}(\text{net}, \theta) = \max\{0, \text{net} - \theta\} \]

softplus function:
\[ f_{\text{act}}(\text{net}, \theta) = \ln(1 + e^{\text{net} - \theta}) \]

- The vanishing gradient problem may be battled with other activation functions.
- These activation functions yield so-called **rectified linear units (ReLUs)**.
- **Rectified maximum/ramp function**
  
  Advantages: simple computation, simple derivative, zeros simplify learning
  
  Disadvantages: no learning \( \leq \theta \), rather inelegant, not continuously differentiable
Reminder: Training a TLU for the Negation

Single input threshold logic unit for the negation $\neg x$.

Modified output error as a function of weight and threshold:

error for $x = 0$  
error for $x = 1$  
sum of errors

A rectified maximum/ramp function yields these errors directly.
Deep Learning: Different Activation Functions

rectified maximum/ramp function:
\[ f_{act}(\text{net}, \theta) = \max\{0, \text{net} - \theta\} \]

softplus function:
\[ f_{act}(\text{net}, \theta) = \ln(1 + e^{\text{net} - \theta}) \]

- **Softplus function**: continuously differentiable, but much more complex

- Alternatives:
  - **Leaky ReLU**: \( f(x) = \begin{cases} x & \text{if } x > 0, \\ \nu x & \text{otherwise.} \end{cases} \) (adds learning \( \leq \theta; \nu \approx 0.01 \))
  - **Noisy ReLU**: \( f(x) = \max\{0, x + \mathcal{N}(0, \sigma(x))\} \) (adds Gaussian noise)
The derivative of the rectified maximum/ramp function is the **step function**.

The derivative of the softplus function is the **logistic function**.

Factor resulting from derivative of activation function can be much larger.

⇒ Larger gradient in early layers of the network; training is (much) faster.

What also helps: **faster hardware, implementations on GPUs** etc.
Reminder: Networks of threshold logic units cannot be trained, because

- there are no desired values for the neurons of the first layer(s),
- the problem can usually be solved with several different functions computed by the neurons of the first layer(s) (non-unique solution).

Although multi-layer perceptrons, even with many hidden layers, can be trained, the same problems still affect the effectiveness and efficiency of training.

Alternative: **Build network layer by layer,**

\textit{train only newly added layer in each step.}

Popular: Build the network as **stacked auto-encoders.**

An **auto-encoder** is a 3-layer perceptron that maps its inputs to approximations of these inputs.

- The hidden layer forms an encoder into some form of internal representation.
- The output layer forms a decoder that (approximately) reconstructs the inputs.
An auto-encoder/decoder (left), of which only the encoder part (right) is later used. The $x_i$ are the given inputs, the $\hat{x}_i$ are the reconstructed inputs, and the $f_i$ are the constructed features; the error is $e = \sum_{i=1}^{n}(\hat{x}_i - x_i)^2$. Training is conducted with error backpropagation.
• Rationale of training an auto-encoder: hidden layer is expected to construct features.

• Hope: Features capture the information contained in the input in a compressed form (encoder), so that the input can be well reconstructed from it (decoder).

• Note: this implicitly assumes that features that are well suited to represent the inputs in a compressed way are also useful to predict some desired output. Experience shows that this assumption is often justified.

• Main problems:
  ◦ how many units should be chosen for the hidden layer?
  ◦ how should this layer be treated during training?

• If there are as many (or even more) hidden units as there are inputs, it is likely that it will merely pass through its inputs to the output layer.

• The most straightforward solution are sparse auto-encoders: There should be (considerably) fewer hidden neurons than there are inputs.
• Few hidden neurons force the auto-encoder to learn relevant features (since it is not possible to simply pass through the inputs).

• How many hidden neurons are a good choice? Note that cross-validation not necessarily a good approach!

• Alternative to few hidden neurons: **sparse activation scheme**
The number of active neurons in the hidden layer is restricted to a small number. May be enforced by either adding a regularization term to the error function that punishes a larger number of active hidden neurons or by explicitly deactivating all but a few neurons with the highest activations.

• A third approach is to add noise (that is, random variations) to the input (but not to the copy of the input used to evaluate the reconstruction error): **denoising auto-encoders**.
  (The auto-encoder to be trained is expected to map the input with noise to (a copy of) the input without noise, thus preventing simple passing through.)
Deep Learning: Stacked Auto-Encoders

- A popular way to initialize/pre-train a stacked auto-encoder is a **greedy layer-wise training approach**.

- In the first step, a (sparse) auto-encoder is trained for the raw input features. The hidden layer constructs **primary features** useful for reconstruction.

- A primary feature data set is obtained by propagating the raw input features up to the hidden layer and recording the hidden neuron activations.

- In the second step, a (sparse) auto-encoder is trained for the obtained feature data set. The hidden layer constructs **secondary features** useful for reconstruction.

- A secondary feature data set is obtained by propagating the primary features up to the hidden layer and recording the hidden neuron activations.

- This process is repeated as many times as hidden layers are desired.

- Finally the encoder parts of the trained **auto-encoders are stacked** and the resulting network is **fine-tuned with error backpropagation**.
1. Train auto-encoder for the raw input; this yields a primary feature set.
2. Train auto-encoder for the primary features; this yields a secondary feature set.
3. A classifier/predictor for the output is trained from the secondary feature set.
4. The resulting networks are stacked.
Deep Learning: Convolutional Neural Networks (CNNs)

- Multilayer perceptrons with several hidden layers built in the way described have been applied very successfully for handwritten digit recognition.

- In such an application it is assumed, though, that the handwriting has already been preprocessed in order to separate the digits (or, in other cases, the letters) from each other.

- However, one would like to use similar networks also for more general applications, for example, recognizing whole lines of handwriting or analyzing photos to identify their parts as sky, landscape, house, pavement, tree, human being etc.

- For such applications it is advantageous that the features constructed in hidden layers are not localized to a specific part of the image.

- Special form of deep learning multi-layer perceptron: **convolutional neural network**.

- Inspired by the human retina, where sensory neurons have a **receptive field**, that is, a limited region in which they respond to a (visual) stimulus.
- Each neuron of the (first) hidden layer is connected to a small number of input neurons that refer to a contiguous region of the input image (left).

- Connection weights are shared, same network is evaluated at different locations. The input field is “moved” step by step over the whole image (right).

- Equivalent to a convolution with a small size kernel.
Reminder: Recognition of Handwritten Digits

- Idea: local connections can implement feature detectors.
- In addition: analogous connections are forced to have identical weights to allow for features appearing in different parts of the input (weight sharing).
- Equivalent to a convolution with a small size kernel, followed by an application of the activation function.

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• Network with four hidden layers implementing two convolution and two averaging stages.
• Developed in 1990, so training took several days, despite its simplicity.
• Nowadays networks working on much larger images and having many more layers are possible.
• **ImageNet**
  is an image database organized according to WordNet nouns.

• **WordNet**
  is a lexical database [of English]; nouns, verbs, adjectives and adverbs are grouped into sets of cognitive synonyms (synsets) each expressing a distinct concept; contains $>100,000$ synsets, of which $>80,000$ are nouns.

• The ImageNet database contains hundreds and thousands of images for each noun/synset.
ImageNet Large Scale Visual Recognition Challenge (ILSVRC)

- Uses a subset of the ImageNet database (1.2 million images, 1,000 categories)
- Evaluates algorithms for **object detection** and **image classification**.
- Yearly challenges/competitions with corresponding workshops since 2010.

**Classification error (top5)** of ILSVRC winners

- Hardly possible 10 years ago; rapid improvement
- Often used: network ensembles
- Very deep learning: ResNet (2015) had > 150 layers
• Structure of the AlexNet deep neural network (winner 2012) [Krizhevsky 2012].
• 60 million parameters, 650,000 neurons, efficient GPU implementation.

picture not available in online version
German Traffic Sign Recognition Benchmark (GTSRB)

pictures not available in online version

- Competition at Int. Joint Conference on Neural Networks (IJCNN) 2011 [Stallkamp et al. 2012] benchmark.ini.rub.de/?section=gtsrb

- Single image, multi-class classification problem (that is, each image can belong to multiple classes).

- More than 40 classes, more than 50,000 images.

- Physical traffic sign instances are unique within the data set (that is, each real-world traffic sign occurs only once).

- Winning entry (committee of CNNs) outperforms humans! error rate winner (convolutional neural networks): 0.54%
  runner up (not a neural network): 1.12%
  human recognition: 1.16%
One way to visualize what goes on in a neural network is to “turn the network upside down” and ask it to enhance an input image in such a way as to elicit a particular interpretation.

Impose a prior constraint that the image should have similar statistics to natural images, such as neighboring pixels needing to be correlated.

Result: NNs trained to discriminate between different kinds of images contain information needed to generate images.
Deep Learning: Board Game Go

- board with $19 \times 19$ grid lines, stones are placed on line crossings
- objective: surround territory/crossings (and enemy stones: capture)
- special “ko” rules for certain situations to prevent infinite retaliation
- a player can pass his/her turn (usually disadvantageous)
- game ends after both players subsequently pass a turn
- winner is determined by counting (controlled area plus captured stones)

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<th>$b$</th>
<th>$d$</th>
<th>$b^d$</th>
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<tbody>
<tr>
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<td>$\approx 35$</td>
<td>$\approx 80$</td>
<td>$\approx 10^{123}$</td>
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<tr>
<td>Go</td>
<td>$\approx 250$</td>
<td>$\approx 150$</td>
<td>$\approx 10^{359}$</td>
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</table>

$b$ number of moves per ply/turn
$d$ length of game in plies/turns
$b^d$ number of possible sequences
Deep Learning: AlphaGo

- **AlphaGo** is a computer program developed by Alphabet Inc.’s Google DeepMind to play the board game Go.

- It uses a combination of machine learning and tree search techniques, combined with extensive training, both from human and computer play.

- AlphaGo uses Monte Carlo tree search, guided by a “value network” and a “policy network”, both of which are implemented using **deep neural networks**.

- A limited amount of game-specific feature detection is applied to the input before it is sent to the neural networks.

- The neural networks were bootstrapped from human gameplay experience. Later AlphaGo was set up to play against other instances of itself, using reinforcement learning to improve its play.

Deep Learning: AlphaGo

Match against **Fan Hui**
(Elo 3016 on 01-01-2016,
#512 world ranking list),
best European player
at the time of the match

AlphaGo wins 5 : 0

Match against **Lee Sedol**
(Elo 3546 on 01-01-2016,
#1 world ranking list 2007–2010,
#3 at the time of the match)

AlphaGo wins 4 : 1

www.goratings.org

<table>
<thead>
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<th>GPUs</th>
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First game of the match between AlphaGo and Lee Sedol
Lee Sedol: black pieces (first move), AlphaGo: white pieces; AlphaGo won

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Radial Basis Function Networks
A radial basis function network (RBFN) is a neural network with a graph $G = (U, C)$ that satisfies the following conditions:

(i) $U_{\text{in}} \cap U_{\text{out}} = \emptyset$,

(ii) $C = (U_{\text{in}} \times U_{\text{hidden}}) \cup C', \quad C' \subseteq (U_{\text{hidden}} \times U_{\text{out}})$

The network input function of each hidden neuron is a distance function of the input vector and the weight vector, that is,

$$\forall u \in U_{\text{hidden}} : \quad f_{\text{net}}^{(u)}(\vec{w}_u, \vec{i}_u) = d(\vec{w}_u, \vec{i}_u),$$

where $d : \mathbb{R}^n \times \mathbb{R}^n \rightarrow \mathbb{R}_0^+$ is a function satisfying $\forall \vec{x}, \vec{y}, \vec{z} \in \mathbb{R}^n$:

(i) $d(\vec{x}, \vec{y}) = 0 \iff \vec{x} = \vec{y}$,

(ii) $d(\vec{x}, \vec{y}) = d(\vec{y}, \vec{x})$ (symmetry),

(iii) $d(\vec{x}, \vec{z}) \leq d(\vec{x}, \vec{y}) + d(\vec{y}, \vec{z})$ (triangle inequality).
**Distance Functions**

**Illustration of distance functions: Minkowski Family**

\[
d_k(x, y) = \left( \sum_{i=1}^{n} |x_i - y_i|^k \right)^{\frac{1}{k}}
\]

Well-known special cases from this family are:

- \( k = 1 \): Manhattan or city block distance,
- \( k = 2 \): Euclidean distance,
- \( k \to \infty \): maximum distance, that is, \( d_\infty(x, y) = \max_{i=1}^{n} |x_i - y_i| \).
The network input function of the output neurons is the weighted sum of their inputs:

\[ \forall u \in U_{\text{out}} : \quad f^{(u)}(\vec{w}_u, \vec{in}_u) = \vec{w}_u^\top \vec{in}_u = \sum_{v \in \text{pred}(u)} w_{uv} \text{out}_v. \]

The activation function of each hidden neuron is a so-called \textbf{radial function}, that is, a monotonically decreasing function

\[ f : \mathbb{R}_0^+ \rightarrow [0, 1] \quad \text{with} \quad f(0) = 1 \quad \text{and} \quad \lim_{x \to \infty} f(x) = 0. \]

The activation function of each output neuron is a linear function, namely

\[ f^{(u)}_{\text{act}}(\text{net}_u, \theta_u) = \text{net}_u - \theta_u. \]

(The linear activation function is important for the initialization.)
Radial Activation Functions

rectangle function:
\[ f_{\text{act}}(\text{net}, \sigma) = \begin{cases} 
0, & \text{if net} > \sigma, \\
1, & \text{otherwise}. 
\end{cases} \]

triangle function:
\[ f_{\text{act}}(\text{net}, \sigma) = \begin{cases} 
0, & \text{if net} > \sigma, \\
1 - \frac{\text{net}}{\sigma}, & \text{otherwise}. 
\end{cases} \]

cosine until zero:
\[ f_{\text{act}}(\text{net}, \sigma) = \begin{cases} 
0, & \text{if net} > 2\sigma, \\
\frac{\cos\left(\frac{\pi}{2\sigma} \text{net}\right)+1}{2}, & \text{otherwise}. 
\end{cases} \]

Gaussian function:
\[ f_{\text{act}}(\text{net}, \sigma) = e^{-\frac{\text{net}^2}{2\sigma^2}} \]
Radial basis function networks for the conjunction $x_1 \land x_2$
Radial basis function networks for the biimplication $x_1 \leftrightarrow x_2$

Idea: logical decomposition

$$x_1 \leftrightarrow x_2 \equiv (x_1 \land x_2) \lor \neg(x_1 \lor x_2)$$
Approximation of a function by rectangular pulses, each of which can be represented by a neuron of an radial basis function network.
A radial basis function network that computes the step function on the preceding slide and the piecewise linear function on the next slide (depends on activation function).
Approximation of a function by triangular pulses, each of which can be represented by a neuron of an radial basis function network.
Approximation of a function by Gaussian functions with radius $\sigma = 1$. It is $w_1 = 1$, $w_2 = 3$ and $w_3 = -2$. 
Radial Basis Function Networks: Function Approximation

Radial basis function network for a sum of three Gaussian functions

- The weights of the connections from the input neuron to the hidden neurons determine the locations of the Gaussian functions.
- The weights of the connections from the hidden neurons to the output neuron determine the height/direction (upward or downward) of the Gaussian functions.
Training Radial Basis Function Networks
Let $L_{\text{fixed}} = \{l_1, \ldots, l_m\}$ be a fixed learning task, consisting of $m$ training patterns $l = (\vec{\mathbf{1}}(l), \vec{\mathbf{o}}(l))$.

**Simple radial basis function network:**
One hidden neuron $v_k$, $k = 1, \ldots, m$, for each training pattern:

$$\forall k \in \{1, \ldots, m\} : \quad \vec{w}_v = \vec{\mathbf{1}}(l_k).$$

If the activation function is the Gaussian function, the radii $\sigma_k$ are chosen heuristically

$$\forall k \in \{1, \ldots, m\} : \quad \sigma_k = \frac{d_{\text{max}}}{\sqrt{2m}},$$

where

$$d_{\text{max}} = \max_{l_j, l_k \in L_{\text{fixed}}} d\left(\vec{\mathbf{1}}(l_j), \vec{\mathbf{1}}(l_k)\right).$$
Initializing the connections from the hidden to the output neurons

\[ \forall u : \sum_{k=1}^{m} w_{uw} \text{out}_{v_m}^{(l)} - \theta_u = o_u^{(l)} \quad \text{or abbreviated} \quad \mathbf{A} \cdot \mathbf{w}_u = \mathbf{o}_u, \]

where \( \mathbf{o}_u = (o_u^{(l_1)}, \ldots, o_u^{(l_m)})^\top \) is the vector of desired outputs, \( \theta_u = 0 \), and

\[
\mathbf{A} = \begin{pmatrix}
\text{out}^{(l_1)}_{v_1} & \text{out}^{(l_1)}_{v_2} & \ldots & \text{out}^{(l_1)}_{v_m} \\
\text{out}^{(l_2)}_{v_1} & \text{out}^{(l_2)}_{v_2} & \ldots & \text{out}^{(l_2)}_{v_m} \\
\vdots & \vdots & \ddots & \vdots \\
\text{out}^{(l_m)}_{v_1} & \text{out}^{(l_m)}_{v_2} & \ldots & \text{out}^{(l_m)}_{v_m}
\end{pmatrix}.
\]

This is a linear equation system, that can be solved by inverting the matrix \( \mathbf{A} \):

\[ \mathbf{w}_u = \mathbf{A}^{-1} \cdot \mathbf{o}_u. \]
Simple radial basis function network for the biimplication $x_1 \leftrightarrow x_2$
Simple radial basis function network for the biimplication \( x_1 \leftrightarrow x_2 \)

\[
A = \begin{pmatrix}
1 & e^{-2} & e^{-2} & e^{-4} \\
e^{-2} & 1 & e^{-4} & e^{-2} \\
e^{-2} & e^{-4} & 1 & e^{-2} \\
e^{-4} & e^{-2} & e^{-2} & 1
\end{pmatrix}
\]

\[
A^{-1} = \begin{pmatrix}
a & b & b & c \\
\bar{D} & \bar{D} & \bar{D} & \bar{D} \\
b & a & c & b \\
\bar{D} & \bar{D} & a & b \\
c & b & b & \bar{a} \\
\bar{D} & \bar{D} & \bar{D} & \bar{D}
\end{pmatrix}
\]

where

\[
\bar{D} = 1 - 4e^{-4} + 6e^{-8} - 4e^{-12} + e^{-16} \approx 0.9287
\]

\[
a = 1 - 2e^{-4} + e^{-8} \approx 0.9637
\]

\[
b = -e^{-2} + 2e^{-6} - e^{-10} \approx -0.1304
\]

\[
c = e^{-4} - 2e^{-8} + e^{-12} \approx 0.0177
\]

\[
\bar{w}_u = A^{-1} \cdot \bar{o}_u = \frac{1}{\bar{D}} \begin{pmatrix}
a + c \\
2b \\
a + c
\end{pmatrix} \approx \begin{pmatrix}
1.0567 \\
-0.2809 \\
-0.2809
\end{pmatrix}
\]
RBFN Initialization: Example

Simple radial basis function network for the biimplication $x_1 \leftrightarrow x_2$

- Initialization leads already to a perfect solution of the learning task.
- Subsequent training is not necessary.
Normal radial basis function networks:
Select subset of $k$ training patterns as centers.

$$
A = \begin{pmatrix}
1 & \text{out}_{v_1}^{(l_1)} & \text{out}_{v_2}^{(l_1)} & \ldots & \text{out}_{v_k}^{(l_1)} \\
1 & \text{out}_{v_1}^{(l_2)} & \text{out}_{v_2}^{(l_2)} & \ldots & \text{out}_{v_k}^{(l_2)} \\
\vdots & \vdots & \vdots & \ddots & \vdots \\
1 & \text{out}_{v_1}^{(l_m)} & \text{out}_{v_2}^{(l_m)} & \ldots & \text{out}_{v_k}^{(l_m)} \\
\end{pmatrix}
$$

Compute (Moore–Penrose) pseudo inverse:

$$
A^+ = (A^T A)^{-1} A^T.
$$

The weights can then be computed by

$$
\tilde{w}_u = A^+ \cdot \tilde{o}_u = (A^T A)^{-1} A^T \cdot \tilde{o}_u.
$$
Normal radial basis function network for the biimplication $x_1 \leftrightarrow x_2$

Select two training patterns:

- $l_1 = (\vec{i}(l_1), \vec{o}(l_1)) = ((0, 0), (1))$
- $l_4 = (\vec{i}(l_4), \vec{o}(l_4)) = ((1, 1), (1))$
Normal radial basis function network for the biimplication $x_1 \leftrightarrow x_2$

$$A = \begin{pmatrix} 1 & 1 & e^{-4} \\ 1 & e^{-2} & e^{-2} \\ 1 & e^{-2} & e^{-2} \\ 1 & e^{-4} & 1 \end{pmatrix}$$

$$A^+ = (A^T A)^{-1} A^T = \begin{pmatrix} a & b & b & a \\ c & d & d & e \\ e & d & d & c \end{pmatrix}$$

where

$$a \approx -0.1810, \quad b \approx 0.6810, \quad c \approx 1.1781, \quad d \approx -0.6688, \quad e \approx 0.1594.$$ 

Resulting weights:

$$\vec{w}_u = \begin{pmatrix} -\theta \\ w_1 \\ w_2 \end{pmatrix} = A^+ \cdot \vec{o}_u \approx \begin{pmatrix} -0.3620 \\ 1.3375 \\ 1.3375 \end{pmatrix}.$$
RBFN Initialization: Example

Normal radial basis function network for the biimplication $x_1 \leftrightarrow x_2$

- Basis function $(0,0)$
- Basis function $(1,1)$
- Output

- Initialization leads already to a perfect solution of the learning task.
- This is an accident, because the linear equation system is not over-determined, due to linearly dependent equations.
How to choose the radial basis function centers?

- Use **all data points** as centers for the radial basis functions.
  - Advantages: Only radius and output weights need to be determined; desired output values can be achieved exactly (unless there are inconsistencies).
  - Disadvantage: Often much too many radial basis functions; computing the weights to the output neuron via a pseudo-inverse can become infeasible.

- Use a **random subset** of data points as centers for the radial basis functions.
  - Advantages: Fast; only radius and output weights need to be determined.
  - Disadvantages: Performance depends heavily on the choice of data points.

- Use the **result of clustering** as centers for the radial basis functions, e.g.
  - $c$-means clustering (on the next slides)
  - Learning vector quantization (to be discussed later)
RBFN Initialization: $c$-means Clustering

- Choose a number $c$ of clusters to be found (user input).

- Initialize the cluster centers randomly
  (for instance, by randomly selecting $c$ data points).

- **Data point assignment:**
  Assign each data point to the cluster center that is closest to it
  (that is, closer than any other cluster center).

- **Cluster center update:**
  Compute new cluster centers as the mean vectors of the assigned data points.
  (Intuitively: center of gravity if each data point has unit weight.)

- Repeat these two steps (data point assignment and cluster center update)
  until the clusters centers do not change anymore.

It can be shown that this scheme must converge,
that is, the update of the cluster centers cannot go on forever.
**c-Means Clustering: Example**

Data set to cluster.
Choose $c = 3$ clusters.
(From visual inspection, can be difficult to determine in general.)

Initial position of cluster centers.
Randomly selected data points.
(Alternative methods include e.g. latin hypercube sampling)
Delaunay Triangulations and Voronoi Diagrams

- Dots represent cluster centers.
- **Left:** Delaunay Triangulation
  The circle through the corners of a triangle does not contain another point.
- **Right:** Voronoi Diagram / Tessellation
  Midperpendiculars of the Delaunay triangulation: boundaries of the regions of points that are closest to the enclosed cluster center (Voronoi cells).
• **Delaunay Triangulation**: simple triangle (shown in gray on the left)

• **Voronoi Diagram**: midperpendiculars of the triangle’s edges (shown in blue on the left, in gray on the right)
c-Means Clustering: Example
Training radial basis function networks:
Derivation of update rules is analogous to that of multi-layer perceptrons.

Weights from the hidden to the output neurons.

Gradient:

$$\nabla \tilde{w}_u e^{(l)}_u = \frac{\partial e^{(l)}_u}{\partial \tilde{w}_u} = -2(o^{(l)}_u - \text{out}^{(l)}_u) \tilde{\text{in}}^{(l)}_u,$$

Weight update rule:

$$\Delta \tilde{w}^{(l)}_u = -\eta \frac{3}{2} \nabla \tilde{w}_u e^{(l)}_u = \eta_3(o^{(l)}_u - \text{out}^{(l)}_u) \tilde{\text{in}}^{(l)}_u$$

Typical learning rate: $\eta_3 \approx 0.001$.

(Two more learning rates are needed for the center coordinates and the radii.)
Training radial basis function networks:
Center coordinates (weights from the input to the hidden neurons).

Gradient:
\[
\nabla_{\vec{w}_v} e^{(l)} = \frac{\partial e^{(l)}}{\partial \vec{w}_v} = -2 \sum_{s \in \text{succ}(v)} (o_s^{(l)} - \text{out}_s^{(l)}) w_{su} \frac{\partial \text{out}_v^{(l)}}{\partial \text{net}_v^{(l)}} \frac{\partial \text{net}_v^{(l)}}{\partial \vec{w}_v}
\]

Weight update rule:
\[
\Delta \vec{w}_v^{(l)} = -\frac{\eta_1}{2} \nabla_{\vec{w}_v} e^{(l)} = \eta_1 \sum_{s \in \text{succ}(v)} (o_s^{(l)} - \text{out}_s^{(l)}) w_{sv} \frac{\partial \text{out}_v^{(l)}}{\partial \text{net}_v^{(l)}} \frac{\partial \text{net}_v^{(l)}}{\partial \vec{w}_v}
\]

Typical learning rate: \( \eta_1 \approx 0.02 \).
Training radial basis function networks:
Center coordinates (weights from the input to the hidden neurons).

Special case: **Euclidean distance**

\[
\frac{\partial \text{net}_v^{(l)}}{\partial \vec{w}_v} = \left( \sum_{i=1}^{n} (w_{vp_i} - \text{out}_{p_i}^{(l)})^2 \right)^{-\frac{1}{2}} (\vec{w}_v - \vec{in}_v^{(l)}).
\]

Special case: **Gaussian activation function**

\[
\frac{\partial \text{out}_v^{(l)}}{\partial \text{net}_v^{(l)}} = \frac{\partial f_{\text{act}}(\text{net}_v^{(l)}, \sigma_v)}{\partial \text{net}_v^{(l)}} = \frac{\partial}{\partial \text{net}_v^{(l)}} e^{-\frac{\left(\text{net}_v^{(l)}\right)^2}{2\sigma_v^2}} = -\frac{\text{net}_v^{(l)}}{\sigma_v^2} e^{-\frac{\left(\text{net}_v^{(l)}\right)^2}{2\sigma_v^2}}.
\]
Training radial basis function networks:
Radii of radial basis functions.

Gradient:
\[
\frac{\partial e^{(l)}}{\partial \sigma_v} = -2 \sum_{s \in \text{succ}(v)} (o_s^{(l)} - \text{out}_s^{(l)}) w_{su} \frac{\partial \text{out}_v^{(l)}}{\partial \sigma_v}.
\]

Weight update rule:
\[
\Delta \sigma_v^{(l)} = -\frac{\eta_2}{2} \frac{\partial e^{(l)}}{\partial \sigma_v} = \eta_2 \sum_{s \in \text{succ}(v)} (o_s^{(l)} - \text{out}_s^{(l)}) w_{sv} \frac{\partial \text{out}_v^{(l)}}{\partial \sigma_v}.
\]

Typical learning rate: \( \eta_2 \approx 0.01 \).
Training radial basis function networks:
Radii of radial basis functions.

Special case: **Gaussian activation function**

\[
\frac{\partial \text{out}_v^{(l)}}{\partial \sigma_v} = \frac{\partial}{\partial \sigma_v} e^{-\frac{(\text{net}_v^{(l)})^2}{2\sigma_v^2}} = \frac{(\text{net}_v^{(l)})^2}{\sigma_v^3} e^{-\frac{(\text{net}_v^{(l)})^2}{2\sigma_v^2}}.
\]

(The distance function is irrelevant for the radius update, since it only enters the network input function.)
Generalization of the distance function

Idea: Use anisotropic (direction dependent) distance function.

Example: **Mahalanobis distance**

\[ d(\vec{x}, \vec{y}) = \sqrt{(\vec{x} - \vec{y})^\top \Sigma^{-1} (\vec{x} - \vec{y})}. \]

Example: **biimplication**

\[
\begin{align*}
\Sigma &= \begin{pmatrix} 9 & 8 \\ 8 & 9 \end{pmatrix} \\
\end{align*}
\]
Application: Recognition of Handwritten Digits

- Images of 20,000 handwritten digits (2,000 per class), split into training and test data set of 10,000 samples each (1,000 per class).
- Represented in a normalized fashion as $16 \times 16$ gray values in $\{0, \ldots, 255\}$.
- Data was originally used in the StatLog project [Michie et al. 1994].
Application: Recognition of Handwritten Digits

- **Comparison of various classifiers:**
  - Nearest Neighbor (1NN)
  - Decision Tree (C4.5)
  - Multi-Layer Perceptron (MLP)
  - Learning Vector Quantization (LVQ)
  - Radial Basis Function Network (RBF)
  - Support Vector Machine (SVM)

- **Distinction of the number of RBF training phases:**
  - 1 phase: find output connection weights e.g. with pseudo-inverse.
  - 2 phase: find RBF centers e.g. with some clustering plus 1 phase.
  - 3 phase: 2 phase plus error backpropagation training.

- **Initialization of radial basis function centers:**
  - Random choice of data points
  - $c$-means Clustering
  - Learning Vector Quantization
  - Decision Tree (one RBF center per leaf)
• The 60 cluster centers (6 per class) resulting from $c$-means clustering. (Clustering was conducted with $c = 6$ for each class separately.)

• Initial cluster centers were selected randomly from the training data.

• The weights of the connections to the output neuron were computed with the pseudo-inverse method.
The 60 cluster centers (6 per class) after training the radial basis function network with error backpropagation.

Differences between the initial and the trained centers of the radial basis functions appear to be fairly small, but ...
Application: Recognition of Handwritten Digits

- Distance matrices showing the Euclidean distances of the 60 radial basis function centers before and after training.
- Centers are sorted by class/digit: first 6 rows/columns refer to digit 0, next 6 rows/columns to digit 1 etc.
- Distances are encoded as gray values: darker means smaller distance.

picture not available in online version
Application: Recognition of Handwritten Digits

- Before training (left): many distances between centers of different classes/digits are small (e.g. 2-3, 3-8, 3-9, 5-8, 5-9), which increases the chance of misclassifications.

- After training (right): only very few small distances between centers of different classes/digits; basically all small distances between centers of same class/digit.
## Application: Recognition of Handwritten Digits

### Classification results:

<table>
<thead>
<tr>
<th>Classifier</th>
<th>Accuracy</th>
</tr>
</thead>
<tbody>
<tr>
<td>Nearest Neighbor (1NN)</td>
<td>97.68%</td>
</tr>
<tr>
<td>Learning Vector Quantization (LVQ)</td>
<td>96.99%</td>
</tr>
<tr>
<td>Decision Tree (C4.5)</td>
<td>91.12%</td>
</tr>
<tr>
<td>2-Phase-RBF (data points)</td>
<td>95.24%</td>
</tr>
<tr>
<td>2-Phase-RBF (c-means)</td>
<td>96.94%</td>
</tr>
<tr>
<td>2-Phase-RBF (LVQ)</td>
<td>95.86%</td>
</tr>
<tr>
<td>2-Phase-RBF (C4.5)</td>
<td>92.72%</td>
</tr>
<tr>
<td>3-Phase-RBF (data points)</td>
<td>97.23%</td>
</tr>
<tr>
<td>3-Phase-RBF (c-means)</td>
<td>98.06%</td>
</tr>
<tr>
<td>3-Phase-RBF (LVQ)</td>
<td>98.49%</td>
</tr>
<tr>
<td>3-Phase-RBF (C4.5)</td>
<td>94.83%</td>
</tr>
<tr>
<td>Support Vector Machine (SVM)</td>
<td>98.76%</td>
</tr>
<tr>
<td>Multi-Layer Perceptron (MLP)</td>
<td>97.59%</td>
</tr>
</tbody>
</table>

- LVQ: 200 vectors (20 per class)
- C4.5: 505 leaves
- c-means: 60 centers (6 per class)
- SVM: 10 classifiers, ≈ 4200 vectors
- MLP: 1 hidden layer with 200 neurons

- Results are medians of three training/test runs.
- Error backpropagation improves RBF results.
Learning Vector Quantization
Learning Vector Quantization

• Up to now: **fixed learning tasks**
  ○ The data consists of input/output pairs.
  ○ The objective is to produce desired output for given input.
  ○ This allows to describe training as error minimization.

• Now: **free learning tasks**
  ○ The data consists only of input values/vectors.
  ○ The objective is to produce similar output for similar input (clustering).

• **Learning Vector Quantization**
  ○ Find a suitable quantization (many-to-few mapping, often to a finite set) of the input space, e.g. a tessellation of a Euclidean space.
  ○ Training adapts the coordinates of so-called reference or codebook vectors, each of which defines a region in the input space.
Dots represent vectors that are used for quantizing the area.

Left: **Delaunay Triangulation**
(The circle through the corners of a triangle does not contain another point.)

Right: **Voronoi Diagram / Tessellation**
(Midperpendiculars of the Delaunay triangulation: boundaries of the regions of points that are closest to the enclosed cluster center (Voronoi cells)).
Finding clusters in a given set of data points

- Data points are represented by empty circles (○).
- Cluster centers are represented by full circles (●).
A learning vector quantization network (LVQ) is a neural network with a graph $G = (U, C)$ that satisfies the following conditions

(i) $U_{\text{in}} \cap U_{\text{out}} = \emptyset$, $U_{\text{hidden}} = \emptyset$

(ii) $C = U_{\text{in}} \times U_{\text{out}}$

The network input function of each output neuron is a distance function of the input vector and the weight vector, that is,

$$\forall u \in U_{\text{out}} : \quad f^{(u)}_{\text{net}}(\vec{w}_u, \vec{u}) = d(\vec{w}_u, \vec{u}),$$

where $d : \mathbb{R}^n \times \mathbb{R}^n \to \mathbb{R}_0^+$ is a function satisfying $\forall \vec{x}, \vec{y}, \vec{z} \in \mathbb{R}^n :$

(i) $d(\vec{x}, \vec{y}) = 0 \iff \vec{x} = \vec{y},$

(ii) $d(\vec{x}, \vec{y}) = d(\vec{y}, \vec{x})$ (symmetry),

(iii) $d(\vec{x}, \vec{z}) \leq d(\vec{x}, \vec{y}) + d(\vec{y}, \vec{z})$ (triangle inequality).
Illustration of distance functions: Minkowski family

\[ d_k(\vec{x}, \vec{y}) = \left( \sum_{i=1}^{n} |x_i - y_i|^k \right)^{\frac{1}{k}} \]

Well-known special cases from this family are:

- \( k = 1 \): Manhattan or city block distance,
- \( k = 2 \): Euclidean distance,
- \( k \to \infty \): maximum distance, that is, \( d_\infty(\vec{x}, \vec{y}) = \max_{i=1}^{n} |x_i - y_i| \).
The activation function of each output neuron is a so-called radial function, that is, a monotonically decreasing function

\[ f : \mathbb{R}_0^+ \rightarrow [0, \infty] \quad \text{with} \quad f(0) = 1 \quad \text{and} \quad \lim_{x \to \infty} f(x) = 0. \]

Sometimes the range of values is restricted to the interval \([0, 1]\). However, due to the special output function this restriction is irrelevant.

The output function of each output neuron is not a simple function of the activation of the neuron. Rather it takes into account the activations of all output neurons:

\[
f^{(u)}_{\text{out}}(\text{act}_u) = \begin{cases} 
1, & \text{if } \text{act}_u = \max_{v \in U_{\text{out}}} \text{act}_v, \\
0, & \text{otherwise}.
\end{cases}
\]

If more than one unit has the maximal activation, one is selected at random to have an output of 1, all others are set to output 0: \textit{winner-takes-all principle}. 

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rectangle function:
\[ f_{\text{act}}(\text{net}, \sigma) = \begin{cases} 
0, & \text{if net} > \sigma, \\
1, & \text{otherwise}. 
\end{cases} \]

cosine until zero:
\[ f_{\text{act}}(\text{net}, \sigma) = \begin{cases} 
0, & \text{if net} > 2\sigma, \\
\frac{\cos\left(\frac{\pi}{2\sigma}\text{net}\right) + 1}{2}, & \text{otherwise}. 
\end{cases} \]

triangle function:
\[ f_{\text{act}}(\text{net}, \sigma) = \begin{cases} 
0, & \text{if net} > \sigma, \\
1 - \frac{\text{net}}{\sigma}, & \text{otherwise}. 
\end{cases} \]

Gaussian function:
\[ f_{\text{act}}(\text{net}, \sigma) = e^{-\frac{\text{net}^2}{2\sigma^2}}. \]
Adaptation of reference vectors / codebook vectors

- For each training pattern find the closest reference vector.
- Adapt only this reference vector (winner neuron).
- For classified data the class may be taken into account: Each reference vector is assigned to a class.

**Attraction rule** (data point and reference vector have same class)

\[
\vec{r}^{\text{(new)}} = \vec{r}^{\text{(old)}} + \eta (\vec{x} - \vec{r}^{\text{(old)}}),
\]

**Repulsion rule** (data point and reference vector have different class)

\[
\vec{r}^{\text{(new)}} = \vec{r}^{\text{(old)}} - \eta (\vec{x} - \vec{r}^{\text{(old)}}).
\]
Learning Vector Quantization

Adaptation of reference vectors / codebook vectors

- $\vec{x}$: data point, $\vec{r}_i$: reference vector
- $\eta = 0.4$ (learning rate)
Adaptation of reference vectors / codebook vectors

- Left: Online training with learning rate $\eta = 0.1$,
- Right: Batch training with learning rate $\eta = 0.05$. 
Problem: fixed learning rate can lead to oscillations

Solution: time dependent learning rate

\[ \eta(t) = \eta_0 \alpha^t, \quad 0 < \alpha < 1, \quad \text{or} \quad \eta(t) = \eta_0 t^{\kappa}, \quad \kappa < 0. \]
Improved update rule for classified data

- **Idea:** Update not only the one reference vector that is closest to the data point (the winner neuron), but **update the two closest reference vectors.**

- Let $\vec{x}$ be the currently processed data point and $c$ its class. Let $\vec{r}_j$ and $\vec{r}_k$ be the two closest reference vectors and $z_j$ and $z_k$ their classes.

- Reference vectors are updated only if $z_j \neq z_k$ and either $c = z_j$ or $c = z_k$. (Without loss of generality we assume $c = z_j$.)

The **update rules** for the two closest reference vectors are:

$$
\begin{align*}
\vec{r}_j^{(\text{new})} &= \vec{r}_j^{(\text{old})} + \eta(\vec{x} - \vec{r}_j^{(\text{old})}) \\
\vec{r}_k^{(\text{new})} &= \vec{r}_k^{(\text{old})} - \eta(\vec{x} - \vec{r}_k^{(\text{old})}),
\end{align*}
$$

while all other reference vectors remain unchanged.
It was observed in practical tests that standard learning vector quantization may drive the reference vectors further and further apart.

To counteract this undesired behavior a **window rule** was introduced: update only if the data point $\vec{x}$ is close to the classification boundary.

“Close to the boundary” is made formally precise by requiring

$$\min\left(\frac{d(\vec{x}, \vec{r}_j)}{d(\vec{x}, \vec{r}_k)}, \frac{d(\vec{x}, \vec{r}_k)}{d(\vec{x}, \vec{r}_j)}\right) > \theta,$$

where

$$\theta = \frac{1 - \xi}{1 + \xi}.$$

$\xi$ is a parameter that has to be specified by a user.

Intuitively, $\xi$ describes the “width” of the window around the classification boundary, in which the data point has to lie in order to lead to an update.

Using it prevents divergence, because the update ceases for a data point once the classification boundary has been moved far enough away.
Soft Learning Vector Quantization

- **Idea:** Use soft assignments instead of winner-takes-all (approach described here: [Seo and Obermayer 2003]).

- **Assumption:** Given data was sampled from a mixture of normal distributions. Each reference vector describes one normal distribution.

- Closely related to clustering by estimating a **mixture of Gaussians**.
  - (Crisp or hard) learning vector quantization can be seen as an “online version” of $c$-means clustering.
  - Soft learning vector quantization can be seen as an “online version” of estimating a mixture of Gaussians (that is, of normal distributions). (In the following: brief review of the Expectation Maximization (EM) Algorithm for estimating a mixture of Gaussians.)

- Hardening soft learning vector quantization (by letting the “radii” of the Gaussians go to zero, see below) yields a version of (crisp or hard) learning vector quantization that works well without a window rule.
Expectation Maximization: Mixture of Gaussians

- **Assumption:** Data was generated by sampling a set of normal distributions.
  (The probability density is a mixture of Gaussian distributions.)

- **Formally:** We assume that the probability density can be described as

\[
f_{\vec{X}}(\vec{x}; C) = \sum_{y=1}^{c} f_{\vec{X},Y}(\vec{x}, y; C) = \sum_{y=1}^{c} p_Y(y; C) \cdot f_{\vec{X}|Y}(\vec{x}|y; C).
\]

- $\mathbf{C}$ is the set of cluster parameters
- $\vec{X}$ is a random vector that has the data space as its domain
- $Y$ is a random variable that has the cluster indices as possible values (i.e., $\text{dom}(\vec{X}) = \mathbb{R}^m$ and $\text{dom}(Y) = \{1, \ldots, c\}$)
- $p_Y(y; C)$ is the probability that a data point belongs to (is generated by) the $y$-th component of the mixture
- $f_{\vec{X}|Y}(\vec{x}|y; C)$ is the conditional probability density function of a data point given the cluster (specified by the cluster index $y$)
**Expectation Maximization**

- **Basic idea**: Do a maximum likelihood estimation of the cluster parameters.

- **Problem**: The likelihood function,

\[
L(X; C) = \prod_{j=1}^{n} f_{\vec{X}_j}(\vec{x}_j; C) = \prod_{j=1}^{n} c \sum_{y=1}^{c} p_Y(y; C) \cdot f_{\vec{X}|Y}(\vec{x}_j|y; C),
\]

is difficult to optimize, even if one takes the natural logarithm (cf. the maximum likelihood estimation of the parameters of a normal distribution), because

\[
\ln L(X; C) = \sum_{j=1}^{n} \ln \sum_{y=1}^{c} p_Y(y; C) \cdot f_{\vec{X}|Y}(\vec{x}_j|y; C)
\]

contains the natural logarithms of complex sums.

- **Approach**: Assume that there are “hidden” variables \( Y_j \) stating the clusters that generated the data points \( \vec{x}_j \), so that the sums reduce to one term.

- **Problem**: Since the \( Y_j \) are hidden, we do not know their values.
• **Formally:** Maximize the likelihood of the “completed” data set $(X, \vec{y})$, where $\vec{y} = (y_1, \ldots, y_n)$ combines the values of the variables $Y_j$. That is,

$$
L(X, \vec{y}; C) = \prod_{j=1}^{n} f_{\vec{X}_j,Y_j}(\vec{x}_j, y_j; C) = \prod_{j=1}^{n} p_{Y_j}(y_j; C) \cdot f_{\vec{X}_j|Y_j}(\vec{x}_j|y_j; C).
$$

• **Problem:** Since the $Y_j$ are hidden, the values $y_j$ are unknown (and thus the factors $p_{Y_j}(y_j; C)$ cannot be computed).

• **Approach to find a solution nevertheless:**
  
  ◦ See the $Y_j$ as random variables (the values $y_j$ are not fixed) and consider a probability distribution over the possible values.
  
  ◦ As a consequence $L(X, \vec{y}; C)$ becomes a random variable, even for a fixed data set $X$ and fixed cluster parameters $C$.

  ◦ Try to **maximize the expected value** of $L(X, \vec{y}; C)$ or $\ln L(X, \vec{y}; C)$ (hence the name **expectation maximization**).
• **Formally:** Find the cluster parameters as

\[
\hat{C} = \arg\max_C E(\ln L(X, \vec{y}; C) | X; C),
\]

that is, maximize the expected likelihood

\[
E(L(X, \vec{y}; C) | X; C) = \sum_{\vec{y} \in \{1, \ldots, c\}^n} p_{Y|X}(\vec{y}|X; C) \cdot \prod_{j=1}^n f_{X_j,Y_j}(x_j, y_j; C)
\]

or, alternatively, maximize the expected log-likelihood

\[
E(\ln L(X, \vec{y}; C) | X; C) = \sum_{\vec{y} \in \{1, \ldots, c\}^n} p_{Y|X}(\vec{y}|X; C) \cdot \sum_{j=1}^n \ln f_{X_j,Y_j}(x_j, y_j; C).
\]

• Unfortunately, these functionals are still difficult to optimize directly.

• **Solution:** Use the equation as an iterative scheme, fixing \(C\) in some terms (iteratively compute better approximations, similar to Heron’s algorithm).
Excursion: Heron’s Algorithm

- **Task:** Find the square root of a given number \( x \), i.e., find \( y = \sqrt{x} \).

- **Approach:** Rewrite the defining equation \( y^2 = x \) as follows:

\[
y^2 = x \iff 2y^2 = y^2 + x \iff y = \frac{1}{2y}(y^2 + x) \iff y = \frac{1}{2} \left( y + \frac{x}{y} \right).
\]

- Use the resulting equation as an iteration formula, i.e., compute the sequence

\[
y_{k+1} = \frac{1}{2} \left( y_k + \frac{x}{y_k} \right) \quad \text{with} \quad y_0 = 1.
\]

- It can be shown that \( 0 \leq y_k - \sqrt{x} \leq y_{k-1} - y_n \) for \( k \geq 2 \). Therefore this iteration formula provides increasingly better approximations of the square root of \( x \) and thus is a safe and simple way to compute it.

Ex.: \( x = 2 \): \( y_0 = 1, y_1 = 1.5, y_2 \approx 1.41667, y_3 \approx 1.414216, y_4 \approx 1.414213 \).

- Heron’s algorithm converges very quickly and is often used in pocket calculators and microprocessors to implement the square root.
Iterative scheme for expectation maximization:

Choose some initial set \( C_0 \) of cluster parameters and then compute

\[
C_{k+1} = \arg\max_C E(\ln L(X, \vec{y}; C) \mid X; C_k)
\]

\[
= \arg\max_C \sum_{\vec{y} \in \{1, \ldots, c\}^n} p_{\vec{y}|X}(\vec{y} \mid X; C_k) \sum_{j=1}^n \ln f_{\vec{X}_j, Y_j}(\vec{x}_j, y_j; C)
\]

\[
= \arg\max_C \sum_{\vec{y} \in \{1, \ldots, c\}^n} \left( \prod_{l=1}^n p_{Y_l|X_l}(y_l \mid \vec{x}_l; C_k) \right) \sum_{j=1}^n \ln f_{\vec{X}_j, Y_j}(\vec{x}_j, y_j; C)
\]

\[
= \arg\max_C \sum_{i=1}^c \sum_{j=1}^n p_{Y_j|X_j}(i \mid \vec{x}_j; C_k) \cdot \ln f_{\vec{X}_j, Y_j}(\vec{x}_j, i; C).
\]

It can be shown that each EM iteration increases the likelihood of the data and that the algorithm converges to a local maximum of the likelihood function (i.e., EM is a safe way to maximize the likelihood function).
Expectation Maximization

Justification of the last step on the previous slide:

\[
\sum_{\bar{y} \in \{1, \ldots, c\}^n} \left( \prod_{l=1}^{n} p_{Y_l|X_l}(y_l|\bar{x}_l; C_k) \right) \sum_{j=1}^{n} \ln f_{\bar{X}_j,Y_j}(\bar{x}_j,y_j; C) \\
= \sum_{y_1=1}^{c} \cdots \sum_{y_n=1}^{c} \left( \prod_{l=1}^{n} p_{Y_l|X_l}(y_l|\bar{x}_l; C_k) \right) \sum_{j=1}^{n} \sum_{i=1}^{c} \delta_{i,y_j} \ln f_{\bar{X}_j,Y_j}(\bar{x}_j,i; C) \\
= \sum_{i=1}^{c} \sum_{j=1}^{n} \ln f_{\bar{X}_j,Y_j}(\bar{x}_j,i; C) \sum_{y_1=1}^{c} \cdots \sum_{y_n=1}^{c} \delta_{i,y_j} \prod_{l=1}^{n} p_{Y_l|X_l}(y_l|\bar{x}_l; C_k) \\
= \sum_{i=1}^{c} \sum_{j=1}^{n} p_{Y_j|X_j}(i|\bar{x}_j; C_k) \cdot \ln f_{\bar{X}_j,Y_j}(\bar{x}_j,i; C) \\
\sum_{y_1=1}^{c} \cdots \sum_{y_{j-1}=1}^{c} \sum_{y_{j+1}=1}^{c} \cdots \sum_{y_n=1}^{c} \prod_{l=1}^{n} p_{Y_l|X_l}(y_l|\bar{x}_l; C_k) \\
= \prod_{l=1,l \neq j}^{n} \sum_{y_l=1}^{c} p_{Y_l|X_l}(y_l|\bar{x}_l; C_k) = \prod_{l=1,l \neq j}^{n} 1 = 1
\]
The probabilities $p_{Y_j|\tilde{X}_j}(i|\tilde{x}_j; C_k)$ are computed as

$$p_{Y_j|\tilde{X}_j}(i|\tilde{x}_j; C_k) = \frac{f_{\tilde{X}_j,Y_j}(\tilde{x}_j, i; C_k)}{f_{\tilde{X}_j}(\tilde{x}_j; C_k)} = \frac{f_{\tilde{X}_j|Y_j}(\tilde{x}_j|i; C_k) \cdot p_{Y_j}(i; C_k)}{\sum_{l=1}^{c} f_{\tilde{X}_j|Y_j}(\tilde{x}_j|l; C_k) \cdot p_{Y_j}(l; C_k)},$$

that is, as the relative probability densities of the different clusters (as specified by the cluster parameters) at the location of the data points $\tilde{x}_j$.

The $p_{Y_j|\tilde{X}_j}(i|\tilde{x}_j; C_k)$ are the posterior probabilities of the clusters given the data point $\tilde{x}_j$ and a set of cluster parameters $C_k$.

They can be seen as case weights of a “completed” data set:

- Split each data point $\tilde{x}_j$ into $c$ data points $((\tilde{x}_j, i), i = 1, \ldots, c$.
- Distribute the unit weight of the data point $\tilde{x}_j$ according to the above probabilities, i.e., assign to $(\tilde{x}_j, i)$ the weight $p_{Y_j|\tilde{X}_j}(i|\tilde{x}_j; C_k), i = 1, \ldots, c$. 
Core Iteration Formula

\[ C_{k+1} = \arg\max_C \sum_{c} \sum_{i=1}^{n} \sum_{j=1}^{n} p_{Y_j|\vec{X}_j}(i|\vec{x}_j; C_k) \cdot \ln f_{\vec{X}_j,Y_j}(\vec{x}_j, i; C) \]

Expectation Step

- For all data points \( \vec{x}_j \):
  Compute for each normal distribution the probability \( p_{Y_j|\vec{X}_j}(i|\vec{x}_j; C_k) \)
  that the data point was generated from it
  (ratio of probability densities at the location of the data point).
  \( \rightarrow \) “weight” of the data point for the estimation.

Maximization Step

- For all normal distributions:
  Estimate the parameters by standard maximum likelihood estimation
  using the probabilities (“weights”) assigned to the data points
  w.r.t. the distribution in the expectation step.
Expectation Maximization: Mixture of Gaussians

**Expectation Step:** Use Bayes’ rule to compute

\[
p_{C|\vec{X}}(i|\vec{x}; C) = \frac{p_C(i; c_i) \cdot f_{\vec{X}|C}(\vec{x}|i; c_i)}{f_{\vec{X}}(\vec{x}; C)} = \frac{p_C(i; c_i) \cdot f_{\vec{X}|C}(\vec{x}|i; c_i)}{\sum_{k=1}^{C} p_C(k; c_k) \cdot f_{\vec{X}|C}(\vec{x}|k; c_k)}.
\]

→ “weight” of the data point \( \vec{x} \) for the estimation.

**Maximization Step:** Use maximum likelihood estimation to compute

\[
\varrho_i^{(t+1)} = \frac{1}{n} \sum_{j=1}^{n} p_{C|\vec{X}_j}(i|\vec{x}_j; C^{(t)}), \quad \mu_i^{(t+1)} = \frac{\sum_{j=1}^{n} p_{C|\vec{X}_j}(i|\vec{x}_j; C^{(t)}) \cdot \vec{x}_j}{\sum_{j=1}^{n} p_{C|\vec{X}_j}(i|\vec{x}_j; C^{(t)})},
\]

and

\[
\Sigma_i^{(t+1)} = \frac{\sum_{j=1}^{n} p_{C|\vec{X}_j}(i|\vec{x}_j; C^{(t)}) \cdot (\vec{x}_j - \mu_i^{(t+1)}) (\vec{x}_j - \mu_i^{(t+1)})^\top}{\sum_{j=1}^{n} p_{C|\vec{X}_j}(i|\vec{x}_j; C^{(t)})}
\]

Iterate until convergence (checked, e.g., by change of mean vector).
Expectation Maximization: Technical Problems

- If a fully general mixture of Gaussian distributions is used, the likelihood function is truly optimized if
  - all normal distributions except one are contracted to single data points and
  - the remaining normal distribution is the maximum likelihood estimate for the remaining data points.

- This undesired result is rare, because the algorithm gets stuck in a local optimum.

- Nevertheless it is recommended to take countermeasures, which consist mainly in reducing the degrees of freedom, like
  - Fix the determinants of the covariance matrices to equal values.
  - Use a diagonal instead of a general covariance matrix.
  - Use an isotropic variance instead of a covariance matrix.
  - Fix the prior probabilities of the clusters to equal values.
Soft Learning Vector Quantization

**Idea:** Use soft assignments instead of winner-takes-all (approach described here: [Seo and Obermayer 2003]).

**Assumption:** Given data was sampled from a mixture of normal distributions. Each reference vector describes one normal distribution.

**Objective:** Maximize the log-likelihood ratio of the data, that is, maximize

\[
\ln L_{\text{ratio}} = \sum_{j=1}^{n} \ln \sum_{\vec{r} \in R(c_j)} \exp \left( -\frac{(\vec{x}_j - \vec{r})^\top (\vec{x}_j - \vec{r})}{2\sigma^2} \right) \\
- \sum_{j=1}^{n} \ln \sum_{\vec{r} \in Q(c_j)} \exp \left( -\frac{(\vec{x}_j - \vec{r})^\top (\vec{x}_j - \vec{r})}{2\sigma^2} \right).
\]

Here \(\sigma\) is a parameter specifying the “size” of each normal distribution. \(R(c)\) is the set of reference vectors assigned to class \(c\) and \(Q(c)\) its complement.

Intuitively: at each data point the probability density for its class should be as large as possible while the density for all other classes should be as small as possible.
Update rule derived from a maximum log-likelihood approach:

\[
\vec{r}_i^{(\text{new})} = \vec{r}_i^{(\text{old})} + \eta \cdot \begin{cases} 
    u_{ij}^\oplus (\vec{x}_j - \vec{r}_i^{(\text{old})}), & \text{if } c_j = z_i, \\
    -u_{ij}^\ominus (\vec{x}_j - \vec{r}_i^{(\text{old})}), & \text{if } c_j \neq z_i,
\end{cases}
\]

where \(z_i\) is the class associated with the reference vector \(\vec{r}_i\) and

\[
u_{ij}^\oplus = \frac{\exp \left( -\frac{1}{2\sigma^2} (\vec{x}_j - \vec{r}_i^{(\text{old})})^\top (\vec{x}_j - \vec{r}_i^{(\text{old})}) \right)}{\sum_{\vec{r}\in R(c_j)} \exp \left( -\frac{1}{2\sigma^2} (\vec{x}_j - \vec{r}^{(\text{old})})^\top (\vec{x}_j - \vec{r}^{(\text{old})}) \right)} \quad \text{and} \quad \nu_{ij}^\ominus = \frac{\exp \left( -\frac{1}{2\sigma^2} (\vec{x}_j - \vec{r}_i^{(\text{old})})^\top (\vec{x}_j - \vec{r}_i^{(\text{old})}) \right)}{\sum_{\vec{r}\in Q(c_j)} \exp \left( -\frac{1}{2\sigma^2} (\vec{x}_j - \vec{r}^{(\text{old})})^\top (\vec{x}_j - \vec{r}^{(\text{old})}) \right)}.
\]

\(R(c)\) is the set of reference vectors assigned to class \(c\) and \(Q(c)\) its complement.
Hard Learning Vector Quantization

**Idea:** Derive a scheme with hard assignments from the soft version.

**Approach:** Let the size parameter $\sigma$ of the Gaussian function go to zero.

The resulting update rule is in this case:

\[
\vec{r}_i^{\text{(new)}} = \vec{r}_i^{\text{(old)}} + \eta \cdot \begin{cases} 
  u_{ij}^{\oplus} \cdot (\vec{x}_j - \vec{r}_i^{\text{(old)}}), & \text{if } c_j = z_i, \\
  -u_{ij}^{\ominus} \cdot (\vec{x}_j - \vec{r}_i^{\text{(old)}}), & \text{if } c_j \neq z_i,
\end{cases}
\]

where

\[
u_{ij}^{\oplus} = \begin{cases} 
  1, & \text{if } \vec{r}_i = \arg\min_{\vec{r} \in R(c_j)} d(\vec{x}_j, \vec{r}), \\
  0, & \text{otherwise},
\end{cases} \quad \nu_{ij}^{\ominus} = \begin{cases} 
  1, & \text{if } \vec{r}_i = \arg\min_{\vec{r} \in Q(c_j)} d(\vec{x}_j, \vec{r}), \\
  0, & \text{otherwise}.
\end{cases}
\]

$\vec{r}_i$ is closest vector of same class $\vec{r}_i$ is closest vector of different class

This update rule is stable without a *window rule* restricting the update.
Learning Vector Quantization: Extensions

- **Frequency Sensitive Competitive Learning**
  - The distance to a reference vector is modified according to the number of data points that are assigned to this reference vector.

- **Fuzzy Learning Vector Quantization**
  - Exploits the close relationship to fuzzy clustering.
  - Can be seen as an online version of fuzzy clustering.
  - Leads to faster clustering.

- **Size and Shape Parameters**
  - Associate each reference vector with a cluster radius. Update this radius depending on how close the data points are.
  - Associate each reference vector with a covariance matrix. Update this matrix depending on the distribution of the data points.
Demonstration of learning vector quantization:

- Visualization of the training process
- Arbitrary datasets, but training only in two dimensions
- [http://www.borgelt.net/lvqd.html](http://www.borgelt.net/lvqd.html)
Self-Organizing Maps
A self-organizing map or Kohonen feature map is a neural network with a graph \( G = (U, C) \) that satisfies the following conditions

(i) \( U_{\text{hidden}} = \emptyset, U_{\text{in}} \cap U_{\text{out}} = \emptyset \),

(ii) \( C = U_{\text{in}} \times U_{\text{out}} \).

The network input function of each output neuron is a distance function of input and weight vector. The activation function of each output neuron is a radial function, that is, a monotonically decreasing function

\[
f : \mathbb{R}^+_0 \to [0, 1] \quad \text{with} \quad f(0) = 1 \quad \text{and} \quad \lim_{x \to \infty} f(x) = 0.
\]

The output function of each output neuron is the identity. The output is often discretized according to the "winner takes all" principle. On the output neurons a neighborhood relationship is defined:

\[
d_{\text{neurons}} : U_{\text{out}} \times U_{\text{out}} \to \mathbb{R}^+_0.
\]
Self-Organizing Maps: Neighborhood

Neighborhood of the output neurons: neurons form a grid

- Thin black lines: Indicate nearest neighbors of a neuron.
- Thick gray lines: Indicate regions assigned to a neuron for visualization.
- Usually two-dimensional grids are used to be able to draw the map easily.
Topology Preserving Mapping

Images of points close to each other in the original space should be close to each other in the image space.

Example: **Robinson projection** of the surface of a sphere (maps from 3 dimensions to 2 dimensions)

- Robinson projection is/was frequently used for world maps.
- The topology is preserved, although distances, angles, areas may be distorted.
Self-Organizing Maps: Topology Preserving Mapping

**neuron space/grid**
usually 2-dimensional
quadratic or
hexagonal grid

**input/data space**
usually high-dim.
(here: only 3-dim.)
blue: ref. vectors

Connections may be drawn between vectors corresponding to adjacent neurons.
Self-Organizing Maps: Neighborhood

Find topology preserving mapping by respecting the neighborhood

Reference vector update rule:
\[
\vec{r}_u^{(\text{new})} = \vec{r}_u^{(\text{old})} + \eta(t) \cdot f_{\text{nb}}(d_{\text{neurons}}(u, u_*), \varrho(t)) \cdot (\vec{x} - \vec{r}_u^{(\text{old})}),
\]

- \(u_*\) is the winner neuron (reference vector closest to data point).
- The neighborhood function \(f_{\text{nb}}\) is a radial function.

Time dependent learning rate
\[
\eta(t) = \eta_0 \alpha_\eta^t, \quad 0 < \alpha_\eta < 1, \quad \text{or} \quad \eta(t) = \eta_0 t^{\kappa_\eta}, \quad \kappa_\eta < 0.
\]

Time dependent neighborhood radius
\[
\varrho(t) = \varrho_0 \alpha_\varrho^t, \quad 0 < \alpha_\varrho < 1, \quad \text{or} \quad \varrho(t) = \varrho_0 t^{\kappa_\varrho}, \quad \kappa_\varrho < 0.
\]
The neighborhood size is reduced over time: (here: step function)

Note that a neighborhood function that is not a step function has a “soft” border and thus allows for a smooth reduction of the neighborhood size (larger changes of the reference vectors are restricted more and more to the close neighborhood).
Self-Organizing Maps: Training Procedure

- Initialize the weight vectors of the neurons of the self-organizing map, that is, place initial reference vectors in the input/data space.

- This may be done by randomly selecting training examples (provided there are fewer neurons than training examples, which is the usual case) or by sampling from some probability distribution on the data space.

- For the actual training, repeat the following steps:
  - Choose a training sample / data point (traverse the data points, possibly shuffling after each epoch).
  - Find the winner neuron with the distance function in the data space, that is, find the neuron with the closest reference vector.
  - Compute the time dependent radius and learning rate and adapt the corresponding neighbors of the winner neuron (severity of weight changes depend by neighborhood and learning rate).
Example: **Unfolding of a two-dimensional self-organizing map.**

- Self-organizing map with $10 \times 10$ neurons (quadratic grid) that is trained with random points chosen uniformly from the square $[-1, 1] \times [-1, 1]$.

- Initialization with random reference vectors chosen uniformly from $[-0.5, 0.5] \times [-0.5, 0.5]$.

- Gaussian neighborhood function

$$f_{nb}(d_{\text{neurons}}(u, u_*), \varrho(t)) = \exp \left( -\frac{d_{\text{neurons}}^2(u, u_*)}{2\varrho(t)^2} \right).$$

- Time-dependent neighborhood radius $\varrho(t) = 2.5 \cdot t^{-0.1}$

- Time-dependent learning rate $\eta(t) = 0.6 \cdot t$.

- The next slides show the SOM state after 10, 20, 40, 80 and 160 training steps. In each training step one training sample is processed. Shading of the neuron grid shows neuron activations for $(-0.5, -0.5)$. 
Unfolding of a two-dimensional self-organizing map. (data space)
Self-Organizing Maps: Examples

Unfolding of a two-dimensional self-organizing map. (neuron grid)
Example: Unfolding of a two-dimensional self-organizing map.

Training a self-organizing map may fail if

- the (initial) learning rate is chosen too small or
- or the (initial) neighborhood radius is chosen too small.
**Example:** Unfolding of a two-dimensional self-organizing map.

![Images of unfolded maps](image)

Self-organizing maps that have been trained with random points from (a) a rotation parabola, (b) a simple cubic function, (c) the surface of a sphere. Since the data points come from a two-dimensional subspace, training works well.

- In this case original space and image space have different dimensionality. (In the previous example they were both two-dimensional.)

- Self-organizing maps can be used for dimensionality reduction (in a quantized fashion, but interpolation may be used for smoothing).
Demonstration of self-organizing map training:

- Visualization of the training process
- Two-dimensional areas and three-dimensional surfaces
- http://www.borgelt.net/somd.html
Create a Phonotopic Map of Finnish [Kohonen 1988]

- The recorded microphone signal is converted into a spectral representation grouped into 15 channels.
- The 15-dimensional input space is mapped to a hexagonal SOM grid.

pictures not available in online version
Organize Countries based on Poverty Indicators [Kaski et al. 1997]

- 39 indicators describing various quality-of-life factors were used, such as state of health, nutrition, educational services etc.

picture not available in online version
Organize Countries based on Poverty Indicators [Kaski et al. 1997]

- Map of the world with countries colored by their poverty type.
- Countries in gray were not considered (possibly insufficient data).

picture not available in online version
Classify News Messages on Neural Networks [Kaski et al. 1998]

picture not available in online version
Organize Music Collections (Sound and Tags) [Mörchen et al. 2005/2007]

- Uses an Emergent Self-organizing Map to organize songs (no fixed shape, larger number of neurons than data points).
- Creates semantic features with regression and feature selection from 140 different short-term features (short time windows with stationary sound) 284 long term features (e.g. temporal statistics etc.) and user-assigned tags.

pictures not available in online version
Hopfield Networks and Boltzmann Machines
A **Hopfield network** is a neural network with a graph $G = (U, C)$ that satisfies the following conditions:

(i) $U_{\text{hidden}} = \emptyset$, $U_{\text{in}} = U_{\text{out}} = U$,

(ii) $C = U \times U - \{(u, u) \mid u \in U\}$.

- In a Hopfield network all neurons are input as well as output neurons.
- There are no hidden neurons.
- Each neuron receives input from all other neurons.
- A neuron is not connected to itself.

The connection weights are symmetric, that is,

$$\forall u, v \in U, u \neq v : \quad w_{uv} = w_{vu}.$$
Hopfield Networks

The network input function of each neuron is the weighted sum of the outputs of all other neurons, that is,

$$\forall u \in U : \quad f_{\text{net}}^{(u)}(\vec{w}_u, \vec{\text{in}}_u) = \vec{w}_u^\top \vec{\text{in}}_u = \sum_{v \in U - \{u\}} w_{uv} \text{out}_v.$$  

The activation function of each neuron is a threshold function, that is,

$$\forall u \in U : \quad f_{\text{act}}^{(u)}(\text{net}_u, \theta_u) = \begin{cases} 1, & \text{if } \text{net}_u \geq \theta, \\ -1, & \text{otherwise}. \end{cases}$$  

The output function of each neuron is the identity, that is,

$$\forall u \in U : \quad f_{\text{out}}^{(u)}(\text{act}_u) = \text{act}_u.$$
Alternative activation function

\[ \forall u \in U : \quad f^{(u)}_{\text{act}}(\text{net}_u, \theta_u, \text{act}_u) = \begin{cases} 
1, & \text{if } \text{net}_u > \theta, \\
-1, & \text{if } \text{net}_u < \theta, \\
\text{act}_u, & \text{if } \text{net}_u = \theta. 
\end{cases} \]

This activation function has advantages w.r.t. the physical interpretation of a Hopfield network.

General weight matrix of a Hopfield network

\[
W = \begin{pmatrix}
0 & w_{u_1u_2} & \ldots & w_{u_1u_n} \\
w_{u_1u_2} & 0 & \ldots & w_{u_2u_n} \\
\vdots & \vdots & \ddots & \vdots \\
w_{u_1u_n} & w_{u_1u_n} & \ldots & 0
\end{pmatrix}
\]
Hopfield Networks: Examples

Very simple Hopfield network

\[
\begin{align*}
&v_1 \\
v_2 \\
\end{align*}
\]

\[
\mathbf{W} = \begin{pmatrix}
0 & 1 \\
1 & 0
\end{pmatrix}
\]

The behavior of a Hopfield network can depend on the update order.

- Computations can oscillate if neurons are updated in parallel.
- Computations always converge if neurons are updated sequentially.
Parallel update of neuron activations

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<th>$u_1$</th>
<th>$u_2$</th>
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<tbody>
<tr>
<td>input</td>
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<td>1</td>
</tr>
<tr>
<td>work</td>
<td>1</td>
<td>−1</td>
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<tr>
<td>−1</td>
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<td>−1</td>
<td>1</td>
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- The computations oscillate, no stable state is reached.
- Output depends on when the computations are terminated.
Sequential update of neuron activations

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<th>$u_2$</th>
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</thead>
<tbody>
<tr>
<td>input phase</td>
<td>$-1$</td>
<td>$1$</td>
</tr>
<tr>
<td>work phase</td>
<td>$1$</td>
<td>$1$</td>
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<td>$-1$</td>
<td>$-1$</td>
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</tbody>
</table>

- Update order $u_1, u_2, u_1, u_2, \ldots$ (left) or $u_2, u_1, u_2, u_1, \ldots$ (right)
- Regardless of the update order a stable state is reached.
- However, *which* state is reached depends on the update order.
Simplified representation of a Hopfield network

\[
W = \begin{pmatrix}
0 & 1 & 2 \\
1 & 0 & 1 \\
2 & 1 & 0
\end{pmatrix}
\]

- Symmetric connections between neurons are combined.
- Inputs and outputs are not explicitly represented.
Graph of activation states and transitions:
(for the Hopfield network shown on the preceding slide)

“+”/”−” encode the neuron activations: “+” means +1 and “−” means −1.

Labels on arrows indicate the neurons, whose updates (activation changes) lead to the corresponding state transitions.

States shown in gray:
stable states, cannot be left again

States shown in white:
unstable states, may be left again.

Such a state graph captures all imaginable update orders.
Convergence Theorem: If the activations of the neurons of a Hopfield network are updated sequentially (asynchronously), then a stable state is reached in a finite number of steps.

If the neurons are traversed cyclically in an arbitrary, but fixed order, at most $n \cdot 2^n$ steps (updates of individual neurons) are needed, where $n$ is the number of neurons of the Hopfield network.

The proof is carried out with the help of an energy function. The energy function of a Hopfield network with $n$ neurons $u_1, \ldots, u_n$ is defined as

$$E = -\frac{1}{2} \vec{a}_{\text{ct}}^\top W \vec{a}_{\text{ct}} + \vec{\theta}^\top \vec{a}_{\text{ct}}$$

$$= -\frac{1}{2} \sum_{u,v \in U, u \neq v} w_{uv} \text{act}_u \text{act}_v + \sum_{u \in U} \theta_u \text{act}_u.$$
Hopfield Networks: Convergence

Consider the energy change resulting from an update that changes an activation:

$$\Delta E = E^{(\text{new})} - E^{(\text{old})} = \left( - \sum_{v \in U - \{u\}} w_{uv} \text{act}_{u}^{(\text{new})} \text{act}_{v} + \theta_{u} \text{act}_{u}^{(\text{new})} \right)$$

$$- \left( - \sum_{v \in U - \{u\}} w_{uv} \text{act}_{u}^{(\text{old})} \text{act}_{v} + \theta_{u} \text{act}_{u}^{(\text{old})} \right)$$

$$= \left( \text{act}_{u}^{(\text{old})} - \text{act}_{u}^{(\text{new})} \right) \left( \sum_{v \in U - \{u\}} w_{uv} \text{act}_{v} - \theta_{u} \right).$$

- $\text{net}_{u} < \theta_{u}$: Second factor is less than 0.  
  $\text{act}_{u}^{(\text{new})} = -1$ and $\text{act}_{u}^{(\text{old})} = 1$, therefore first factor greater than 0.  
  **Result:** $\Delta E < 0$.

- $\text{net}_{u} \geq \theta_{u}$: Second factor greater than or equal to 0.  
  $\text{act}_{u}^{(\text{new})} = 1$ and $\text{act}_{u}^{(\text{old})} = -1$, therefore first factor less than 0.  
  **Result:** $\Delta E \leq 0$.  

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It takes at most $n \cdot 2^n$ update steps to reach convergence.

- Provided that the neurons are updated in an arbitrary, but fixed order, since this guarantees that the neurons are traversed cyclically, and therefore each neuron is updated every $n$ steps.

- If in a traversal of all $n$ neurons no activation changes: a stable state has been reached.

- If in a traversal of all $n$ neurons at least one activation changes: the previous state cannot be reached again, because
  - either the new state has a smaller energy than the old (no way back: updates cannot increase the network energy)
  - or the number of +1 activations has increased (no way back: equal energy is possible only for $\text{net}_u \geq \theta_u$).

- The number of possible states of the Hopfield network is $2^n$, at least one of which must be rendered unreachable in each traversal of the $n$ neurons.
Hopfield Networks: Examples

Arrange states in state graph according to their energy

Energy function for example Hopfield network:

\[ E = -a_{1}a_{2} - 2a_{1}a_{3} - a_{2}a_{3} \]
The state graph need not be symmetric
Physical interpretation: Magnetism

A Hopfield network can be seen as a (microscopic) model of magnetism (so-called Ising model, [Ising 1925]).

<table>
<thead>
<tr>
<th>Physical</th>
<th>Neural</th>
</tr>
</thead>
<tbody>
<tr>
<td>atom</td>
<td>neuron</td>
</tr>
<tr>
<td>magnetic moment (spin)</td>
<td>activation state</td>
</tr>
<tr>
<td>strength of outer magnetic field</td>
<td>threshold value</td>
</tr>
<tr>
<td>magnetic coupling of the atoms</td>
<td>connection weights</td>
</tr>
<tr>
<td>Hamilton operator of the magnetic field</td>
<td>energy function</td>
</tr>
</tbody>
</table>
Hopfield Networks: Associative Memory

Idea: Use stable states to store patterns

First: Store only one pattern \( \vec{x} = (\text{act}_{u_1}^{(l)}, \ldots, \text{act}_{u_n}^{(l)})^\top \in \{-1, 1\}^n, n \geq 2 \), that is, find weights, so that pattern is a stable state.

Necessary and sufficient condition:

\[
S(\mathbf{W} \vec{x} - \vec{\theta}) = \vec{x},
\]

where

\[
S : \mathbb{R}^n \rightarrow \{-1, 1\}^n,
\]

\[
\vec{x} \mapsto \vec{y}
\]

with

\[
\forall i \in \{1, \ldots, n\} : \quad y_i = \begin{cases} 
1, & \text{if } x_i \geq 0, \\
-1, & \text{otherwise}.
\end{cases}
\]
If $\vec{\theta} = \vec{0}$ an appropriate matrix $\mathbf{W}$ can easily be found. It suffices

$$\mathbf{W}\vec{x} = c\vec{x} \quad \text{with } c \in \mathbb{R}^+.$$ 

Algebraically: Find a matrix $\mathbf{W}$ that has a positive eigenvalue w.r.t. $\vec{x}$.

Choose

$$\mathbf{W} = \vec{x}\vec{x}^\top - \mathbf{E}$$

where $\vec{x}\vec{x}^\top$ is the so-called outer product of $\vec{x}$ with itself.

With this matrix we have

$$\mathbf{W}\vec{x} = (\vec{x}\vec{x}^\top)\vec{x} - \mathbf{E}\vec{x} \overset{(*)}{=} \vec{x} \left(\vec{x}^\top \vec{x}\right) - \vec{x} = |\vec{x}|^2 = n$$

$$= n\vec{x} - \vec{x} = (n - 1)\vec{x}.$$ 

$(*)$ holds, because vector/matrix multiplication is associative.
Hopfield Networks: Associative Memory

Hebbian learning rule  [Hebb 1949]

Written in individual weights the computation of the weight matrix reads:

\[ w_{uv} = \begin{cases} 
0, & \text{if } u = v, \\
1, & \text{if } u \neq v, \text{ act}_u^{(p)} = \text{ act}_u^{(v)}, \\
-1, & \text{otherwise.} 
\end{cases} \]

- Originally derived from a biological analogy.
- Strengthen connection between neurons that are active at the same time.

Note that this learning rule also stores the complement of the pattern:

With \( \mathbf{Wx} = (n - 1)x \) it is also \( \mathbf{W}(-\mathbf{x}) = (n - 1)(-\mathbf{x}) \).
Storing several patterns

Choose

\[ W\vec{x}_j = \sum_{i=1}^{m} W_i \vec{x}_j = \left( \sum_{i=1}^{m} (\vec{x}_i \vec{x}_i^\top) \vec{x}_j \right) - m \mathbf{E}\vec{x}_j \]

\[ = \left( \sum_{i=1}^{m} \vec{x}_i (\vec{x}_i^\top \vec{x}_j) \right) - m\vec{x}_j \]

If the patterns are orthogonal, we have

\[ \vec{x}_i^\top \vec{x}_j = \begin{cases} 0, & \text{if } i \neq j, \\ n, & \text{if } i = j, \end{cases} \]

and therefore

\[ W\vec{x}_j = (n - m)\vec{x}_j. \]
Storing several patterns

\[ W \vec{x}_j = (n - m) \vec{x}_j. \]

Result: As long as \( m < n \), \( \vec{x} \) is a stable state of the Hopfield network.

Note that the complements of the patterns are also stored.

With \( W \vec{x}_j = (n - m) \vec{x}_j \) it is also \( W(-\vec{x}_j) = (n - m)(-\vec{x}_j). \)

But: Capacity is very small compared to the number of possible states \( (2^n) \), since at most \( m = n - 1 \) orthogonal patterns can be stored (so that \( n - m > 0 \)).

Furthermore, the requirement that the patterns must be orthogonal is a strong limitation of the usefulness of this result.
Non-orthogonal patterns:

\[ \mathbf{W \tilde{x}_j} = (n - m) \tilde{x}_j + \sum_{\substack{i=1 \\ i \neq j}}^{m} \tilde{x}_i (\tilde{x}_i^\top \tilde{x}_j) \]

“disturbance term”

- The “disturbance term” need not make it impossible to store the patterns.
- The states corresponding to the patterns \( \tilde{x}_j \) may still be stable, if the “disturbance term” is sufficiently small.
- For this term to be sufficiently small, the patterns must be “almost” orthogonal.
- The larger the number of patterns to be stored (that is, the smaller \( n - m \)), the smaller the “disturbance term” must be.
- The theoretically possible maximal capacity of a Hopfield network (that is, \( m = n - 1 \)) is hardly ever reached in practice.
Example: Store patterns $\vec{x}_1 = (+1, +1, -1, -1)^\top$ and $\vec{x}_2 = (-1, +1, -1, +1)^\top$.

$$W = W_1 + W_2 = \vec{x}_1\vec{x}_1^\top + \vec{x}_2\vec{x}_2^\top - 2E$$

where

$$W_1 = \begin{pmatrix} 
 0 & 1 & -1 & -1 \\
 1 & 0 & -1 & -1 \\
 -1 & -1 & 0 & 1 \\
 -1 & -1 & 1 & 0 \\
\end{pmatrix}, \quad W_2 = \begin{pmatrix} 
 0 & -1 & 1 & -1 \\
 -1 & 0 & -1 & 1 \\
 1 & -1 & 0 & -1 \\
 -1 & 1 & -1 & 0 \\
\end{pmatrix}.$$ 

The full weight matrix is:

$$W = \begin{pmatrix} 
 0 & 0 & 0 & -2 \\
 0 & 0 & -2 & 0 \\
 0 & -2 & 0 & 0 \\
 -2 & 0 & 0 & 0 \\
\end{pmatrix}.$$ 

Therefore it is

$$W\vec{x}_1 = (+2, +2, -2, -2)^\top \quad \text{and} \quad W\vec{x}_1 = (-2, +2, -2, +2)^\top.$$
Example: Storing bit maps of numbers

- Left: Bit maps stored in a Hopfield network.
- Right: Reconstruction of a pattern from a random input.
Hopfield Networks: Associative Memory

Training a Hopfield network with the Delta rule

Necessary condition for pattern $\vec{x}$ being a stable state:

$$s(0 + w_{u_1 u_2} \text{act}_{u_2}^{(p)} + \ldots + w_{u_1 u_n} \text{act}_{u_n}^{(p)} - \theta_{u_1}) = \text{act}_{u_1}^{(p)},$$

$$s(w_{u_2 u_1} \text{act}_{u_1}^{(p)} + 0 + \ldots + w_{u_2 u_n} \text{act}_{u_n}^{(p)} - \theta_{u_2}) = \text{act}_{u_2}^{(p)},$$

$$\vdots$$

$$s(w_{u_n u_1} \text{act}_{u_1}^{(p)} + w_{u_n u_2} \text{act}_{u_2}^{(p)} + \ldots + 0 - \theta_{u_n}) = \text{act}_{u_n}^{(p)}.$$

with the standard threshold function

$$s(x) = \begin{cases} 
  1, & \text{if } x \geq 0, \\
  -1, & \text{otherwise.}
\end{cases}$$
Training a Hopfield network with the Delta rule

Turn weight matrix into a weight vector:

\[ \vec{w} = ( w_{u_1u_2}, w_{u_1u_3}, \ldots, w_{u_1u_n}, \\
    w_{u_2u_3}, \ldots, w_{u_2u_n}, \\
    \ldots, \\
    w_{u_{n-1}u_n}, \\
    -\theta_{u_1}, -\theta_{u_2}, \ldots, -\theta_{u_n} ) . \]

Construct input vectors for a threshold logic unit

\[ \vec{z}_2 = ( \text{act}^{(p)}_{u_1}, 0, \ldots, 0, \text{act}^{(p)}_{u_3}, \ldots, \text{act}^{(p)}_{u_n}, 0, 1, 0, \ldots, 0 ) . \]

Apply Delta rule training / Widrow–Hoff procedure until convergence.
Demonstration of Hopfield networks as associative memory:

- Visualization of the association/recognition process
- Two-dimensional networks of arbitrary size
- http://www.borgelt.net/hfnd.html
Hopfield Networks: Solving Optimization Problems

Use energy minimization to solve optimization problems

General procedure:

- Transform function to optimize into a function to minimize.
- Transform function into the form of an energy function of a Hopfield network.
- Read the weights and threshold values from the energy function.
- Construct the corresponding Hopfield network.
- Initialize Hopfield network randomly and update until convergence.
- Read solution from the stable state reached.
- Repeat several times and use best solution found.
A Hopfield network may be defined either with activations $-1$ and $1$ or with activations $0$ and $1$. The networks can be transformed into each other.

From $\text{act}_u \in \{-1, 1\}$ to $\text{act}_u \in \{0, 1\}$:

$$
\begin{align*}
    w^0_{uv} &= 2w^-_{uv} \\
    \theta^0_u &= \theta^-_u + \sum_{v \in U \setminus \{u\}} w^-_{uv}
\end{align*}
$$

From $\text{act}_u \in \{0, 1\}$ to $\text{act}_u \in \{-1, 1\}$:

$$
\begin{align*}
    w^-_{uv} &= \frac{1}{2}w^0_{uv} \\
    \theta^-_u &= \theta^0_u - \frac{1}{2} \sum_{v \in U \setminus \{u\}} w^0_{uv}.
\end{align*}
$$
Combination lemma: Let two Hopfield networks on the same set $U$ of neurons with weights $w_{uv}^{(i)}$, threshold values $\theta_u^{(i)}$ and energy functions

$$E_i = -\frac{1}{2} \sum_{u \in U} \sum_{v \in U - \{u\}} w_{uv}^{(i)} \text{act}_u \text{act}_v + \sum_{u \in U} \theta_u^{(i)} \text{act}_u,$$

$i = 1, 2$, be given. Furthermore let $a, b \in \mathbb{R}$. Then $E = aE_1 + bE_2$ is the energy function of the Hopfield network on the neurons in $U$ that has the weights $w_{uv} = aw_{uv}^{(1)} + bw_{uv}^{(2)}$ and the threshold values $\theta_u = a\theta_u^{(1)} + b\theta_u^{(2)}$.

Proof: Just do the computations.

Idea: Additional conditions can be formalized separately and incorporated later. (One energy function per condition, then apply combination lemma.)
Example: Traveling salesman problem

Idea: Represent tour by a matrix.

An element $m_{ij}$ of the matrix is 1 if the $i$-th city is visited in the $j$-th step and 0 otherwise.

Each matrix element will be represented by a neuron.
Minimization of the tour length

\[ E_1 = \sum_{j_1=1}^{n} \sum_{j_2=1}^{n} \sum_{i=1}^{n} d_{j_1 j_2} \cdot m_{ij_1} \cdot m_{(i \mod n)+1, j_2}. \]

Double summation over steps (index \( i \)) needed:

\[ E_1 = \sum_{(i_1, j_1)\in\{1,\ldots,n\}^2} \sum_{(i_2, j_2)\in\{1,\ldots,n\}^2} d_{j_1 j_2} \cdot \delta(i_1 \mod n)+1, i_2 \cdot m_{i_1 j_1} \cdot m_{i_2 j_2}, \]

where

\[ \delta_{ab} = \begin{cases} 1, & \text{if } a = b, \\ 0, & \text{otherwise}. \end{cases} \]

Symmetric version of the energy function:

\[ E_1 = -\frac{1}{2} \sum_{(i_1, j_1)\in\{1,\ldots,n\}^2} \sum_{(i_2, j_2)\in\{1,\ldots,n\}^2} -d_{j_1 j_2} \cdot (\delta(i_1 \mod n)+1, i_2 + \delta_{i_1, (i_2 \mod n)+1}) \cdot m_{i_1 j_1} \cdot m_{i_2 j_2}. \]
Hopfield Networks: Solving Optimization Problems

Additional conditions that have to be satisfied:

- Each city is visited on exactly one step of the tour:
  \[ \forall j \in \{1, \ldots, n\} : \sum_{i=1}^{n} m_{ij} = 1, \]
  that is, each column of the matrix contains exactly one 1.

- On each step of the tour exactly one city is visited:
  \[ \forall i \in \{1, \ldots, n\} : \sum_{j=1}^{n} m_{ij} = 1, \]
  that is, each row of the matrix contains exactly one 1.

These conditions are incorporated by finding additional functions to optimize.
Hopfield Networks: Solving Optimization Problems

Formalization of first condition as a minimization problem:

\[
E_2^* = \sum_{j=1}^{n} \left( \left( \sum_{i=1}^{n} m_{ij} \right)^2 - 2 \sum_{i=1}^{n} m_{ij} + 1 \right)
\]

\[
= \sum_{j=1}^{n} \left( \left( \sum_{i_1=1}^{n} m_{i_1j} \right) \left( \sum_{i_2=1}^{n} m_{i_2j} \right) - 2 \sum_{i=1}^{n} m_{ij} + 1 \right)
\]

\[
= \sum_{j=1}^{n} \sum_{i_1=1}^{n} \sum_{i_2=1}^{n} m_{i_1j} m_{i_2j} - 2 \sum_{j=1}^{n} \sum_{i=1}^{n} m_{ij} + n.
\]

Double summation over cities (index \(i\)) needed:

\[
E_2 = \sum_{(i_1,j_1)\in\{1,...,n\}^2} \sum_{(i_2,j_2)\in\{1,...,n\}^2} \delta_{j_1 j_2} \cdot m_{i_1 j_1} \cdot m_{i_2 j_2} - 2 \sum_{(i,j)\in\{1,...,n\}^2} m_{ij}.
\]
Resulting energy function:

\[ E_2 = -\frac{1}{2} \sum_{(i_1,j_1)\in\{1,\ldots,n\}^2} -2\delta_{j_1 j_2} \cdot m_{i_1 j_1} \cdot m_{i_2 j_2} + \sum_{(i,j)\in\{1,\ldots,n\}^2} -2m_{ij} \]

Second additional condition is handled in a completely analogous way:

\[ E_3 = -\frac{1}{2} \sum_{(i_1,j_1)\in\{1,\ldots,n\}^2} -2\delta_{i_1 i_2} \cdot m_{i_1 j_1} \cdot m_{i_2 j_2} + \sum_{(i,j)\in\{1,\ldots,n\}^2} -2m_{ij}. \]

Combining the energy functions:

\[ E = aE_1 + bE_2 + cE_3 \quad \text{where} \quad \frac{b}{a} = \frac{c}{a} > 2 \max_{(j_1,j_2)\in\{1,\ldots,n\}^2} d_{j_1 j_2}. \]
From the resulting energy function we can read the weights

\[ w(i_1,j_1)(i_2,j_2) = -ad_{j_1j_2} \cdot (\delta(i_1 \text{mod } n)+1,i_2 + \delta(i_1, (i_2 \text{mod } n)+1) - 2b\delta_{j_1j_2} - 2c\delta_{i_1i_2} \]

from \( E_1 \)

from \( E_2 \) from \( E_3 \)

and the threshold values:

\[ \theta(i,j) = 0a \quad -2b \quad -2c = -2(b + c). \]

from \( E_1 \) from \( E_2 \) from \( E_3 \)

Problem: Random initialization and update until convergence
not always leads to a matrix that represents a tour, let alone an optimal one.
Hopfield Networks: Reasons for Failure

Hopfield network only rarely finds a tour, let alone an optimal one.

- One of the main problems is that the Hopfield network is unable to switch from a found tour to another with a lower total length.

- The reason is that transforming a matrix that represents a tour into another matrix that represents a different tour requires that at least four neurons (matrix elements) change their activations.

- However, each of these changes, if carried out individually, violates at least one of the constraints and thus increases the energy.

- Only all four changes together can result in a smaller energy, but cannot be executed together due to the asynchronous update.

- Therefore the normal activation updates can never change an already found tour into another, even if this requires only a marginal change of the tour.
Results can be somewhat improved if instead of **discrete Hopfield networks** (activations in $\{-1, 1\}$ (or $\{0, 1\}$)) one uses **continuous Hopfield networks** (activations in $[-1, 1]$ (or $[0, 1]$)). However, the fundamental problem is not solved in this way.

More generally, the reason for the difficulties that are encountered if an optimization problem is to be solved with a Hopfield network is:

**The update procedure may get stuck in a local optimum.**

The problem of local optima occurs also with many other optimization methods, for example, gradient descent, hill climbing, alternating optimization etc.

Ideas to overcome this difficulty for other optimization methods may be transferred to Hopfield networks.

One such method, which is very popular, is **simulated annealing**.
Simulated Annealing

May be seen as an extension of random or gradient descent that tries to avoid getting stuck.

**Idea:** transitions from higher to lower (local) minima should be more probable than *vice versa.*

[Metropolis *et al.* 1953; Kirkpatrik *et al.* 1983]

**Principle of Simulated Annealing:**

- Random variants of the current solution (candidate) are created.
- Better solution (candidates) are always accepted.
- Worse solution (candidates) are accepted with a probability that depends on
  - the quality difference between the new and the old solution (candidate) and
  - a temperature parameter that is decreased with time.
Simulated Annealing

• **Motivation:**
  - Physical minimization of the energy (more precisely: atom lattice energy) if a heated piece of metal is cooled slowly.
  - This process is called **annealing**.
  - It serves the purpose to make the metal easier to work or to machine by relieving tensions and correcting lattice malformations.

• **Alternative Motivation:**
  - A ball rolls around on an (irregularly) curved surface; minimization of the potential energy of the ball.
  - In the beginning the ball is endowed with a certain kinetic energy, which enables it to roll up some slopes of the surface.
  - In the course of time, friction reduces the kinetic energy of the ball, so that it finally comes to a rest in a valley of the surface.

• **Attention:** There is no guarantee that the global optimum is found!
Simulated Annealing: Procedure

1. Choose a (random) starting point \( s_0 \in \Omega \) (\( \Omega \) is the search space).

2. Choose a point \( s' \in \Omega \) “in the vicinity” of \( s_i \)
   (for example, by a small random variation of \( s_i \)).

3. Set

   \[
   s_{i+1} = \begin{cases} 
   s' & \text{if } f(s') \geq f(s_i), \\
   s' & \text{with probability } p = e^{-\frac{\Delta f}{kT}} \text{ and} \\
   s_i & \text{with probability } 1 - p \text{ otherwise.}
   \end{cases}
   \]

   \( \Delta f = f(s_i) - f(s') \) quality difference of the solution (candidates)

   \( k = \Delta f_{\text{max}} \) (estimation of the) range of quality values

   \( T \) temperature parameter (is (slowly) decreased over time)

4. Repeat steps 2 and 3 until some termination criterion is fulfilled.

   • For (very) small \( T \) the method approaches a pure random descent.
Applying simulated annealing to Hopfield networks is very simple:

- All neuron activations are initialized randomly.
- The neurons of the Hopfield network are traversed repeatedly (for example, in some random order).
- For each neuron, it is determined whether an activation change leads to a reduction of the network energy or not.
- An activation change that reduces the network energy is always accepted (in the normal update process, only such changes occur).
- However, if an activation change increases the network energy, it is accepted with a certain probability (see preceding slide).
- Note that in this case we have simply

\[ \Delta f = \Delta E = |\text{net}_u - \theta_u| \]
Hopfield Networks: Summary

- Hopfield networks are **restricted recurrent neural networks** (full pairwise connections, symmetric connection weights).

- Synchronous update of the neurons may lead to oscillations, but **asynchronous update is guaranteed to reach a stable state** (asynchronous updates either reduce the energy of the network or increase the number of +1 activations).

- Hopfield networks can be used as **associative memory**, that is, as memory that is addressed by its contents, by using the stable states to store desired patterns.

- Hopfield networks can be used to **solve optimization problems**, if the function to optimize can be reformulated as an energy function (stable states are (local) minima of the energy function).

- Approaches like **simulated annealing** may be needed to prevent that the update gets stuck in a local optimum.
Boltzmann Machines

- Boltzmann machines are closely related to Hopfield networks.
- They differ from Hopfield networks mainly in how the neurons are updated.
- They also rely on the fact that one can define an energy function that assigns a numeric value (an energy) to each state of the network.
- With the help of this energy function a probability distribution over the states of the network is defined based on the Boltzmann distribution (also known as Gibbs distribution) of statistical mechanics, namely

$$P(\vec{s}) = \frac{1}{c} e^{-\frac{E(\vec{s})}{kT}}.$$ 

$\vec{s}$ describes the (discrete) state of the system,
$c$ is a normalization constant,
$E$ is the function that yields the energy of a state $\vec{s}$,
$T$ is the thermodynamic temperature of the system,
$k$ is Boltzmann’s constant ($k \approx 1.38 \cdot 10^{-23} \text{ J/K}$).
Boltzmann Machines

- For Boltzmann machines the product $kT$ is often replaced by merely $T$, combining the temperature and Boltzmann’s constant into a single parameter.
- The state $\vec{s}$ consists of the vector $\vec{act}$ of the neuron activations.
- The energy function of a Boltzmann machine is
  \[ E(\vec{act}) = -\frac{1}{2} \vec{act}^\top \mathbf{W} \vec{act} + \vec{\theta}^\top \vec{act}, \]
  where $\mathbf{W}$: matrix of connection weights; $\vec{\theta}$: vector of threshold values.
- Consider the energy change resulting from the change of a single neuron $u$:
  \[ \Delta E_u = E_{act_u=1} - E_{act_u=0} = \sum_{v \in U - \{u\}} w_{uv} \text{act}_v - \theta_u \]
- Writing the energies in terms of the Boltzmann distribution yields
  \[ \Delta E_u = -kT \ln(P(\text{act}_u = 1)) - (-kT \ln(P(\text{act}_u = 0))). \]
Boltzmann Machines

- Rewrite as
  \[ \frac{\Delta E_u}{kT} = \ln(P(\text{act}_u = 1)) - \ln(P(\text{act}_u = 0)) \]
  \[ = \ln(P(\text{act}_u = 1)) - \ln(1 - P(\text{act}_u = 1)) \]
  (since obviously \( P(\text{act}_u = 0) + P(\text{act}_u = 1) = 1 \)).

- Solving this equation for \( P(\text{act}_u = 1) \) finally yields
  \[ P(\text{act}_u = 1) = \frac{1}{1 + e^{-\frac{\Delta E_u}{kT}}} . \]

- That is: the probability of a neuron being active is a logistic function of the (scaled) energy difference between its active and inactive state.

- Since the energy difference is closely related to the network input, namely as
  \[ \Delta E_u = \sum_{v \in U - \{u\}} w_{uv} \text{act}_v - \theta_u = \text{net}_u - \theta_u, \]
  this formula suggests a stochastic update procedure for the network.
Boltzmann Machines: Update Procedure

- A neuron $u$ is chosen (randomly), its network input, from it the energy difference $\Delta E_u$ and finally the probability of the neuron having activation 1 is computed. The neuron is set to activation 1 with this probability and to 0 otherwise.

- This update is repeated many times for randomly chosen neurons.

- Simulated annealing is carried out by slowly lowering the temperature $T$.

- This update process is a Markov Chain Monte Carlo (MCMC) procedure.

- After sufficiently many steps, the probability that the network is in a specific activation state depends only on the energy of that state. It is independent of the initial activation state the process was started with.

- This final situation is also referred to as thermal equilibrium.

- Therefore: Boltzmann machines are representations of and sampling mechanisms for the Boltzmann distributions defined by their weights and threshold values.
Boltzmann Machines: Training

- **Idea of Training:** Develop a training procedure with which the probability distribution represented by a Boltzmann machine via its energy function can be adapted to a given sample of data points, in order to obtain a **probabilistic model of the data**.

- This objective can only be achieved sufficiently well if the data points are actually a sample from a Boltzmann distribution. (Otherwise the model cannot, in principle, be made to fit the sample data well.)

- In order to mitigate this restriction to Boltzmann distributions, a deviation from the structure of Hopfield networks is introduced.

- The neurons are divided into
  - **visible neurons**, which receive the data points as input, and
  - **hidden neurons**, which are not fixed by the data points.

  (Reminder: Hopfield networks have only visible neurons.)
Boltzmann Machines: Training

- **Objective of Training:**
  Adapt the connection weights and threshold values in such a way that the true distribution underlying a given data sample is approximated well by the probability distribution represented by the Boltzmann machine on its visible neurons.

- **Natural Approach to Training:**
  - Choose a measure for the difference between two probability distributions.
  - Carry out a gradient descent in order to minimize this difference measure.

- **Well-known measure:** **Kullback–Leibler information divergence.**
  For two probability distributions $p_1$ and $p_2$ defined over the same sample space $\Omega$:
  \[
  KL(p_1, p_2) = \sum_{\omega \in \Omega} p_1(\omega) \ln \frac{p_1(\omega)}{p_2(\omega)}.
  \]
  Applied to Boltzmann machines: $p_1$ refers to the data sample, $p_2$ to the visible neurons of the Boltzmann machine.
Boltzmann Machines: Training

- In each training step the Boltzmann machine is ran twice (two “phases”).
- “Positive Phase”: Visible neurons are fixed to a randomly chosen data point; only the hidden neurons are updated until thermal equilibrium is reached.
- “Negative Phase”: All units are updated until thermal equilibrium is reached.
- In the two phases statistics about individual neurons and pairs of neurons (both visible and hidden) being activated (simultaneously) are collected.
- Then update is performed according to the following two equations:

  \[ \Delta w_{uv} = \frac{1}{\eta} (p^+_{uv} - p^-_{uv}) \quad \text{and} \quad \Delta \theta_u = -\frac{1}{\eta} (p^+_u - p^-_u). \]

  - \( p_u \) probability that neuron \( u \) is active,
  - \( p_{uv} \) probability that neurons \( u \) and \( v \) are both active simultaneously,
  - \(+\) \(-\) as upper indices indicate the phase referred to.

  (All probabilities are estimated from observed relative frequencies.)
Boltzmann Machines: Training

- Intuitive explanation of the update rule:
  - If a neuron is more often active when a data sample is presented than when the network is allowed to run freely, the probability of the neuron being active is too low, so the threshold should be reduced.
  - If neurons are more often active together when a data sample is presented than when the network is allowed to run freely, the connection weight between them should be increased, so that they become more likely to be active together.

- This training method is very similar to the Hebbian learning rule.
  Derived from a biological analogy it says: connections between two neurons that are synchronously active are strengthened ("cells that fire together, wire together").
Boltzmann Machines: Training

- Unfortunately, this procedure is impractical unless the networks are very small.
- The main reason is the fact that the larger the network, the more update steps need to be carried out in order to obtain sufficiently reliable statistics for the neuron activation (pairs) needed in the update formulas.
- Efficient training is possible for the restricted Boltzmann machine.
- Restriction consists in using a bipartite graph instead of a fully connected graph:
  - Vertices are split into two groups, the visible and the hidden neurons;
  - Connections only exist between neurons from different groups.
A **restricted Boltzmann Machine (RBM)** or **Harmonium** is a neural network with a graph $G = (U, C)$ that satisfies the following conditions:

(i) $U = U_{\text{in}} \cup U_{\text{hidden}}, \quad U_{\text{in}} \cap U_{\text{hidden}} = \emptyset, \quad U_{\text{out}} = U_{\text{in}},$

(ii) $C = U_{\text{in}} \times U_{\text{hidden}}.$

- In a restricted Boltzmann machine, all input neurons are also output neurons and *vice versa* (all output neurons are also input neurons).
- There are hidden neurons, which are different from input and output neurons.
- Each input/output neuron receives input from all hidden neurons; each hidden neuron receives input from all input/output neurons.

The connection weights between input and hidden neurons are symmetric, that is,

$$\forall u \in U_{\text{in}}, v \in U_{\text{hidden}} : \quad w_{uv} = w_{vu}.$$
Due to the lack of connections within the visible units and within the hidden units, training can proceed by repeating the following three steps:

- Visible units are fixed to a randomly chosen data sample $\vec{x}$; hidden units are updated once and in parallel (result: $\vec{y}$). $\vec{x}\vec{y}^\top$ is called the **positive gradient** for the weight matrix.

- Hidden neurons are fixed to the computed vector $\vec{y}$; visible units are updated once and in parallel (result: $\vec{x}^*$). Visible neurons are fixed to “reconstruction” $\vec{x}^*$; hidden neurons are update once more (result: $\vec{y}^*$). $\vec{x}^*\vec{y}^*\top$ is called the **negative gradient** for the weight matrix.

- Connection weights are updated with difference of positive and negative gradient:

$$\Delta w_{uv} = \eta(\vec{x}_u\vec{y}_v^\top - \vec{x}^*_u\vec{y}^*_v\top)$$

where $\eta$ is a learning rate.
• Many **improvements of this basic procedure** exist [Hinton 2010]:
  o use a momentum term for the training,
  o use actual probabilities instead of binary reconstructions,
  o use online-like training based on small batches etc.

• Restricted Boltzmann machines have also been used to build **deep networks**
  in a fashion similar to stacked auto-encoders for multi-layer perceptrons.

• Idea: train a restricted Boltzmann machine, then create a data set of hidden
  neuron activations by sampling from the trained Boltzmann machine, and build
  another restricted Boltzmann machine from the obtained data set.

• This procedure can be repeated several times and
  the resulting Boltzmann machines can then easily be stacked.

• The obtained stack is fine-tuned with a procedure similar to back-propagation
  [Hinton *et al.* 2006].
Recurrent Neural Networks
Recurrent Networks: Cooling Law

A body of temperature $\vartheta_0$ that is placed into an environment with temperature $\vartheta_A$. The cooling/heating of the body can be described by **Newton’s cooling law**:

$$\frac{d\vartheta}{dt} = \dot{\vartheta} = -k(\vartheta - \vartheta_A).$$

Exact analytical solution:

$$\vartheta(t) = \vartheta_A + (\vartheta_0 - \vartheta_A)e^{-k(t-t_0)}$$

Approximate solution with **Euler-Cauchy polygonal courses**:

$$\vartheta_1 = \vartheta(t_1) = \vartheta(t_0) + \dot{\vartheta}(t_0)\Delta t = \vartheta_0 - k(\vartheta_0 - \vartheta_A)\Delta t.$$  

$$\vartheta_2 = \vartheta(t_2) = \vartheta(t_1) + \dot{\vartheta}(t_1)\Delta t = \vartheta_1 - k(\vartheta_1 - \vartheta_A)\Delta t.$$  

General recursive formula:

$$\vartheta_i = \vartheta(t_i) = \vartheta(t_{i-1}) + \dot{\vartheta}(t_{i-1})\Delta t = \vartheta_{i-1} - k(\vartheta_{i-1} - \vartheta_A)\Delta t.$$
Recurrent Networks: Cooling Law

Euler–Cauchy polygonal courses for different step widths:

The thin curve is the exact analytical solution.

Recurrent neural network:
Recurrent Networks: Cooling Law

More formal derivation of the recursive formula:

Replace differential quotient by **forward difference**

\[
\frac{d\vartheta(t)}{dt} \approx \frac{\Delta \vartheta(t)}{\Delta t} = \frac{\vartheta(t + \Delta t) - \vartheta(t)}{\Delta t}
\]

with sufficiently small \(\Delta t\). Then it is

\[
\vartheta(t + \Delta t) - \vartheta(t) = \Delta \vartheta(t) \approx -k(\vartheta(t) - \vartheta_A)\Delta t,
\]

\[
\vartheta(t + \Delta t) - \vartheta(t) = \Delta \vartheta(t) \approx -k\Delta t\vartheta(t) + k\vartheta_A\Delta t
\]

and therefore

\[
\vartheta_i \approx \vartheta_{i-1} - k\Delta t\vartheta_{i-1} + k\vartheta_A\Delta t.
\]
Governing physical laws:

- **Hooke’s law**: $F = c\Delta l = -cx$ (c is a spring dependent constant)
- **Newton’s second law**: $F = ma = m\ddot{x}$ (force causes an acceleration)

Resulting differential equation:

$$m\ddot{x} = -cx \quad \text{or} \quad \ddot{x} = -\frac{c}{m}x.$$
Recurrent Networks: Mass on a Spring

General analytical solution of the differential equation:

\[ x(t) = a \sin(\omega t) + b \cos(\omega t) \]

with the parameters

\[ \omega = \sqrt{\frac{c}{m}}, \quad a = x(t_0) \sin(\omega t_0) + v(t_0) \cos(\omega t_0), \]
\[ b = x(t_0) \cos(\omega t_0) - v(t_0) \sin(\omega t_0). \]

With given initial values \( x(t_0) = x_0 \) and \( v(t_0) = 0 \) and the additional assumption \( t_0 = 0 \) we get the simple expression

\[ x(t) = x_0 \cos \left( \sqrt{\frac{c}{m}} t \right). \]
Recurrent Networks: Mass on a Spring

Turn differential equation into two coupled equations:

\[ \dot{x} = v \quad \text{and} \quad \dot{v} = -\frac{c}{m} x. \]

Approximate differential quotient by forward difference:

\[ \frac{\Delta x}{\Delta t} = \frac{x(t + \Delta t) - x(t)}{\Delta t} = v \quad \text{and} \quad \frac{\Delta v}{\Delta t} = \frac{v(t + \Delta t) - v(t)}{\Delta t} = -\frac{c}{m} x \]

Resulting recursive equations:

\[ x(t_i) = x(t_{i-1}) + \Delta x(t_{i-1}) = x(t_{i-1}) + \Delta t \cdot v(t_{i-1}) \quad \text{and} \]

\[ v(t_i) = v(t_{i-1}) + \Delta v(t_{i-1}) = v(t_{i-1}) - \frac{c}{m} \Delta t \cdot x(t_{i-1}). \]
Recurrent Networks: Mass on a Spring

Neuron $u_1$:  
\[ f^{(u_1)}_{\text{net}}(v, w_{u1}u_2) = w_{u1}u_2 v = -\frac{c}{m} \Delta t \ v \quad \text{and} \]
\[ f^{(u_1)}_{\text{act}}(\text{act}_{u_1}, \text{net}_{u_1}, \theta_{u_1}) = \text{act}_{u_1} + \text{net}_{u_1} - \theta_{u_1}, \]

Neuron $u_2$:  
\[ f^{(u_2)}_{\text{net}}(x, w_{u2}u_1) = w_{u2}u_1 x = \Delta t \ x \quad \text{and} \]
\[ f^{(u_2)}_{\text{act}}(\text{act}_{u_2}, \text{net}_{u_2}, \theta_{u_2}) = \text{act}_{u_2} + \text{net}_{u_2} - \theta_{u_2}. \]
Recurrent Networks: Mass on a Spring

Some computation steps of the neural network:

<table>
<thead>
<tr>
<th>$t$</th>
<th>$v$</th>
<th>$x$</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.0</td>
<td>0.0000</td>
<td>1.0000</td>
</tr>
<tr>
<td>0.1</td>
<td>−0.5000</td>
<td>0.9500</td>
</tr>
<tr>
<td>0.2</td>
<td>−0.9750</td>
<td>0.8525</td>
</tr>
<tr>
<td>0.3</td>
<td>−1.4012</td>
<td>0.7124</td>
</tr>
<tr>
<td>0.4</td>
<td>−1.7574</td>
<td>0.5366</td>
</tr>
<tr>
<td>0.5</td>
<td>−2.0258</td>
<td>0.3341</td>
</tr>
<tr>
<td>0.6</td>
<td>−2.1928</td>
<td>0.1148</td>
</tr>
</tbody>
</table>

- The resulting curve is close to the analytical solution.
- The approximation gets better with smaller step width.
General representation of explicit $n$-th order differential equation:

$$x^{(n)} = f(t, x, \dot{x}, \ddot{x}, \ldots, x^{(n-1)})$$

Introduce $n - 1$ intermediary quantities

$$y_1 = \dot{x}, \quad y_2 = \ddot{x}, \quad \ldots \quad y_{n-1} = x^{(n-1)}$$

to obtain the system

$$\begin{align*}
\dot{x} &= y_1, \\
\dot{y}_1 &= y_2, \\
\vdots \\
\dot{y}_{n-2} &= y_{n-1}, \\
\dot{y}_{n-1} &= f(t, x, y_1, y_2, \ldots, y_{n-1})
\end{align*}$$

of $n$ coupled first order differential equations.
Recurrent Networks: Differential Equations

Replace differential quotient by forward distance to obtain the recursive equations

\[ x(t_i) = x(t_{i-1}) + \Delta t \cdot y_1(t_{i-1}), \]
\[ y_1(t_i) = y_1(t_{i-1}) + \Delta t \cdot y_2(t_{i-1}), \]
\[ \vdots \]
\[ y_{n-2}(t_i) = y_{n-2}(t_{i-1}) + \Delta t \cdot y_{n-3}(t_{i-1}), \]
\[ y_{n-1}(t_i) = y_{n-1}(t_{i-1}) + f(t_{i-1}, x(t_{i-1}), y_1(t_{i-1}), \ldots, y_{n-1}(t_{i-1})) \]

- Each of these equations describes the update of one neuron.
- The last neuron needs a special activation function.
Recurrent Networks: Differential Equations

\[
\dot{x}_0, \quad \ddot{x}_0, \quad \dddot{x}_0, \quad \ldots, \quad x_0^{(n-1)} \quad \text{to} \quad \theta \quad \text{and} \quad -\Delta t
\]

\[
x_0, \quad \Delta t, \quad x(t)
\]
Recurrent Networks: Diagonal Throw

Diagonal throw of a body.

Two differential equations (one for each coordinate):

\[ \ddot{x} = 0 \quad \text{and} \quad \ddot{y} = -g, \]

where \( g = 9.81 \text{ ms}^{-2} \).

Initial conditions \( x(t_0) = x_0, \ y(t_0) = y_0, \ \dot{x}(t_0) = v_0 \cos \varphi \) and \( \dot{y}(t_0) = v_0 \sin \varphi \).
Introduce intermediary quantities

\[ v_x = \dot{x} \quad \text{and} \quad v_y = \dot{y} \]

to reach the system of differential equations:

\[ \dot{x} = v_x, \quad \dot{v}_x = 0, \]

\[ \dot{y} = v_y, \quad \dot{v}_y = -g, \]

from which we get the system of recursive update formulae

\[ x(t_i) = x(t_{i-1}) + \Delta t \ v_x(t_{i-1}), \quad v_x(t_i) = v_x(t_{i-1}), \]

\[ y(t_i) = y(t_{i-1}) + \Delta t \ v_y(t_{i-1}), \quad v_y(t_i) = v_y(t_{i-1}) - \Delta t \ g. \]
Better description: Use vectors as inputs and outputs

\[ \ddot{\vec{r}} = -g \vec{e}_y, \]

where \( \vec{e}_y = (0, 1) \).

Initial conditions are \( \vec{r}(t_0) = \vec{r}_0 = (x_0, y_0) \) and \( \dot{\vec{r}}(t_0) = \vec{v}_0 = (v_0 \cos \varphi, v_0 \sin \varphi) \).

Introduce one vector-valued intermediary quantity \( \vec{v} = \dot{\vec{r}} \) to obtain

\[ \dot{\vec{r}} = \vec{v}, \quad \dot{\vec{v}} = -g \vec{e}_y \]

This leads to the recursive update rules

\[ \vec{r}(t_i) = \vec{r}(t_{i-1}) + \Delta t \, \vec{v}(t_{i-1}), \]
\[ \vec{v}(t_i) = \vec{v}(t_{i-1}) - \Delta t \, g \vec{e}_y \]
Advantage of vector networks becomes obvious if friction is taken into account:

\[ \ddot{\vec{a}} = -\beta \vec{v} = -\beta \dot{\vec{r}} \]

\( \beta \) is a constant that depends on the size and the shape of the body.

This leads to the differential equation

\[ \ddot{\vec{r}} = -\beta \dot{\vec{r}} - g \vec{e}_y. \]

Introduce the intermediary quantity \( \vec{v} = \dot{\vec{r}} \) to obtain

\[ \dot{\vec{r}} = \vec{v}, \quad \dot{\vec{v}} = -\beta \vec{v} - g \vec{e}_y, \]

from which we obtain the recursive update formulae

\[ \vec{r}(t_i) = \vec{r}(t_{i-1}) + \Delta t \vec{v}(t_{i-1}), \]

\[ \vec{v}(t_i) = \vec{v}(t_{i-1}) - \Delta t \beta \vec{v}(t_{i-1}) - \Delta t g \vec{e}_y. \]
Recurrent Networks: Diagonal Throw

Resulting recurrent neural network:

- There are no strange couplings as there would be in a non-vector network.
- Note the deviation from a parabola that is due to the friction.
Recurrent Networks: Planet Orbit

\[ \ddot{\vec{r}} = -\gamma m \frac{\vec{r}}{|\vec{r}|^3}, \quad \Rightarrow \quad \dot{\vec{r}} = \vec{v}, \quad \dot{\vec{v}} = -\gamma m \frac{\vec{r}}{|\vec{r}|^3}. \]

Recursive update rules:

\[ \vec{r}(t_i) = \vec{r}(t_{i-1}) + \Delta t \vec{v}(t_{i-1}), \]

\[ \vec{v}(t_i) = \vec{v}(t_{i-1}) - \Delta t \gamma m \frac{\vec{r}(t_{i-1})}{|\vec{r}(t_{i-1})|^3}, \]
Recurrent Networks: Training

- All recurrent network computations shown up to now are possible only if the differential equation \textit{and} its parameters are known.
- In practice one often knows only the form of the differential equation.
- If measurement data of the described system are available, one may try to find the system parameters by training a recurrent neural network.
- In principle, recurrent neural networks are trained like multi-layer perceptrons, that is, an output error is computed and propagated back through the network.
- However, a direct application of error backpropagation is problematic due to the backward connections / recurrence.

\textbf{Solution:} The backward connections are eliminated by unfolding the network in time between two training patterns (\textit{error backpropagation through time}).

- Technically: create one (pseudo-)neuron for each intermediate point in time.
Recurrent Networks: Backpropagation through Time

Example: Newton’s cooling law

\[
\frac{d\vartheta}{dt} = \dot{\vartheta} = -k(\vartheta - \vartheta_A)
\]

- Assumption: we have measurements of the temperature of a body at different points in time. We also know the temperature \( \vartheta_A \) of the environment.

- Objective: find the value of the cooling constant \( k \).

- Initialization: like for an MLP, choose random thresholds and weights.

- The time between two consecutive measurement values is split into intervals. Thus the recurrence of the network is *unfolded in time*.

- For example, if there are 4 intervals between two consecutive measurement values \( t_{j+1} = t_j + 4\Delta t \), we obtain
Recurrent Networks: Backpropagation through Time

For a given measurement $\vartheta(t_j)$, an approximation of $\vartheta(t_{j+1})$ is computed.

By comparing this (approximate) output with the actual value $\vartheta(t_{j+1})$, an error signal is obtained that is backpropagated through the network and leads to changes of the thresholds and the connection weights.

Therefore: training is standard backpropagation on the unfolded network.

However: the above example network possesses only one free parameter, namely the cooling constant $k$, which is part of the weight and the threshold.

Therefore: updates can be carried out only after the first neuron is reached.

Generally, training recurrent networks is beneficial if the system of differential equations cannot be solved analytically.
Recurrent Networks: Virtual Laparoscopy

Example: Finding tissue parameters for virtual surgery/laparoscopy

Due to the large number of differential equations, such systems are much too complex to be treated analytically. However, by training recurrent neural networks, remarkable successes could be achieved.
Single Wind Turbine

- Input from sensors:
  - wind speed
  - vibration

- Variables:
  - blade angle
  - generator settings

- Objectives:
  - maximize electricity generation
  - minimize wear and tear
Many Wind Turbines

- Problem: wind turbines create turbulences, which affect other wind turbines placed behind them
- Input: sensor data of all wind turbines
- Variables: settings of all wind turbines
- Objectives:
  - maximize electricity generation
  - minimize wear and tear
- Solution: use a recurrent neural network
Supplementary Topic:
Neuro-Fuzzy Systems
Brief Introduction to Fuzzy Theory

• **Classical Logic:** only two truth values *true* and *false*
  
  **Classical Set Theory:** either *is element of* or *is not element of*

• The bivalence of the classical theories is often inappropriate.

Illustrative example: **Sorites Paradox**  (greek. *sorites*: pile)

  ◦ One billion grains of sand are a pile of sand.  \( (true) \)
  
  ◦ If a grain of sand is taken away from a pile of sand, the remainder is a pile of sand.  \( (true) \)

It follows:

  ◦ 999 999 999 grains of sand are a pile of sand.  \( (true) \)

Repeated application of this inference finally yields

  ◦ A single grain of sand is a pile of sand.  \( (false!) \)

At which number of grains of sand is the inference not truth-preserving?
Brief Introduction to Fuzzy Theory

- Obviously: There is no precise number of grains of sand, at which the inference to the next smaller number is false.

- Problem: Terms of natural language are vague. (e.g. “pile of sand”, “bald”, “warm”, “fast”, “light”, “high pressure”)

- Note: Vague terms are inexact, but nevertheless not useless.
  - Even for vague terms there are situations or objects, to which they are certainly applicable, and others, to which they are certainly not applicable.
  - In between lies a so-called penumbra (lat. for half shadow) of situations, in which it is unclear whether the terms are applicable, or in which they are applicable only with certain restrictions (“small pile of sand”).
  - Fuzzy theory tries to model this penumbra in a mathematical fashion (“soft transition” between applicable and not applicable).
**Fuzzy Logic**

- **Fuzzy Logic** is an extension of classical logic by values between *true* and *false*.

- **Truth values are any values from the real interval* [0, 1], where 0 \equiv *false and 1 \equiv *true.**

- Therefore necessary: **extension of the logical operators**
  - Negation classical: \( \neg a \), fuzzy: \( \sim a \)  
  - Conjunction classical: \( a \land b \), fuzzy: \( \top(a, b) \)  
  - Disjunction classical: \( a \lor b \), fuzzy: \( \bot(a, b) \)

- **Basic principles of this extension:**
  - For the extreme values 0 and 1 the operations should behave exactly like their classical counterparts (border or corner conditions).
  - For the intermediate values the behavior should be monotone.
  - As far as possible, the laws of classical logic should be preserved.
A fuzzy negation is a function \( \sim : [0, 1] \to [0, 1] \), that satisfies the following conditions:

- \( \sim 0 = 1 \) and \( \sim 1 = 0 \) (boundary conditions)
- \( \forall a, b \in [0, 1] : a \leq b \Rightarrow \sim a \geq \sim b \) (monotony)

If in the second condition the relations \(<\) and \(>\) hold instead of merely \(\leq\) and \(\geq\), the fuzzy negation is called a strict negation.

Additional conditions that are sometimes required are:

- \( \sim \) is a continuous function.
- \( \sim \) is involutive, that is, \( \forall a \in [0, 1] : \sim \sim a = a \).

Involutivity corresponds to the classical law of identity \( \neg \neg a = a \).

The above conditions do not uniquely determine a fuzzy negation.
Fuzzy Negations

- **Standard negation**: \( \sim a = 1 - a \)
- **Threshold negation**: \( \sim (a; \theta) = \begin{cases} 1 & \text{if } x \leq \theta, \\ 0 & \text{otherwise.} \end{cases} \)
- **Cosine negation**: \( \sim a = \frac{1}{2}(1 + \cos \pi a) \)
- **Sugeno negation**: \( \sim (a; \lambda) = \frac{1 - a}{1 + \lambda a} \)
- **Yager negation**: \( \sim (a; \lambda) = (1 - a^\lambda)^\frac{1}{\lambda} \)
A 
t-norm or 
fuzzy conjunction

is a function \( T : [0, 1]^2 \rightarrow [0, 1], \)

that satisfies the following conditions:

- \( \forall a \in [0, 1] : \ T(a, 1) = a \)  \hspace{1cm} \text{(boundary condition)}
- \( \forall a, b, c \in [0, 1] : \ b \leq c \Rightarrow \ T(a, b) \leq T(a, c) \)  \hspace{1cm} \text{(monotony)}
- \( \forall a, b \in [0, 1] : \ T(a, b) = T(b, a) \)  \hspace{1cm} \text{(commutativity)}
- \( \forall a, b, c \in [0, 1] : \ T(a, T(b, c)) = T(T(a, b), c) \)  \hspace{1cm} \text{(associativity)}

Additional conditions that are sometimes required are:

- \( T \) is a continuous function  \hspace{1cm} \text{(continuity)}
- \( \forall a \in [0, 1] : \ T(a, a) < a \)  \hspace{1cm} \text{(sub-idempotency)}
- \( \forall a, b, c, d \in [0, 1] : \ a < b \land c < d \Rightarrow \ T(a, b) < T(c, d) \)  \hspace{1cm} \text{(strict monotony)}

The first two of these conditions (in addition to the top four) define the sub-class of so-called \textit{Archimedic t-norms}. 
t-Norms / Fuzzy Conjunctions

standard conjunction: \( T_{\text{min}}(a, b) = \min\{a, b\} \)

algebraic product: \( T_{\text{prod}}(a, b) = a \cdot b \)

Łukasiewicz: \( T_{\text{luka}}(a, b) = \max\{0, a + b - 1\} \)

drastic product: \( T_{-1}(a, b) = \begin{cases} a & \text{if } b = 1, \\ b & \text{if } a = 1, \\ 0 & \text{otherwise.} \end{cases} \)
A **t-conorm** or **fuzzy disjunction** is a function \( \perp : [0, 1]^2 \rightarrow [0, 1] \), that satisfies the following conditions:

- \( \forall a \in [0, 1] : \perp(a, 0) = a \) (boundary condition)
- \( \forall a, b, c \in [0, 1] : b \leq c \Rightarrow \perp(a, b) \leq \perp(a, c) \) (monotony)
- \( \forall a, b \in [0, 1] : \perp(a, b) = \perp(b, a) \) (commutativity)
- \( \forall a, b, c \in [0, 1] : \perp(a, \perp(b, c)) = \perp(\perp(a, b), c) \) (assoziativity)

Additional conditions that are sometimes required are:

- \( \perp \) is a continuous function (continuity)
- \( \forall a \in [0, 1] : \perp(a, a) > a \) (super-idempotency)
- \( \forall a, b, c, d \in [0, 1] : a < b \land c < d \Rightarrow \perp(a, b) < \perp(c, d) \) (strict monotony)

The first two of these conditions (in addition to the top four) define the sub-class of so-called **Archimedic t-conorms**.
t-Conorms / Fuzzy Disjunctions

standard disjunction: \( \perp_{\text{max}}(a, b) = \max\{a, b\} \)

algebraische sum: \( \perp_{\text{sum}}(a, b) = a + b - a \cdot b \)

Łukasiewicz: \( \perp_{\text{luka}}(a, b) = \min\{1, a + b\} \)

drastic sum: \( \perp_{-1}(a, b) = \begin{cases} a & \text{if } b = 0, \\ b & \text{if } a = 0, \\ 1 & \text{otherwise.} \end{cases} \)
Interplay of the Fuzzy Operators

- It is $\forall a, b \in [0, 1] : \top_{-1}(a, b) \leq \top_{\text{luka}}(a, b) \leq \top_{\text{prod}}(a, b) \leq \top_{\text{min}}(a, b)$. All other possible $t$-norms lie between $\top_{-1}$ and $\top_{\text{min}}$ as well.

- It is $\forall a, b \in [0, 1] : \bot_{\text{max}}(a, b) \leq \bot_{\text{sum}}(a, b) \leq \bot_{\text{luka}}(a, b) \leq \bot_{-1}(a, b)$. All other possible $t$-conorms lie between $\bot_{\text{max}}$ and $\bot_{-1}$ as well.

- Note: Generally neither $\top(a, \sim a) = 0$ nor $\bot(a, \sim a) = 1$ holds.

- A set of operators ($\sim$, $\top$, $\bot$) consisting of a fuzzy negation $\sim$, a $t$-norm $\top$, and a $t$-conorm $\bot$ is called a dual triplet if with these operators DeMorgan’s laws hold, that is, if

$$\forall a, b \in [0, 1] : \sim \top(a, b) = \bot(\sim a, \sim b)$$
$$\forall a, b \in [0, 1] : \sim \bot(a, b) = \top(\sim a, \sim b)$$

- The most frequently used set of operators is the dual triplet ($\sim$, $\top_{\text{min}}$, $\bot_{\text{max}}$) with the standard negation $\sim a \equiv 1 - a$. 
Fuzzy Set Theory

- Classical set theory is based on the notion “is element of” (∈). Alternatively the membership in a set can be described with the help of an indicator function:

  Let $X$ be a set (base set). Then

  $$I_M : X \rightarrow \{0, 1\}, \quad I_M(x) = \begin{cases} 
  1 & \text{if } x \in X, \\
  0 & \text{otherwise,}
\end{cases}$$

  is called **indicator function** of the set $M$ w.r.t. the base set $X$.

- In fuzzy set theory the indicator function of classical set theory is replaced by a membership function:

  Let $X$ be a (classical/crisp) set. Then

  $$\mu_M : X \rightarrow [0, 1], \quad \mu_M(x) \equiv \text{degree of membership of } x \text{ to } M,$$

  is called **membership function** of the fuzzy set $M$ w.r.t. the base set $X$. Usually the fuzzy set is identified with its membership function.
Fuzzy Set Theory: Operations

• In analogy to the transition from classical logic to fuzzy logic, the transition from classical set theory to fuzzy set theory requires an **extension of the operators**.

• **Basic principle of the extension:**
  Draw on the logical definition of the operators.
  ⇒ element-wise application of the logical operators.

• Let $A$ and $B$ be (fuzzy) sets w.r.t. the base set $X$.

  **complement**
  \[
  \begin{align*}
  \text{classical} & : \quad \overline{A} = \{x \in X \mid x \notin A\} \\
  \text{fuzzy} & : \quad \forall x \in X : \quad \mu_{\overline{A}}(x) = \sim \mu_A(x)
  \end{align*}
  \]

  **intersection**
  \[
  \begin{align*}
  \text{classical} & : \quad A \cap B = \{x \in X \mid x \in A \land x \in B\} \\
  \text{fuzzy} & : \quad \forall x \in X : \quad \mu_{A \cap B}(x) = \top(\mu_A(x), \mu_B(x))
  \end{align*}
  \]

  **union**
  \[
  \begin{align*}
  \text{classical} & : \quad A \cup B = \{x \in X \mid x \in A \lor x \in B\} \\
  \text{fuzzy} & : \quad \forall x \in X : \quad \mu_{A \cup B}(x) = \bot(\mu_A(x), \mu_B(x))
  \end{align*}
  \]
Fuzzy Set Operations: Examples

- The fuzzy intersection shown on the left and the fuzzy union shown on the right are independent of the chosen $t$-norm or $t$-conorm.
Fuzzy Intersection: Examples

- Note that all fuzzy intersections lie between the one shown at the top right and the one shown at the right bottom.
Fuzzy Union: Examples

- Note that all fuzzy unions lie between the one shown at the top right and the one shown at the right bottom.
In order to describe a domain by linguistic terms, it is fuzzy-partitioned with the help of fuzzy sets. To each fuzzy set of the partition a linguistic term is assigned.

Common condition: At each point the membership values of the fuzzy sets must sum to 1 (partition of unity).

Example: **fuzzy partition for temperatures**

We define a linguistic variable with the values *cold, tepid, warm* and *hot.*
• The knowledge base contains the fuzzy rules of the controller as well as the fuzzy partitions of the domains of the variables.

• A fuzzy rule reads: if $X_1$ is $A_{i_1}^{(1)}$ and \ldots and $X_n$ is $A_{i_n}^{(n)}$ then $Y$ is $B$. $X_1, \ldots, X_n$ are the measurement values and $Y$ is the controller output. $A_{i_k}^{(k)}$ and $B$ are linguistic terms to which fuzzy sets are assigned.
Example Fuzzy Controller: Inverted Pendulum

\[ \dot{\theta} : \begin{array}{cccccccc} \theta : & -60 & -45 & -30 & -15 & 0 & 15 & 30 & 45 & 60 \\ \end{array} \]

\[ \dot{\theta} : \begin{array}{cccccccc} \theta : & -32 & -24 & -16 & -8 & 0 & 8 & 16 & 24 & 32 \\ \end{array} \]

**abbreviations**

- pb – positive big
- pm – positive medium
- ps – positive small
- az – approximately zero
- ns – negative small
- nm – negative medium
- nb – negative big

\begin{array}{|c|c|c|c|c|c|c|c|}
\hline
\dot{\theta} \setminus \theta & nb & nm & ns & az & ps & pm & pb \\
\hline
pb & & & & & ps & pb & \\
\hline
pm & & & & & & pm & \\
\hline
ps & & & & & nm & az & ps \\
\hline
az & & nb & nm & ns & az & ps & pm & pb \\
\hline
ns & & & & & ns & az & pm \\
\hline
nm & & & & & & nm & \\
\hline
nb & & & & & & nb & ns \\
\hline
\end{array}
Mamdani–Assilian Controller

Rule evaluation in a Mamdani–Assilian controller. The input tuple \((25, -4)\) yields the fuzzy output shown on the right. From this fuzzy set the actual output value is computed by defuzzification, e.g. by the mean of maxima method (MOM) or the center of gravity method (COG).
A fuzzy control system with one measurement and one control variable and three fuzzy rules. Each pyramid is specified by one fuzzy rule. The input value $x^*$ leads to the fuzzy output shown in gray.
The evaluation of the fuzzy rules yields an output fuzzy set.
The output fuzzy set has to be turned into a crisp controller output.
This process is called defuzzification.

The most important defuzzification methods are:

- **Center of Gravity** (COG)
  The center of gravity of the area under the output fuzzy set.

- **Center of Area** (COA)
  The point that divides the area under the output fuzzy set into equally large parts.

- **Mean of Maxima** (MOM)
  The arithmetic mean of the points with maximal degree of membership.
Takagi–Sugeno–Kang Controller (TSK Controller)

- The rules of a Takagi–Sugeno–Kang controller have the same kind of antecedent as the rules of a Mamdani–Assilian controller, but a different kind of consequent:

\[ R_i: \text{if } x_1 \text{ is } \mu_i^{(1)} \text{ and } \ldots \text{ and } x_n \text{ is } \mu_i^{(n)}, \text{ then } y = f_i(x_1, \ldots, x_n). \]

The consequent of a Takagi–Sugeno–Kang rule specifies a function of the inputs that is to be computed if the antecedent is satisfied.

- Let \( \tilde{a}_i \) be the activation of the antecedent of the rule \( R_i \), that is,

\[ \tilde{a}_i(x_1, \ldots, x_n) = \top \left( \top \left( \ldots \top \left( \mu_i^{(1)}(x_1), \mu_i^{(2)}(x_2) \right), \ldots \right), \mu_i^{(n)}(x_n) \right). \]

- Then the output of a Takagi–Sugeno–Kang controller is computed as

\[ y(x_1, \ldots, x_n) = \frac{\sum_{i=1}^{r} \tilde{a}_i \cdot f_i(x_1, \ldots, x_n)}{\sum_{i=1}^{r} \tilde{a}_i}, \]

that is, the controller output is a weighted average of the outputs of the individual rules, where the activations of the antecedents provide the weights.
Neuro-Fuzzy Systems

- **Disadvantages of Neural Networks:**
  - Training results are difficult to interpret (black box).
    The result of training an artificial neural network are matrices or vectors of real-valued numbers. Even though the computations are clearly defined, humans usually have trouble understanding what is going on.
  - It is difficult to specify and incorporate prior knowledge.
    Prior knowledge would have to be specified as the mentioned matrices or vectors of real-valued numbers, which are difficult to understand for humans.

- **Possible Remedy:**
  - Use a hybrid system, in which an artificial neural network is coupled with a rule-based system.
    The rule-based system can be interpreted and set up by a human.
  - One such approach are **neuro-fuzzy systems**.
Neuro-fuzzy systems are commonly divided into cooperative and hybrid systems.

- **Cooperative Models:**
  - A neural network and a fuzzy controller work independently.
  - The neural network generates (offline) or optimizes (online) certain parameters.

- **Hybrid Models:**
  - Combine the structure of a neural network and a fuzzy controller.
  - A hybrid neuro-fuzzy controller can be interpreted as a neural network and can be implemented with the help of a neural network.
  - Advantages: integrated structure; no communication between two different models is needed; in principle, both offline and online training are possible;

Hybrid models are more accepted and more popular than cooperative models.
Neuro-Fuzzy Systems: Hybrid Methods

- Hybrid methods map fuzzy sets and fuzzy rules to a neural network structure.
- The activation $\tilde{a}_i$ of the antecedent of a Mamdani–Assilian rule
  $$R_i: \text{if } x_1 \text{ is } \mu^{(1)}_i \text{ and } \ldots \text{ and } x_n \text{ is } \mu^{(n)}_i, \text{ then } y \text{ is } \nu_i.$$ 
  or of a Takagi–Sugeno–Kang rule
  $$R_i: \text{if } x_1 \text{ is } \mu^{(1)}_i \text{ and } \ldots \text{ and } x_n \text{ is } \mu^{(n)}_i, \text{ then } y = f_i(x_1, \ldots, x_n).$$
  is computed with a $t$-norm $\top$ (most commonly $\top_{\min}$).
- For given input values $x_1, \ldots, x_n$ the network structure has to compute:
  $$\tilde{a}_i(x_1, \ldots, x_n) = \top \left( \top \left( \ldots \top \left( \mu^{(1)}_i(x_1), \mu^{(2)}_i(x_2) \right), \ldots \right), \mu^{(n)}_i(x_n) \right).$$
  (Since $t$-norms are associative, it does not matter in which order the membership
degrees are combined by successive pairwise applications of the $t$-norm.)
Computing the activation $\tilde{a}_i$ of the antecedent of a fuzzy rule:

The fuzzy sets appearing in the antecedents of fuzzy rules can be modeled as connection weights:

\[
x_1 \xrightarrow{\mu_i^{(1)}} \top \xrightarrow{\tilde{a}_i} \xrightarrow{\mu_i^{(2)}} x_2
\]

as activation functions:

\[
x_1 \xrightarrow{\mu_i^{(1)}} \top \xrightarrow{\tilde{a}_i} \xrightarrow{\mu_i^{(2)}} x_2
\]

The neurons in the first hidden layer represent the rule antecedents and the connections from the input units represent the fuzzy sets of these antecedents.

The neurons in the first hidden layer represent the fuzzy sets and the neurons in the second hidden layer represent the rule antecedents ($\top$ is a $t$-norm).
From Fuzzy Rule Base to Network Structure

If the fuzzy sets are represented as activation functions (right on previous slide), a fuzzy rule base is turned into a network structure as follows:

1. For every input variable $x_i$: create a neuron in the input layer.

2. For every output variable $y_i$: create a neuron in the output layer.

3. For every fuzzy set $\mu_i^{(j)}$: create a neuron in the first hidden layer and connect it to the input neuron corresponding to $x_i$.

4. For every fuzzy rule $R_i$: create a (rule) neuron in the second hidden layer and specify a $t$-norm for computing the rule (antecedent) activation.

5. Connect each rule neuron to the neurons that represent the fuzzy sets of the antecedent of its corresponding fuzzy rule $R_i$.

6. (This step depends on whether the controller is of the Mamdani–Assilian or of the Takagi–Sugeno–Kang type; see next slide.)
Mamdani–Assilian controller:

6. Connect each rule neuron to the output neuron corresponding to the consequent domain of its fuzzy rule. As connection weight choose the consequent fuzzy set of the fuzzy rule. Furthermore, a $t$-conorm for combining the outputs of the individual rules and a defuzzification method have to be integrated adequately into the output neurons (e.g. as network input and activation functions).

Takagi–Sugeno–Kang controller:

6. For each rule neuron, create a sibling neuron that computes the output function of the corresponding fuzzy rule and connect all input neurons to it (arguments of the consequent function). All rule neurons that refer to the same output domain as well as their sibling neurons are connected to the corresponding output neuron (in order to compute the weighted average of the rule outputs).

The resulting network structure can now be trained with procedures that are analogous to those of standard neural networks (e.g. error backpropagation).
This ANFIS network represents the fuzzy rule base (Takagi–Sugeno–Kang rules):

\[ R_1: \text{if } x_1 \text{ is } \mu_1^{(1)} \text{ and } x_2 \text{ is } \mu_1^{(2)}, \text{ then } y = f_1(x_1, x_2) \]
\[ R_2: \text{if } x_1 \text{ is } \mu_1^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)}, \text{ then } y = f_2(x_1, x_2) \]
\[ R_3: \text{if } x_1 \text{ is } \mu_2^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)}, \text{ then } y = f_3(x_1, x_2) \]
This NEFCON network represents the fuzzy rule base (Mamdani-Assilian rules):

\[ R_1: \text{if } x_1 \text{ is } \mu_1^{(1)} \text{ and } x_2 \text{ is } \mu_1^{(2)}, \text{ then } y \text{ is } \nu_1 \]

\[ R_2: \text{if } x_1 \text{ is } \mu_1^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)}, \text{ then } y \text{ is } \nu_1 \]

\[ R_3: \text{if } x_1 \text{ is } \mu_2^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)}, \text{ then } y \text{ is } \nu_2 \]

\[ R_4: \text{if } x_1 \text{ is } \mu_3^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)}, \text{ then } y \text{ is } \nu_3 \]

\[ R_5: \text{if } x_1 \text{ is } \mu_3^{(1)} \text{ and } x_2 \text{ is } \mu_3^{(2)}, \text{ then } y \text{ is } \nu_3 \]
This NEFCLASS network represents fuzzy rules that predict classes:

\[ \begin{align*}
R_1 &: \text{ if } x_1 \text{ is } \mu_1^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)} , \text{ then } \text{ class } c_1 \\
R_2 &: \text{ if } x_1 \text{ is } \mu_1^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)} , \text{ then } \text{ class } c_1 \\
R_3 &: \text{ if } x_1 \text{ is } \mu_2^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)} , \text{ then } \text{ class } c_2 \\
R_4 &: \text{ if } x_1 \text{ is } \mu_3^{(1)} \text{ and } x_2 \text{ is } \mu_2^{(2)} , \text{ then } \text{ class } c_3 \\
R_5 &: \text{ if } x_1 \text{ is } \mu_3^{(1)} \text{ and } x_2 \text{ is } \mu_3^{(2)} , \text{ then } \text{ class } c_3
\end{align*} \]
NEFCLASS: Initializing the Fuzzy Partitions

- NEFCLASS is based on modified Wang–Mendel procedure. [Nauck 1997]

- NEFCLASS first fuzzy partitions the domain of each variable, usually with a given number of equally sized triangular fuzzy sets; the boundary fuzzy sets are “shouldered” (membership 1 to the boundary).

- Based on the initial fuzzy partitions, the initial rule base is selected.

---

![Diagram of fuzzy partitions](image1)

![Diagram of fuzzy partitions](image2)
**NEFCLASS: Initializing the Rule Base**

\[ A := \emptyset; \]

**for each** training pattern \( p \) **do begin**

- find rule antecedent \( A \) such that \( A(p) \) is maximal;

  **if** \( A \notin A \) **then**

    \[ A := A \cup \{A\}; \]

  **end**

\[ R := \emptyset; \]

**for each** antecedent \( A \in A \) **do begin**

- find best consequent \( C \) for antecedent \( A \);

  - create rule base candidate \( R = (A, C) \);

  - determine performance of \( R \);

  \[ R := R \cup \{R\}; \]

**end**

**return** rule base \( R \);

Fuzzy rule bases may also be created from prior knowledge or using fuzzy cluster analysis, fuzzy decision trees, evolutionary algorithms etc.
NEFCLASS: Selecting the Rule Base

- In order to reduce/limit the number of initial rules, their performance is evaluated.
- The performance of a rule $R_i$ is computed as

$$\text{Perf}(R_i) = \frac{1}{m} \sum_{j=1}^{m} e_{ij} \tilde{a}_i(\vec{x}_j)$$

where $m$ is the number of training patterns and $e_{ij}$ is an error indicator,

$$e_{ij} = \begin{cases} 
+1 & \text{if class}(\vec{x}_j) = \text{cons}(R_i), \\
-1 & \text{otherwise}.
\end{cases}$$

- Sort the rules in the initial rule base by their performance.
- Choose either the best $r$ rules or the best $r/c$ rules per class, where $c$ is the number of classes.
- The number $r$ of rules in the rule base is either provided by a user or is automatically determined in such a way that all patterns are covered.
NEFCLASS: Computing the Error Signal

- Let $o_k^{(l)}$ be the desired output and $\text{out}_k^{(l)}$ the actual output of the $k$-th output neuron (class $c_k$).

- Fuzzy error ($k$-th output/class):
  
  $e_k^{(l)} = 1 - \gamma(\varepsilon_k^{(l)})$,

  where $\varepsilon_k^{(l)} = o_k^{(l)} - \text{out}_k^{(l)}$ and

  $\gamma(z) = e^{-\beta z^2}$,

  where $\beta > 0$ is a sensitivity parameter: larger $\beta$ means larger error tolerance.

- Error signal ($k$-th output/class):
  
  $\delta_k^{(l)} = \text{sgn}(\varepsilon_k^{(l)}) e_k^{(l)}$. 
NEFCLASS: Computing the Error Signal

- Rule error signal (rule $R_i$ for $c_k$):
  \[ \delta_R^{(l)} = \text{out}_{R_i}^{(l)} \left( 1 - \text{out}_{R_i}^{(l)} \right) \delta_k^{(l)} , \]
  (the factor "out(1 − out)" is chosen in analogy to multi-layer perceptrons).

- Find input variable $x_j$ such that
  \[ \mu_{i}^{(j)}(\hat{\nu}^{(l)}) = \tilde{\alpha}_i(\hat{\nu}^{(l)}) = \min_{\nu=1}^{d} \mu_{i}^{(\nu)}(\hat{\nu}^{(l)}) , \]
  where $d$ is the number of inputs (find antecedent term of $R_i$ giving the smallest membership degree, which yields the rule activation).

- Adapt parameters of the fuzzy set $\mu_{i}^{(j)}$ (see next slide for details).
NEFCLASS: Training the Fuzzy Sets

- Triangular fuzzy set as an example:

\[ \mu_{a,b,c}(x) = \begin{cases} \frac{x-a}{b-a} & \text{if } x \in [a, b), \\ \frac{c-x}{c-b} & \text{if } x \in [b, c], \\ 0 & \text{otherwise.} \end{cases} \]

- Parameter changes (learning rate \( \eta \)):

\[
\begin{align*}
\Delta b &= +\eta \cdot \delta^{(l)}_{R_i} \cdot (c - a) \cdot \text{sgn}(i_{j}^{(l)} - b) \\
\Delta a &= -\eta \cdot \delta^{(l)}_{R_i} \cdot (c - a) + \Delta b \\
\Delta c &= +\eta \cdot \delta^{(l)}_{R_i} \cdot (c - a) + \Delta b
\end{align*}
\]

- Heuristics: the fuzzy set to train is moved away from \( x^* \) (towards \( x^* \)) and its support is reduced (increased) in order to reduce (increase) the degree of membership of \( x^* \).
When fuzzy sets of a fuzzy partition are trained, restrictions apply, so correction procedures are needed to

- ensure valid parameter values
- ensure non-empty intersections of neighboring fuzzy sets
- preserve relative positions
- preserve symmetry
- ensure partition of unity (membership degrees sum to 1 everywhere)

On the right: example of a correction of a fuzzy partition with three fuzzy sets.
Objective: Remove variables, rules and fuzzy sets, in order to improve interpretability and generalization ability.

\textbf{repeat}

select pruning method;
\textbf{repeat}

execute pruning step;
train fuzzy sets;
\textbf{if} no improvement\textbf{then} undo step;
\textbf{until} there is no improvement;
\textbf{until} no further method;

1. Remove variables (correlation, information gain etc.)
2. Remove rules (performance of a rule)
3. Remove antecedent terms (satisfaction of a rule)
4. Remove fuzzy sets

- After each pruning step the fuzzy sets need to be retrained, in order to obtain the optimal parameter values for the new structure.

- A pruning step that does not improve performance, the system is reverted to its state before the pruning step.
NEFCLASS: Full Procedure

fuzzy partitions

initial rule base

trained rule base

pruned rule base
picture not available in online version
Stock Index Prediction (DAX) [Siekmann 1999]

- Prediction of the daily relative changes of the German stock index (Deutscher Aktienindex, DAX).
- Based on time series of stock indices and other quantities between 1986 and 1997.

**Input Variables:**

- DAX (Germany)
- Composite DAX (Germany)
- Dow Jones industrial index (USA)
- Nikkei index (Japan)
- Morgan–Stanley index Germany
- Morgan–Stanley index Europe
- German 3 month interest rate
- return Germany
- US treasure bonds
- price to income ratio
- exchange rate DM / US-$
- gold price
DAX Prediction: Example Rules

- **trend rule:**
  - **if** DAX is decreasing **and** US-$ is decreasing
  - **then** DAX prediction is decreasing
  - **with** high certainty

- **turning point rule:**
  - **if** DAX is decreasing **and** US-$ is increasing
  - **then** DAX prediction is increasing
  - **with** low certainty

- **delay rule:**
  - **if** DAX is stable **and** US-$ is decreasing
  - **then** DAX prediction is decreasing
  - **with** very high certainty

- **general form:**
  - **if** $x_1$ is $\mu_1$ **and** $x_2$ is $\mu_2$ **and** ... **and** $x_n$ is $\mu_n$
  - **then** $y$ is $\nu$
  - **with** certainty $c$

Initial rules may be provided by financial experts.
DAX Prediction: Architecture

Input $x_1$ and $x_2$ are fed into the network as antecedents. The membership functions determine whether the antecedents are increasing, stable, or decreasing. These membership functions feed into the rules layer, which generates increasing, stable, and decreasing consequents. The consequents are then combined to produce the output $y$. The architecture shows how the input values are processed through the network to predict the output.
DAX Prediction: From Rules to Neural Network

- Finding the membership values (evaluate membership functions):

- Evaluating the rules (computing the rule activation for \( r \) rules):

\[
\forall j \in \{1, \ldots, r\} : \quad \tilde{a}_j(x_1, \ldots, x_d) = \prod_{i=1}^{d} \mu_{j}^{(i)}(x_i).
\]

- Accumulation of \( r \) rule activations, normalization:

\[
y = \sum_{j=1}^{r} w_j \frac{c_j \tilde{a}_j(x_1, \ldots, x_d)}{\sum_{k=1}^{r} c_k \tilde{a}_k(x_1, \ldots, x_d)}, \quad \text{where} \quad \sum_{j=1}^{r} w_j = 1.
\]
DAX Prediction: Training the Network

- Membership degrees of different inputs share their parameters, e.g.
  \[ \mu_{\text{stable}}^{(\text{DAX})} = \mu_{\text{stable}}^{(\text{Composite DAX})} \]
  Advantage: number of free parameters is reduced.

- Membership functions of the same input variable must not “pass each other”, but must preserve their original order:
  \[ \mu_{\text{negative}} < \mu_{\text{zero}} < \mu_{\text{positive}} \quad \mu_{\text{decreasing}} < \mu_{\text{stable}} < \mu_{\text{increasing}} \]
  Advantage: optimized rule base remains interpretable.

- The parameters of the fuzzy sets, the rule certainties, and the rule weights are optimized with a backpropagation approach.

- Pruning methods are employed to simplify the rules and the rule base.
DAX Prediction: Trading Performance

- Different prediction/trading models for the DAX: naive, Buy&Hold, linear model, multi-layer perceptron, neuro-fuzzy system
- Profit and loss obtained from trading according to prediction.
- Validation period: March 1994 to April 1997

pictures not available in online version
Surface Control of Car Body Parts (BMW)

- **Previous Approach:**
  - surface control is done manually
  - experienced employee treats surface with a grinding block
  - human experts classify defects by linguistic terms
  - cumbersome, subjective, error-prone, time-consuming

- **Suggested Approach:**
  - digitization of the surface with optical measurement systems
  - characterization of the shape defects by mathematical properties (close to the subjective features)
Surface Control: Topometric Measurement System

picture not available in online version
Surface Control: Data Processing

picture not available in online version
Surface Control: Color Coded Representation

picture not available in online version
Surface Control: 3D Representation

- **Sink mark**
  - Slight flat-based sink inwards

- **Uneven surface**
  - Several neighboring sink marks

- **Press mark**
  - Local smoothing of the surface

- **Wavy surface**
  - Several severe foldings in series
Data Characteristics

- 9 master pieces with a total of 99 defects were analyzed.
- 42 features were computed for each defect.
- Defect types are fairly unbalanced, rare types were dropped.
- Some extremely correlated features were dropped ⇒ 31 features remain.
- Remaining 31 features were ranked by their importance.
- Experiment was conducted with 4-fold stratified cross validation.

Accuracy of different classifiers:

<table>
<thead>
<tr>
<th>accuracy</th>
<th>DC</th>
<th>NBC</th>
<th>DT</th>
<th>NN</th>
<th>NEFCLASS</th>
</tr>
</thead>
<tbody>
<tr>
<td>training</td>
<td>46.8%</td>
<td>89.0%</td>
<td>94.7%</td>
<td>90.0%</td>
<td>81.6%</td>
</tr>
<tr>
<td>test</td>
<td>46.8%</td>
<td>75.6%</td>
<td>75.6%</td>
<td>85.5%</td>
<td>79.9%</td>
</tr>
</tbody>
</table>
Surface Control: Fuzzy Rule Base

\[
R_1: \text{if } \text{max\_dist\_to\_cog is fun\_2 and min\_extrema is fun\_1 and max\_extrema is fun\_1 then type is press\_mark}
\]

\[
R_2: \text{if } \text{max\_dist\_to\_cog is fun\_2 and all\_extrema is fun\_1 and max\_extrema is fun\_2 then type is sink\_mark}
\]

\[
R_3: \text{if } \text{max\_dist\_to\_cog is fun\_3 and min\_extrema is fun\_1 and max\_extrema is fun\_1 then type is uneven\_surface}
\]

\[
R_4: \text{if } \text{max\_dist\_to\_cog is fun\_2 and min\_extrema is fun\_1 and max\_extrema is fun\_1 then type is uneven\_surface}
\]

\[
R_5: \text{if } \text{max\_dist\_to\_cog is fun\_2 and all\_extrema is fun\_1 and min\_extrema is fun\_1 then type is press\_mark}
\]

\[
R_6: \text{if } \text{max\_dist\_to\_cog is fun\_3 and all\_extrema is fun\_1 and max\_extrema is fun\_1 then type is uneven\_surface}
\]

\[
R_7: \text{if } \text{max\_dist\_to\_cog is fun\_3 and min\_extrema is fun\_1 then type is uneven\_surface}
\]

NEFCLASS rules for surface defect classification
Neuro-Fuzzy Systems: Summary

- Neuro-fuzzy systems can be useful for **discovering knowledge** in the form of rules and rule bases.

- The fact that they are **interpretable**, allows for plausibility checks and improves acceptance.

- Neuro-fuzzy systems exploit tolerances in the underlying system in order to find near-optimal solutions.

- Training procedures for neuro-fuzzy systems have to be able to cope with restrictions in order to preserve the semantics of the original model.

- **No (fully) automatic model generation.**
  \[\Rightarrow\] A user has to work and interact with the system.

- Simple training methods support **exploratory data analysis**.