

# Artificial Neural Networks

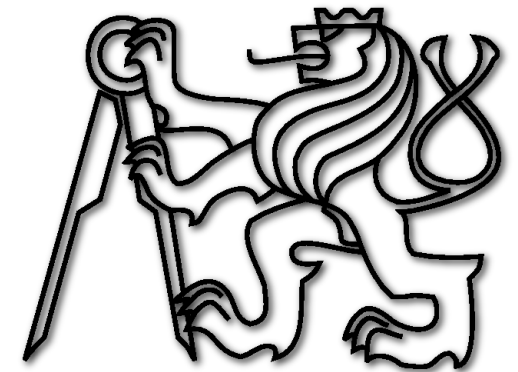
## Unsupervised learning: SOM



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Czech Technical University in Prague*



# Outline

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- Competitive learning.
- Self-organization, Vector Quantization, Cluster Analysis.
- SOM architecture and learning.
- SOM visualizations.
- SOM evaluation.

# Competitive Learning

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- Nature inspired.
- No arbiter needed – unsupervised learning.
- Individuals (units, neurons) learn from examples.
- System **self-organizes**.
- Now we are going to apply this to **cluster analysis**.

# SOM

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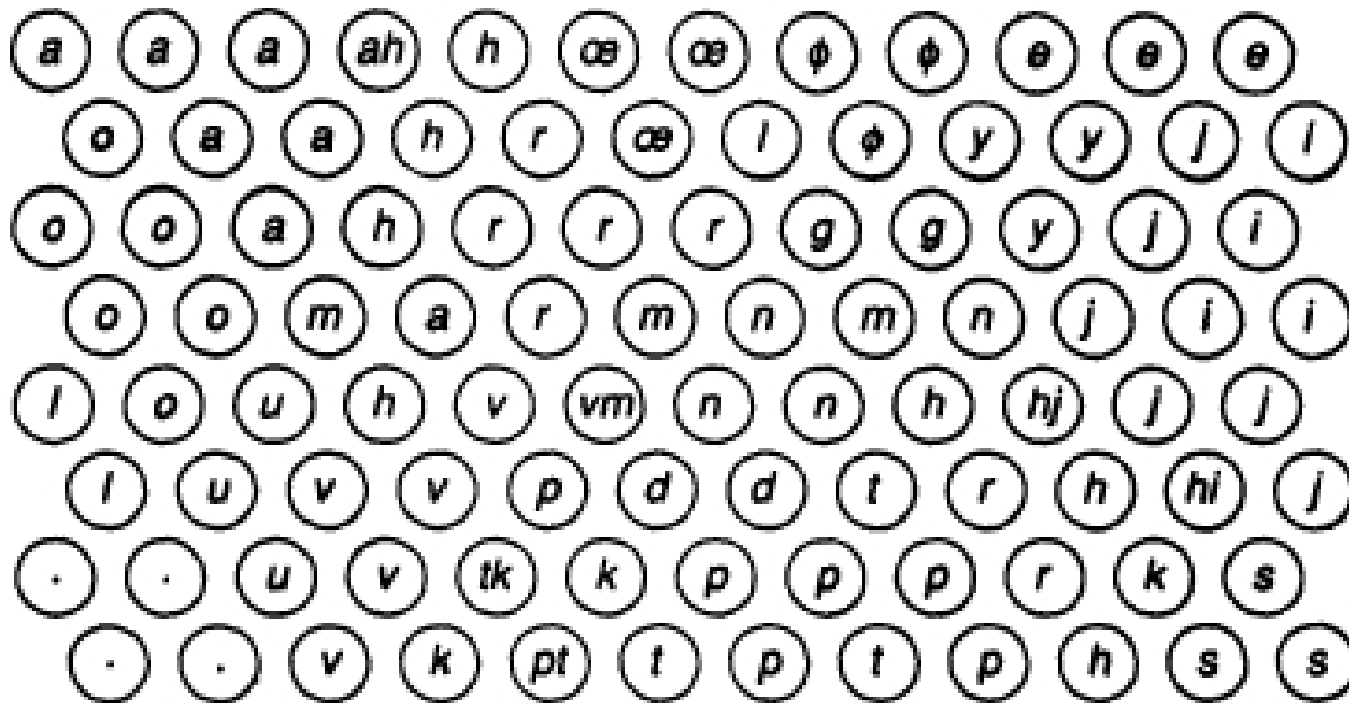
- SOM = Self Organizing Maps.
- Prof. Teuvo Kohonen, Finsko, TU Helsinki, 1981, several thousands scientific publications since...



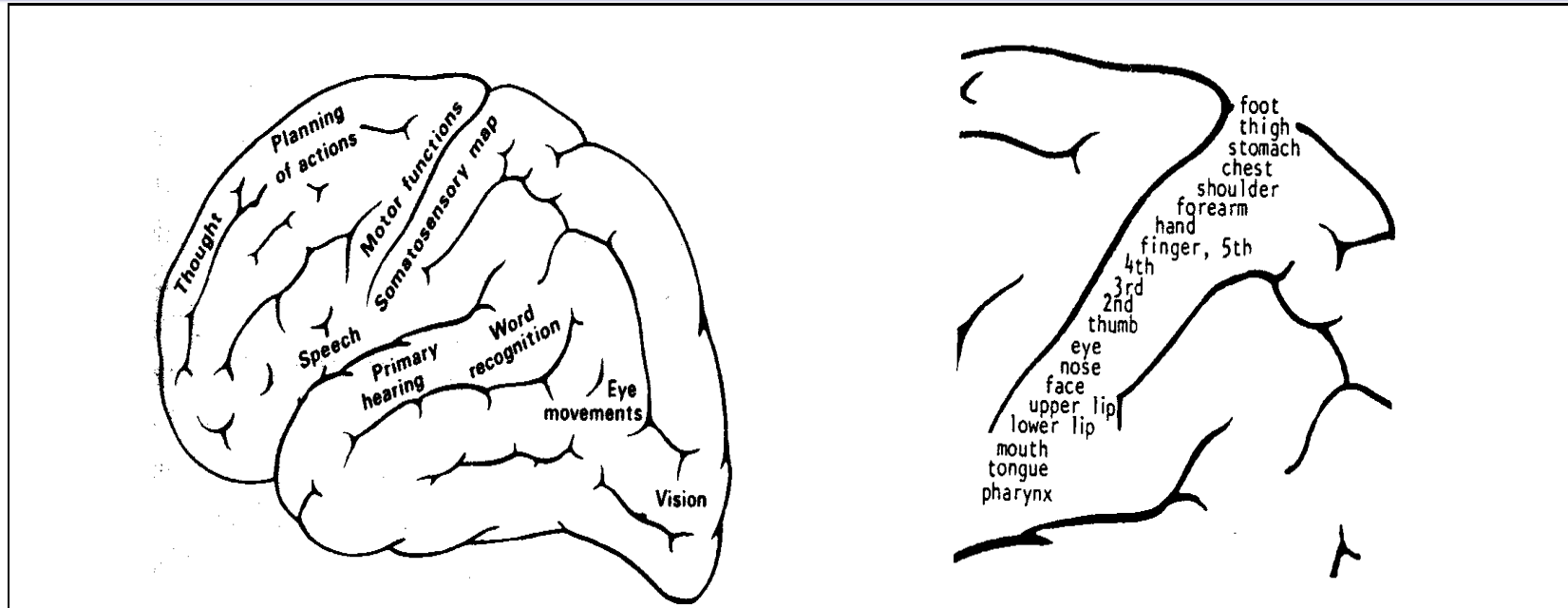
# SOM – Kohonen's Application

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- Original application: phonetic “typewriter”:
  - Finish language.



# SOM Inspiration

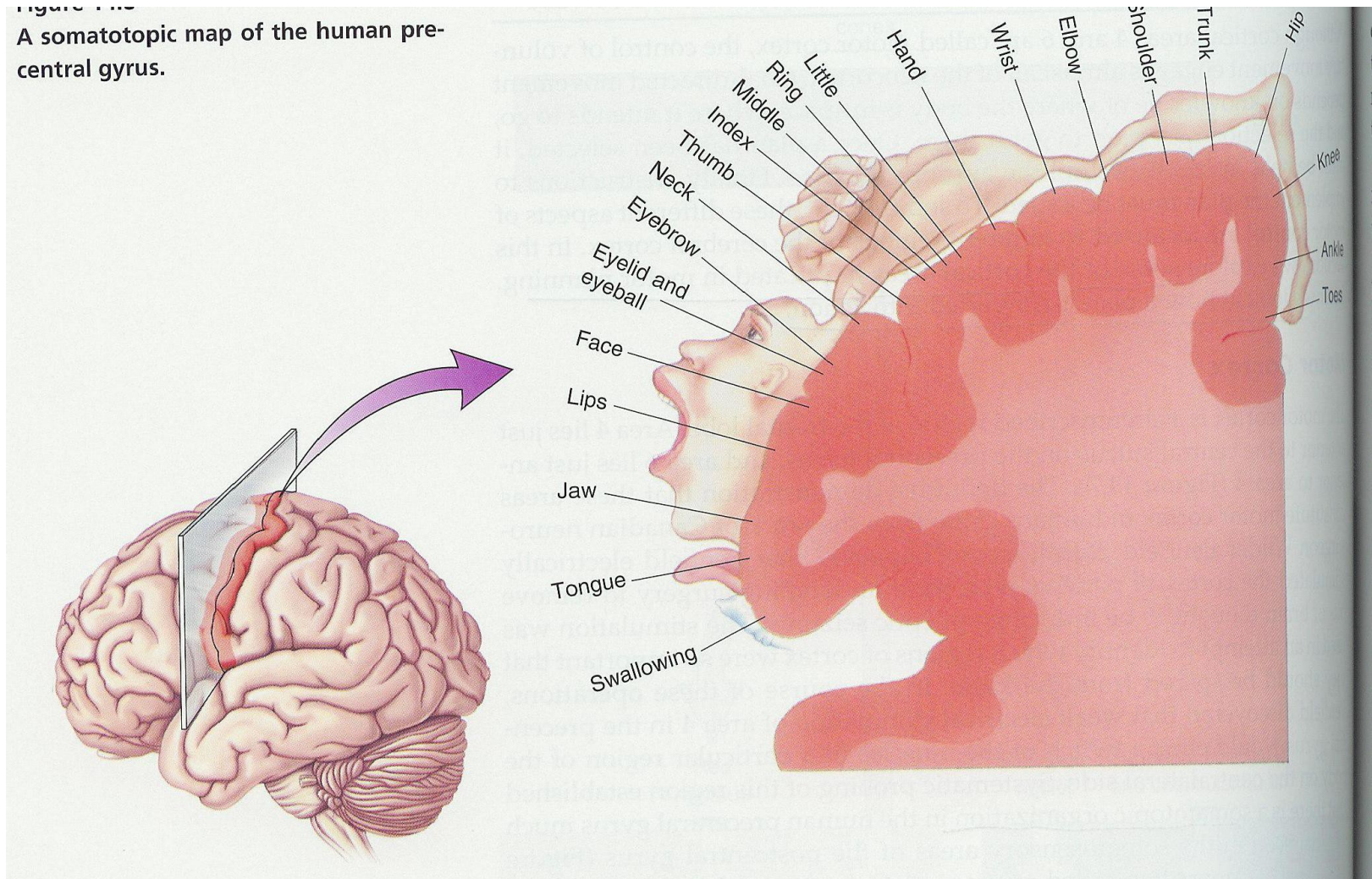


- Brain represents the world in a **topological way**.
- Exterior spatial relations are mapped to similar spatial relations in the brain:
  - i.e. signals from hand and arm are processed nearby.



# SOM Inspiration II

Figure 1.16  
A somatotopic map of the human pre-central gyrus.



Bear, Connors & Paradiso (2001). *Neuroscience: Exploring The Brain*. Pg. 474.

# SOM Overview

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- Single layer, feed-forward.
- Unsupervised, **self-organization**.
- No output, instead **Winner-takes-all**.
- Used for **cluster analysis**.
- Performs **vector quantization**.
- Not a classifier!
  - But can be simply transformed into one by adding another layer.



# What is Self-Organization?

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- Self-organization of a system is a process which leads to a rise of a quality of its inner configuration while not using any information from outside.
- Self-organization clears up relationships between parts of a system.

# What is Cluster Analysis

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- Assignment of a set of observations into subsets (clusters).
- A measure of similarity is defined:
  - observations in the same cluster are similar,
  - observations between two clusters are dissimilar.
- Classic cluster analysis works with  $R^n$  input space observations.

See: [http://en.wikipedia.org/wiki/Cluster\\_analysis](http://en.wikipedia.org/wiki/Cluster_analysis)

# What is Vector Quantization

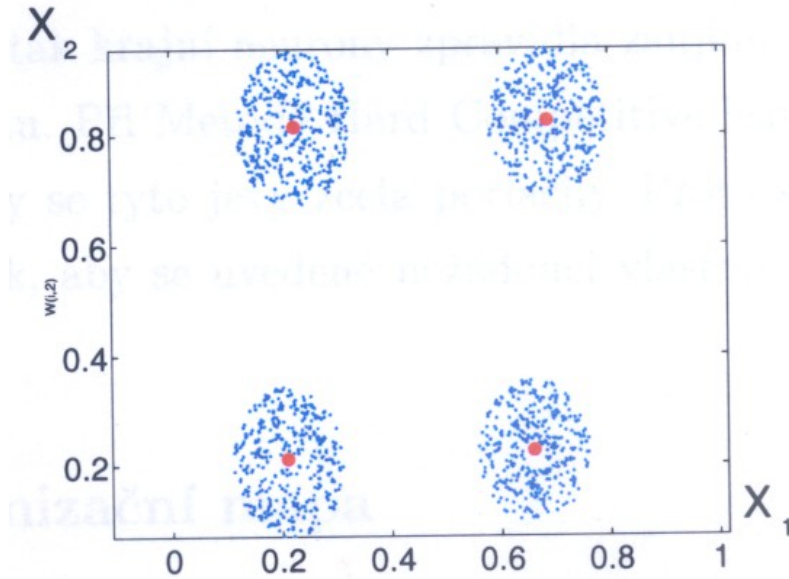
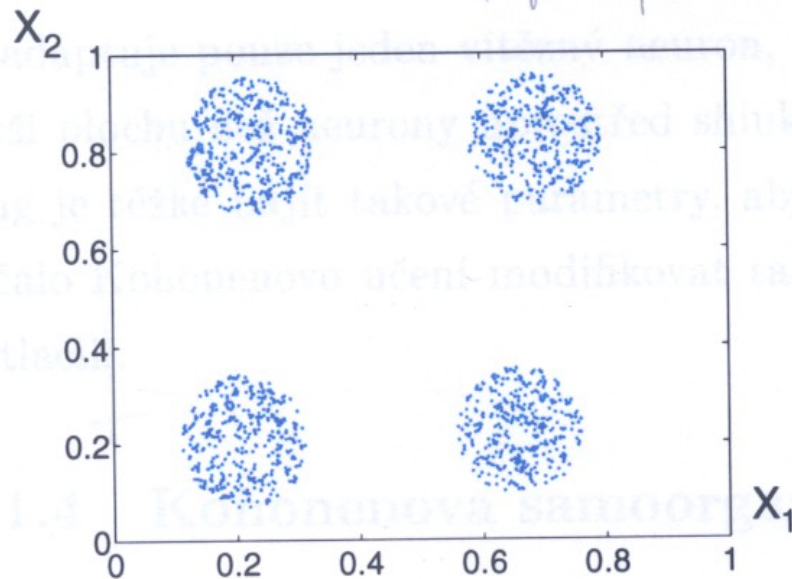
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The goal of Vector Quantization is to approximate the probability density  $p(x)$  of real input vectors  $\mathbf{x} \in \mathbf{R}^n$  distribution using finite number of representatives  $\mathbf{w}_i \in \mathbf{R}^n$ .

The representative vectors tend to drift there where the data is dense, while there tends to be only a few of them where data is sparsely located. In this manner, the net tends to approximate the probability density of the input data. *Hollmen '96*

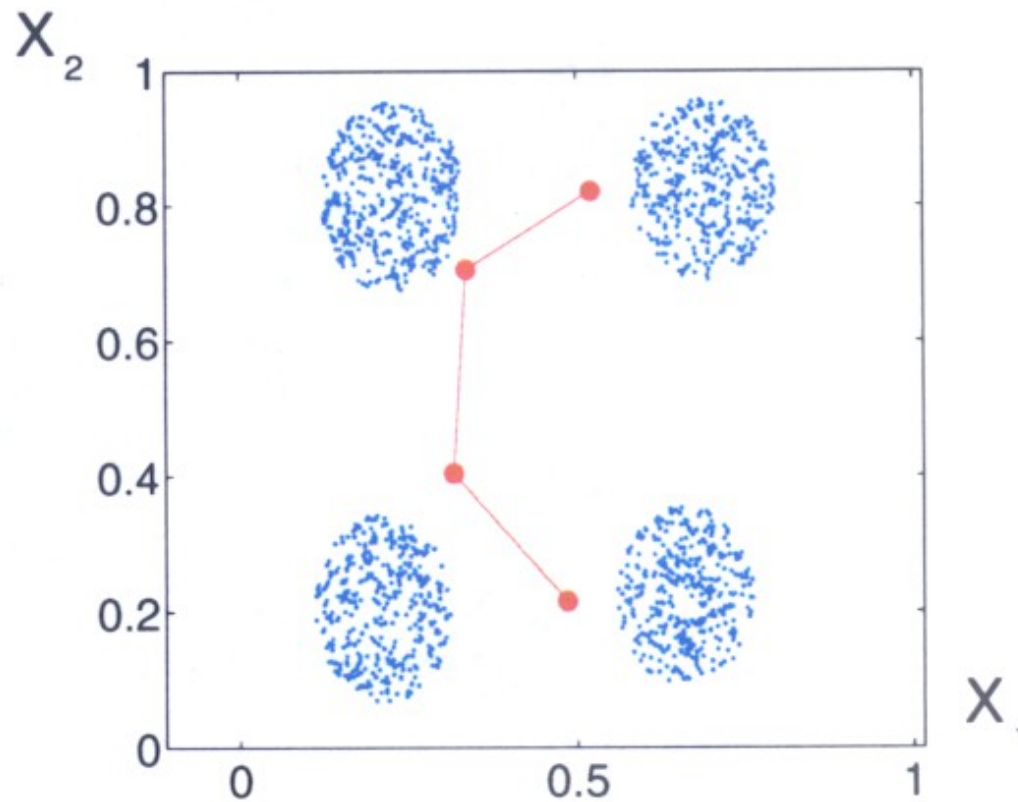
# Vector Quantization Example

Blue points are the input vectors.



Red points are the representatives.

# VQ by SOM

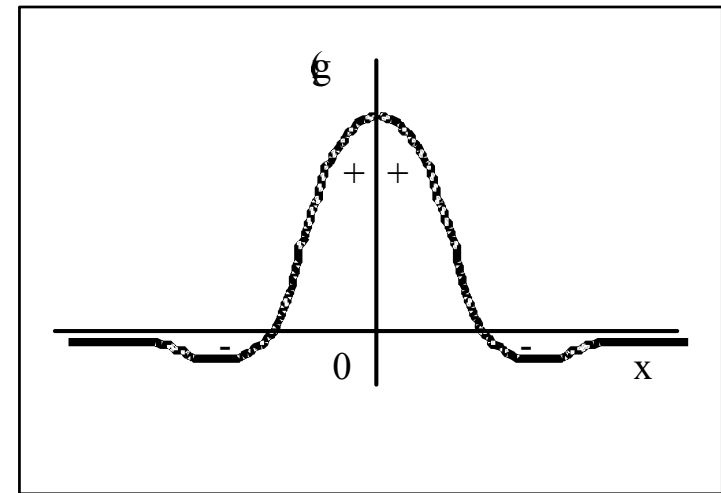


1D SOM of 4 neurons

# Why the Different Result?

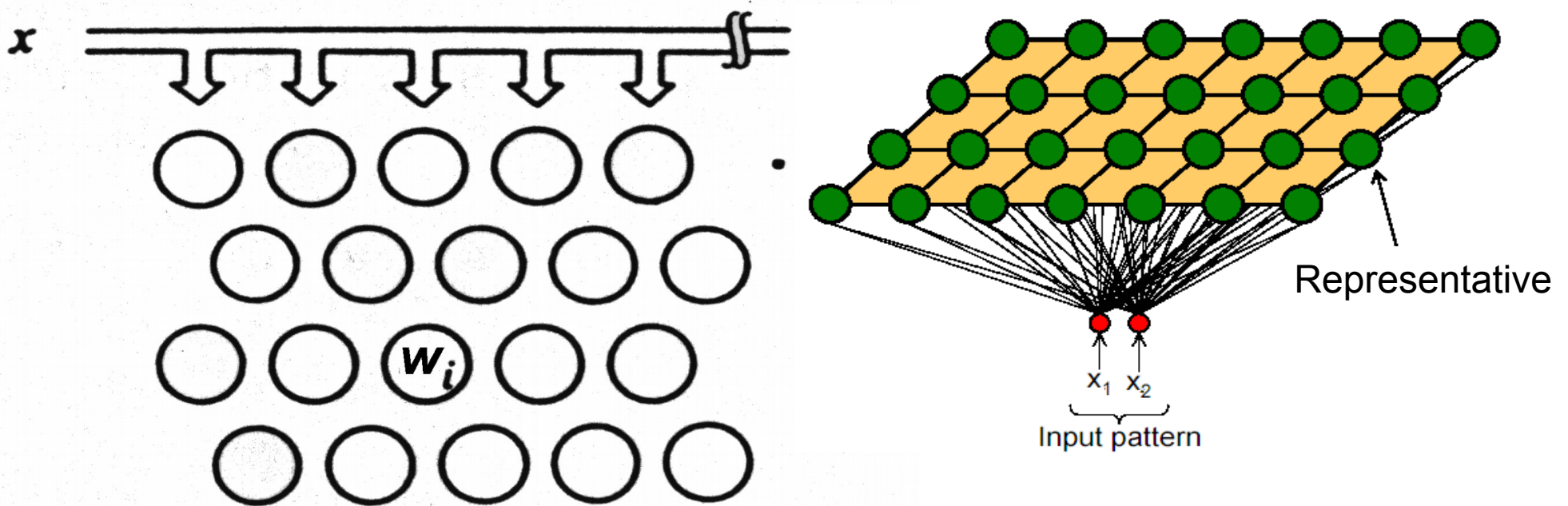
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- SOM works with neighbourhood.
- Representatives influence each other.
- They form “elastic”:
  - chain for 1D SOM,
  - mesh for higher dimensions.





# SOM Architecture 1/3



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- Typically: 2D mesh of representatives (neurons)
-

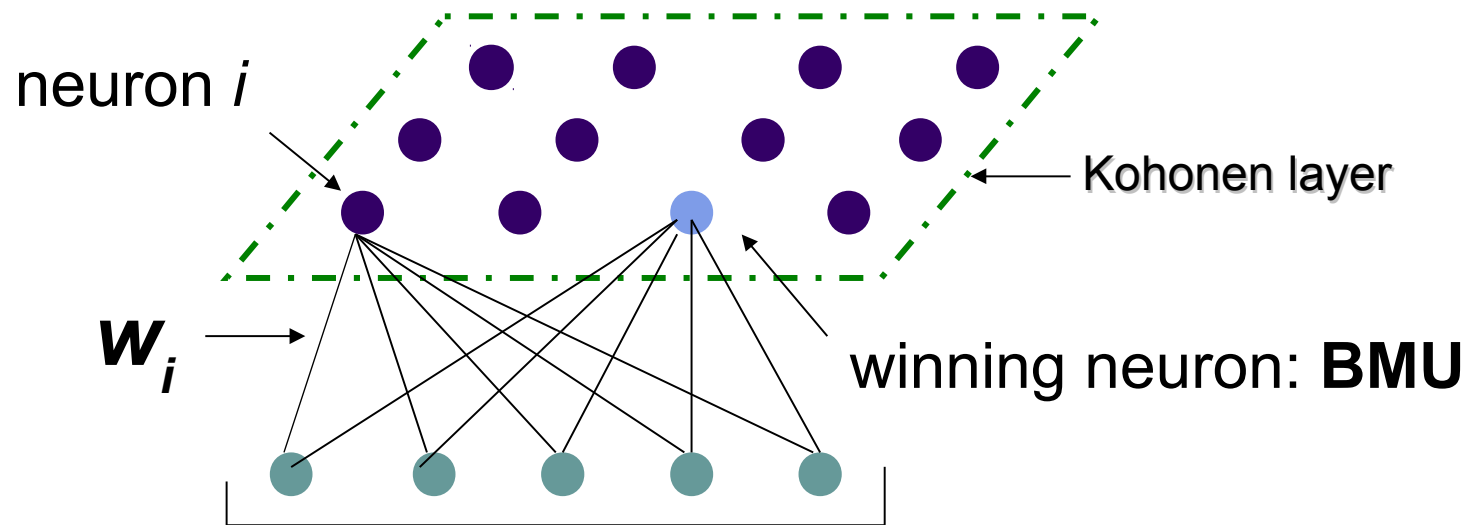
# SOM Architecture 2/3

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- Arrangements:
  - 1D linear quite often,
  - 2D mesh most frequently,
  - 3D (and higher dimensions) exceptionally – problematic visualization.
- The arrangement defines **neighbourhood** of a neuron.
- Kohonen suggests: rectangular SOM!

# SOM Architecture 3/3

- Input vector  $\mathbf{x}$  has a dimension  $N$ .
- Each neuron has a weight vector  $\mathbf{w}$  of the same dimension  $N$ .
- Weight vectors of all neurons are compared to  $\mathbf{x}$ .
- The most similar is chosen  $\rightarrow$  BMU (Best Matching Unit).
- BMU becomes a representative of vector  $\mathbf{x}$ .



# SOM Neuron 1/2

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Evaluates the similarity of input vector  $\mathbf{x}$  and weight vector  $\mathbf{w}_i$ .

Similarity: i.e., Euclidean

The most similar neuron to a input vector is chosen (BMU):

$$j^* = \underset{i}{\operatorname{argmin}} \left\{ \|\mathbf{x} - \mathbf{w}_i\| \right\},$$

**SOM neuron is a representative of a cluster.**

# SOM Neuron 2/2

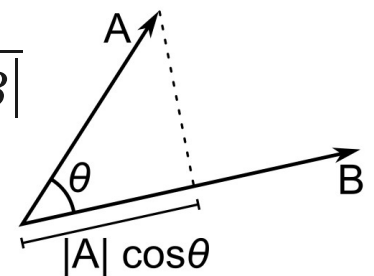
- Note, we don't have to use Euclidean distance.
- We can use directional similarity expressed by the dot product:

$$j^* = \arg \max_i \{ x^T(t) w_i(t) \} .$$

Why max here?

Note:

$$\cos \theta = \frac{A \cdot B}{|A||B|}$$



[http://en.wikipedia.org/wiki/Dot\\_product](http://en.wikipedia.org/wiki/Dot_product)

# Learning SOM

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- Initialization (random weights).
- Apply input pattern  $\mathbf{x} = (x_1, x_2, \dots, x_N)$ .
- Compute distances.
- Select BMU – neuron  $j$ .
- Adjust weights for all neurons  $i$ :

$$w_i(t+1) = w_i(t) + \eta_{ij}(t) [x(t) - w_i(t)]$$

- Continue with next pattern.

Neighbourhood function



# Example

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$$\mathbf{X} = \begin{bmatrix} 0.52 \\ 0.12 \end{bmatrix}$$

$$\mathbf{W}_1 = \begin{bmatrix} 0.27 \\ 0.81 \end{bmatrix}$$

$$\mathbf{W}_2 = \begin{bmatrix} 0.42 \\ 0.70 \end{bmatrix}$$

$$\mathbf{W}_3 = \begin{bmatrix} 0.43 \\ 0.21 \end{bmatrix}$$

$$d_1 = \sqrt{(x_1 - w_{11})^2 + (x_2 - w_{21})^2} = \sqrt{(0.52 - 0.27)^2 + (0.12 - 0.81)^2} = 0.73$$

$$d_2 = \sqrt{(x_1 - w_{12})^2 + (x_2 - w_{22})^2} = \sqrt{(0.52 - 0.42)^2 + (0.12 - 0.70)^2} = 0.59$$

$$d_3 = \sqrt{(x_1 - w_{13})^2 + (x_2 - w_{23})^2} = \sqrt{(0.52 - 0.43)^2 + (0.12 - 0.21)^2} = 0.13$$

The third vector is the winner (BMU).

# Example contd.

Let's move the neuron closer to the input pattern:  $w_{ij}(t+1) = w_{ij}(t) + \eta(t)[x_i(t) - w_{ij}(t)]$

$$\Delta w_{13} = \eta(t)(x_1 - w_{13}) = 0.1(0.52 - 0.43) = 0.01$$

$$\Delta w_{23} = \eta(t)(x_2 - w_{23}) = 0.1(0.12 - 0.21) = -0.01$$

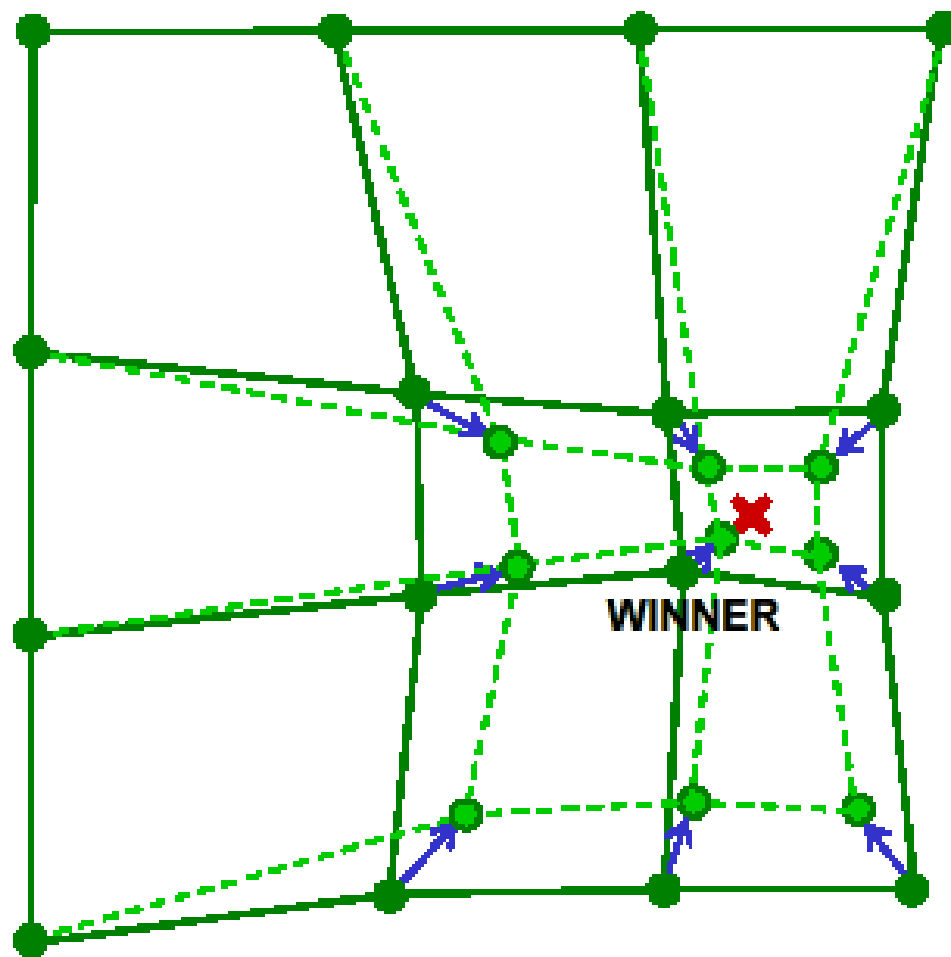
$$W_3(t+1) = W_3(t) + \Delta W_3(t) = \begin{bmatrix} 0.43 \\ 0.21 \end{bmatrix} + \begin{bmatrix} 0.01 \\ -0.01 \end{bmatrix} = \begin{bmatrix} 0.44 \\ 0.2 \end{bmatrix}$$

We adjusted only BMU weights.

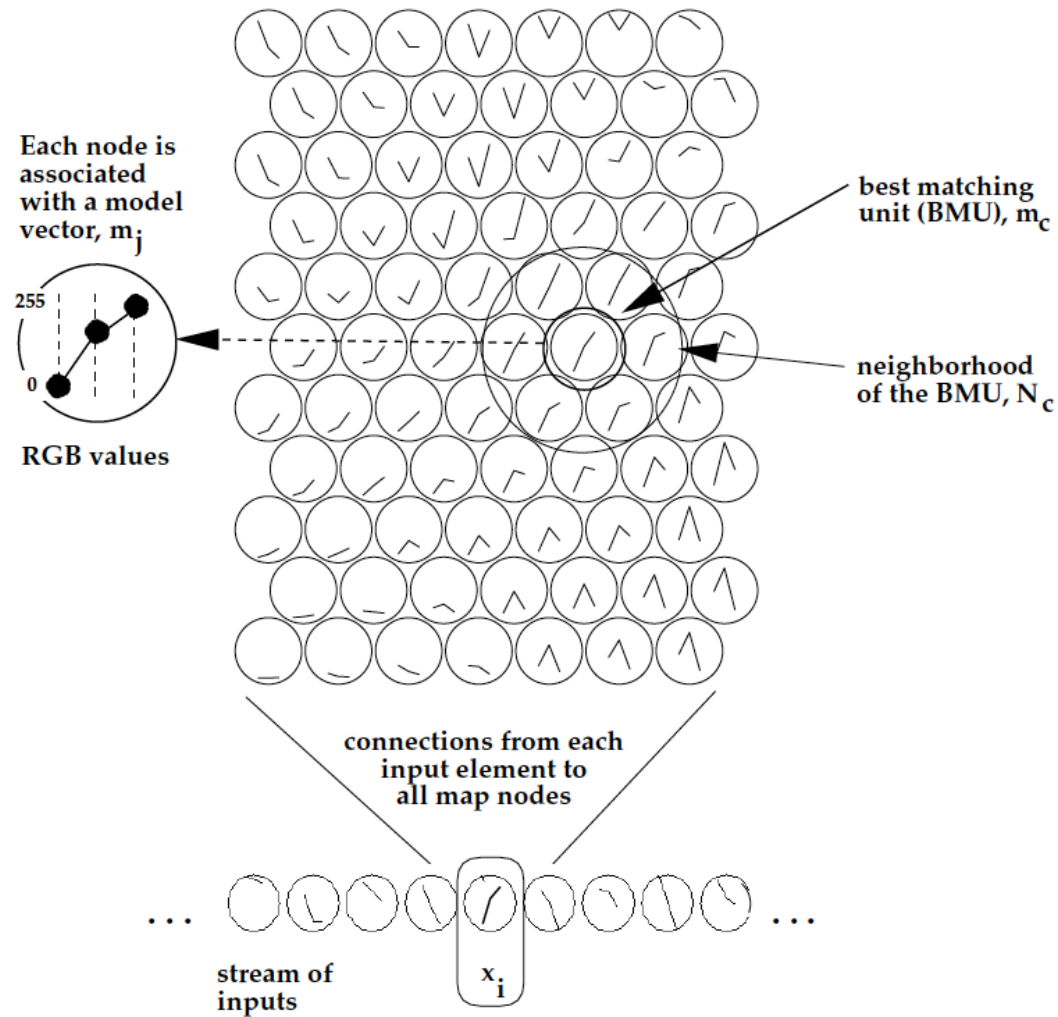
Here, the winner takes all.

# What About Updating Also Neurons in the Neighbourhood?

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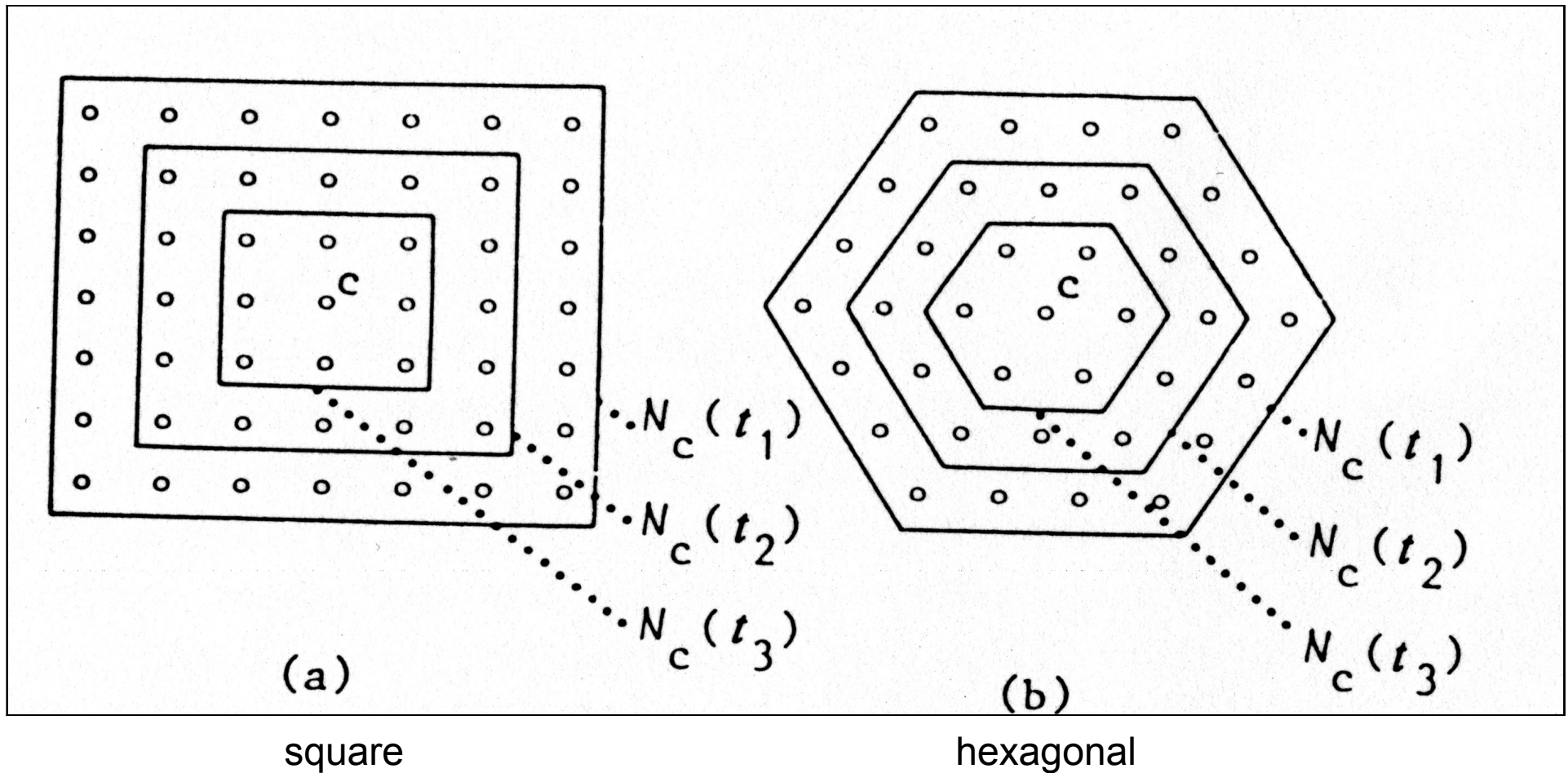


# Neighbourhood for SOM



Timo Honkela (*Description of Kohonen's Self-Organizing Map*)

# Common Neighbourhoods



*T. Kohonen: Self Organizing Maps*

# Learning SOM II

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- The neighbourhood plays important role when learning SOM:
  - topological arrangement,
  - neighbour distances.
- Neighbourhood changes in time:
  - its “diameter” decreases (to zero).
- The change is realised by neighbourhood function  $\eta(t)$ .



# Gaussian Neighbourhood

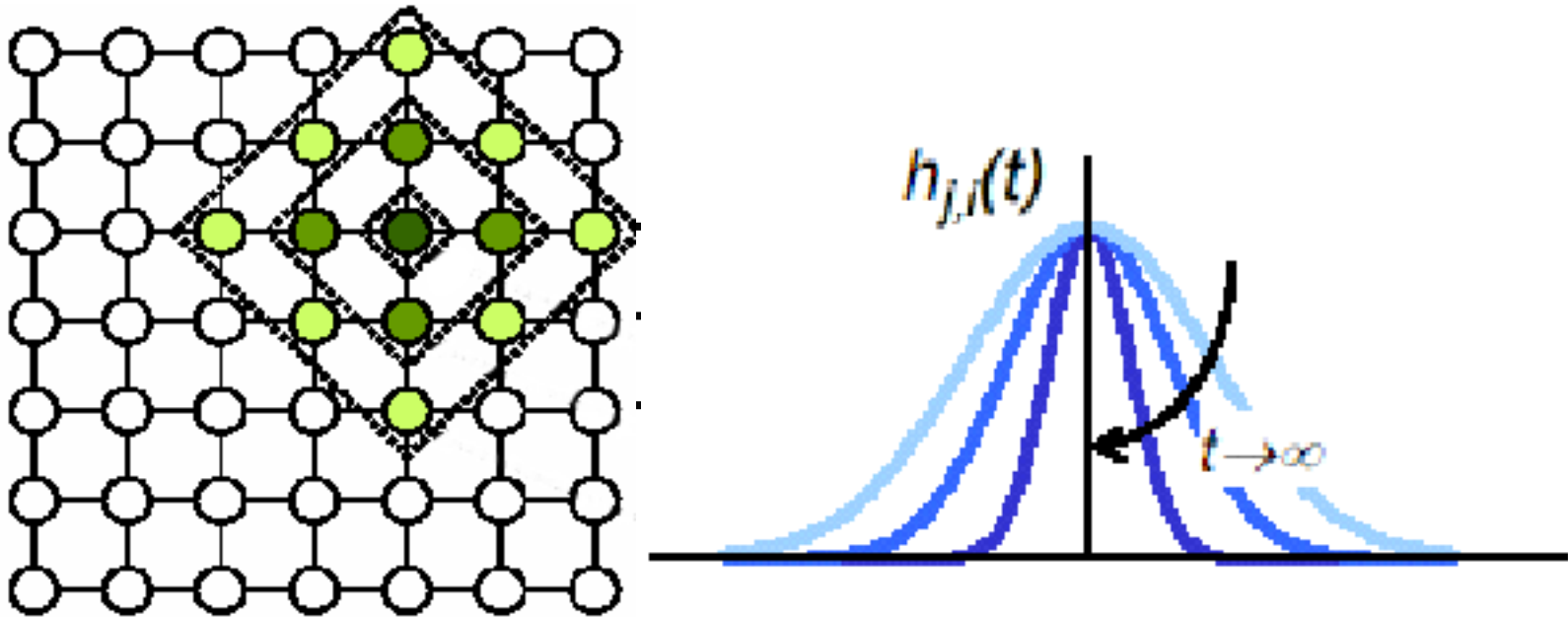
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- Neighbourhood function for neuron  $i$ .

$$\eta_{ij^*}(t) = \alpha(t) \cdot \exp\left(-\frac{\|r_{j^*} - r_i\|^2}{2\sigma^2(t)}\right)$$

- Where  $j^*$  is the BMU,  
 $r$  the position of neuron in map,  
and function  $\alpha(t)$ : learning rate.
- The *exp* expression represents neighbourhood shape.

# Gaussian Neighbourhood



Distance related learning

# Neighbourhood Related Functions

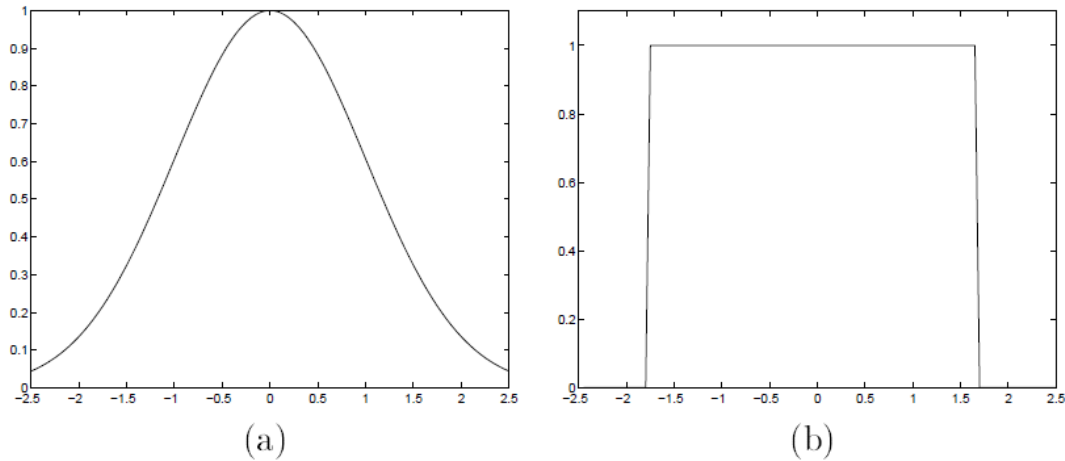


Figure 2.6: Neighborhood function values

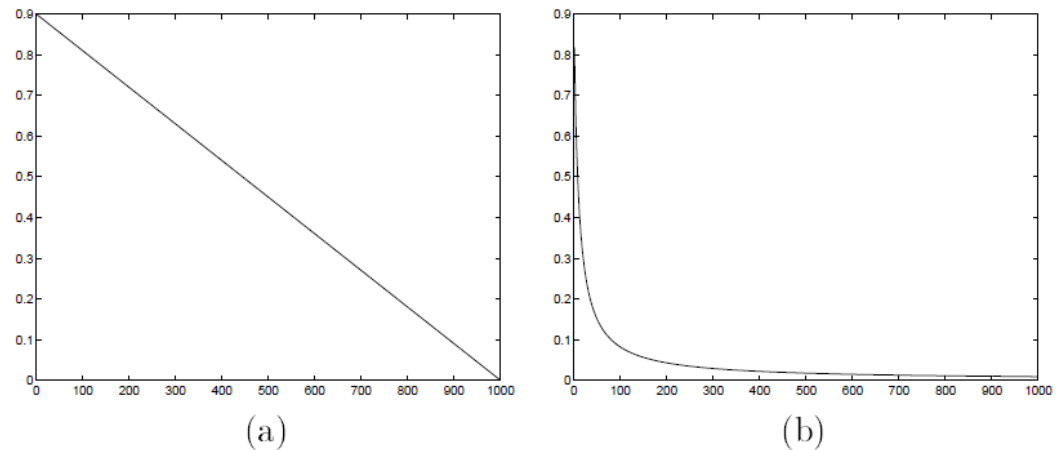
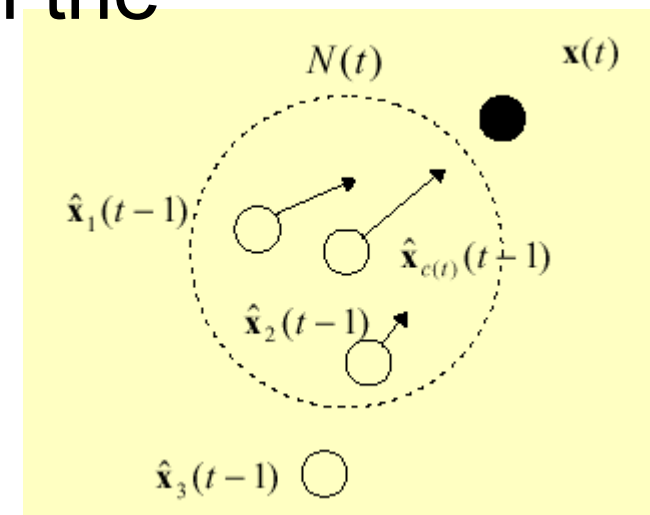


Figure 2.7: Learning rates as functions of time

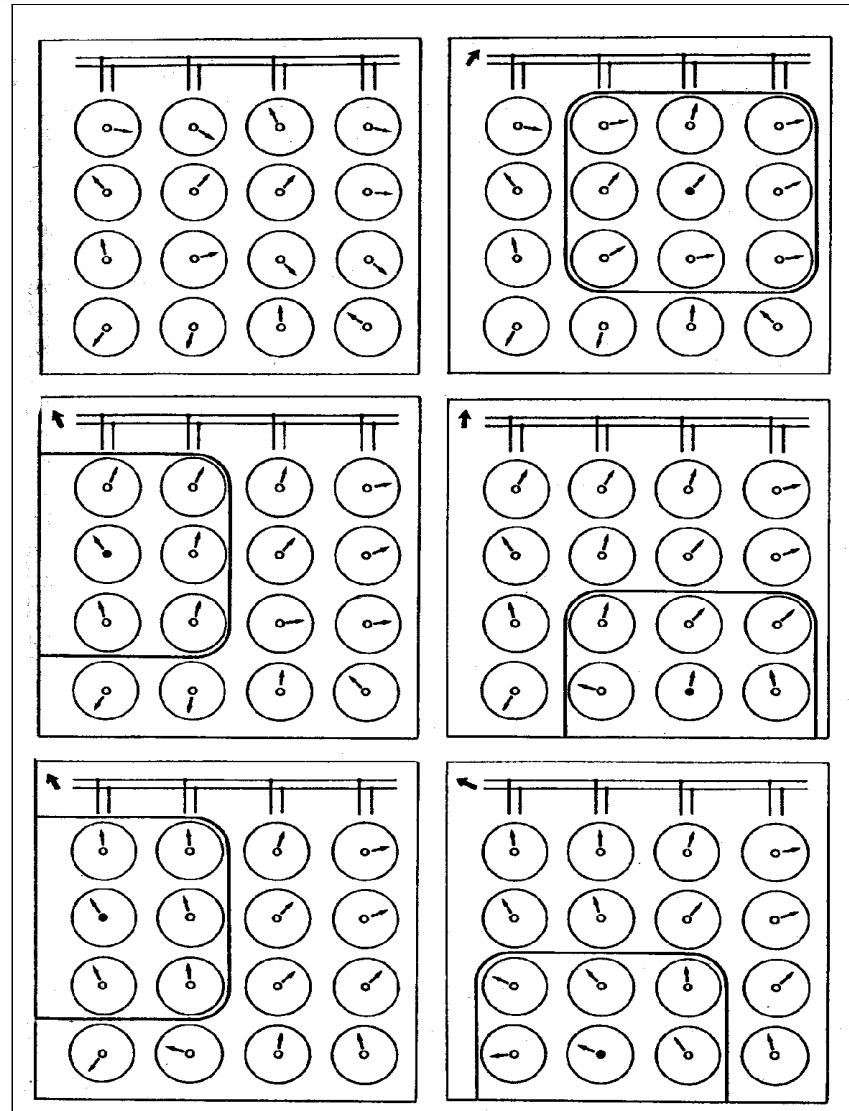
Hollmen '96, MSc.

# Learning Process

- During the learning the BMU (and its neighbours) is adapted to get closer to the input pattern which have caused its activation.
- Neurons are moving towards the input pattern.
- What influences the magnitude of the approach?



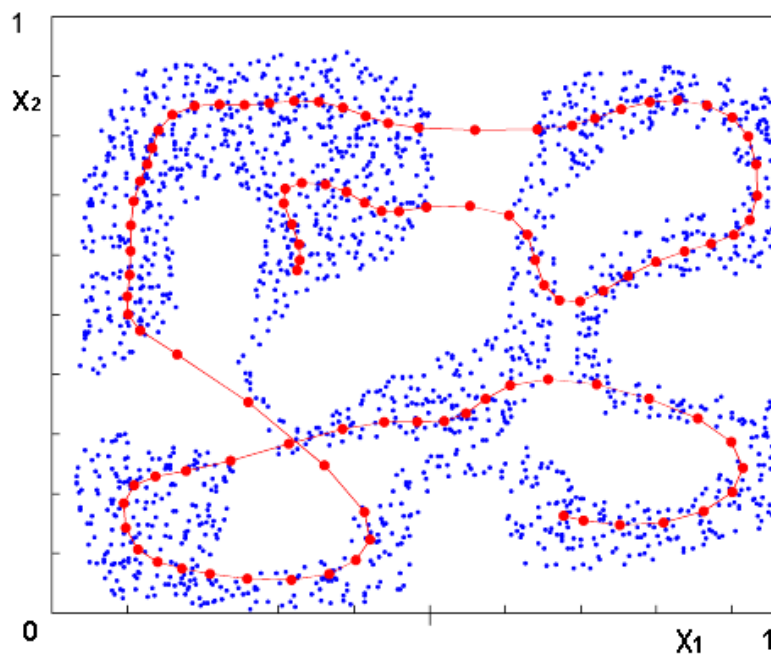
# Example: Learning Dot-Product SOM



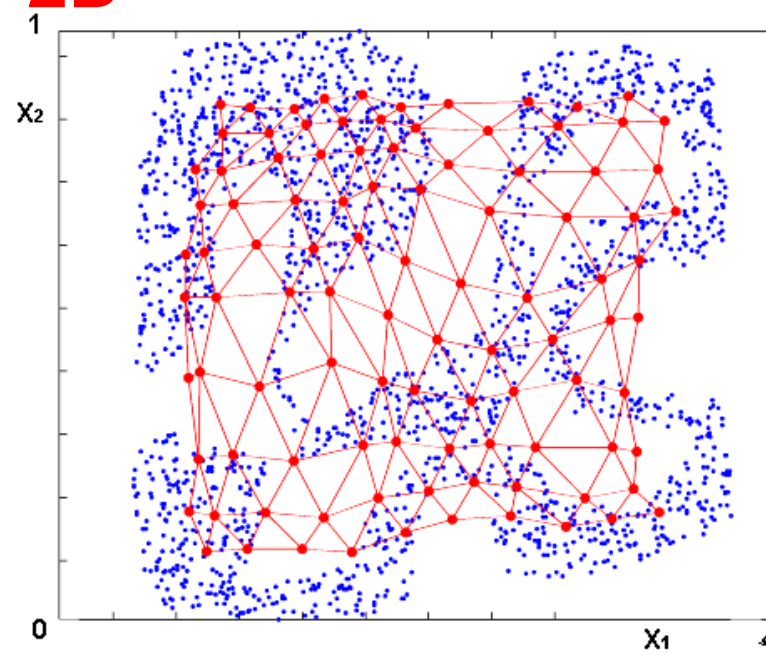
# SOM Applications

- To visualize data.
- To cover the input space by representatives.

**1D**



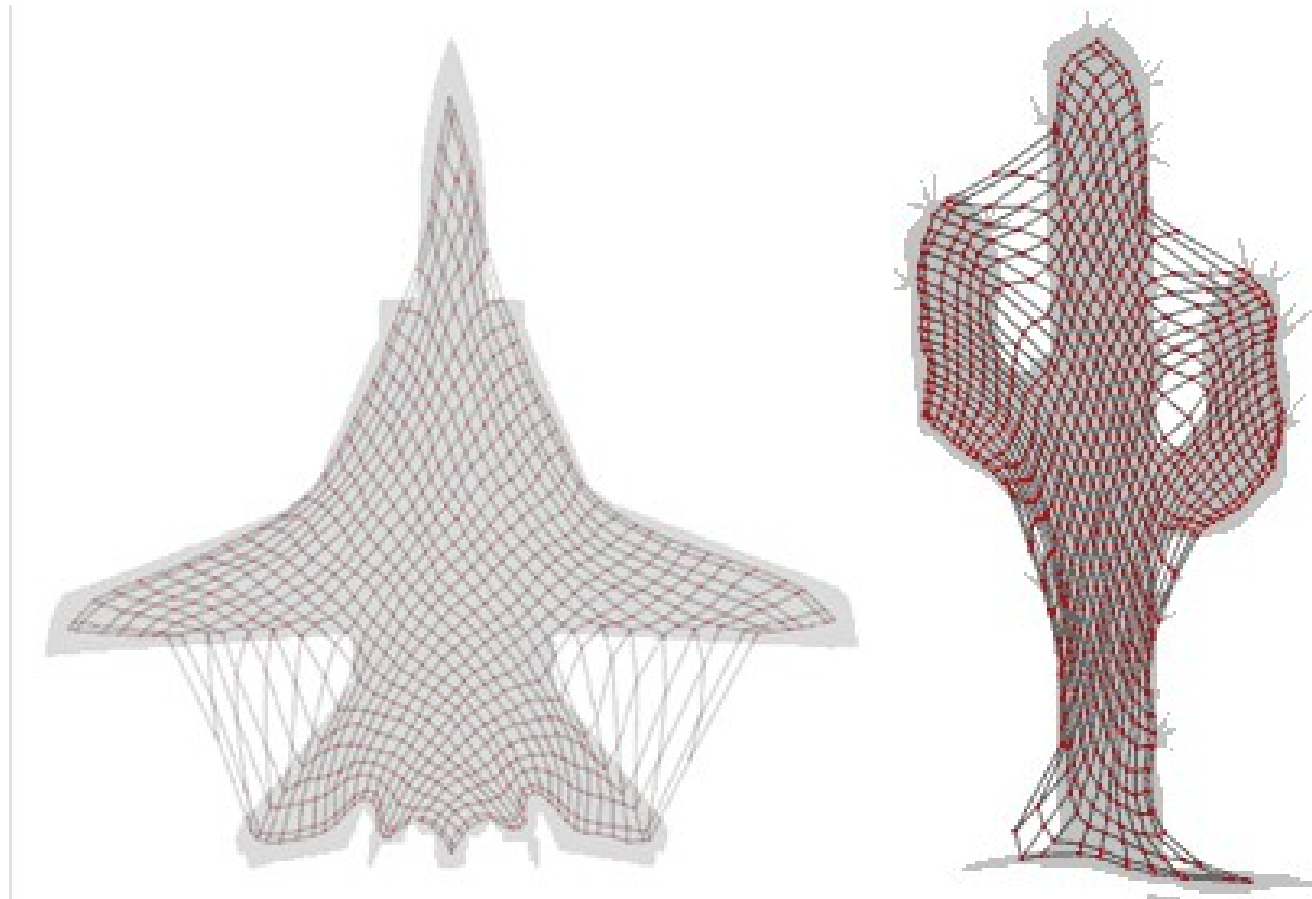
**2D**





# Or ...

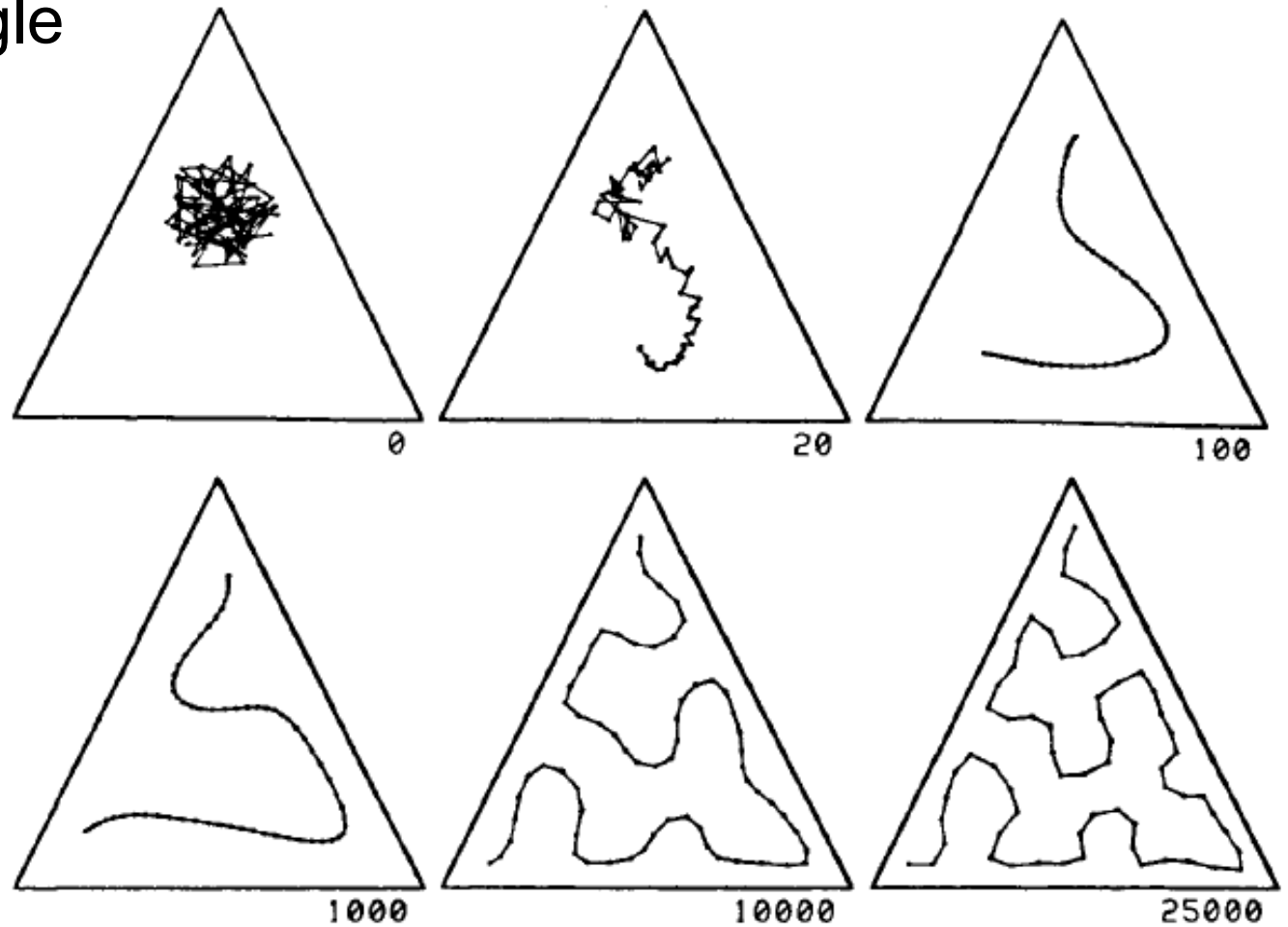
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Slide by Johan Everts

# More Examples

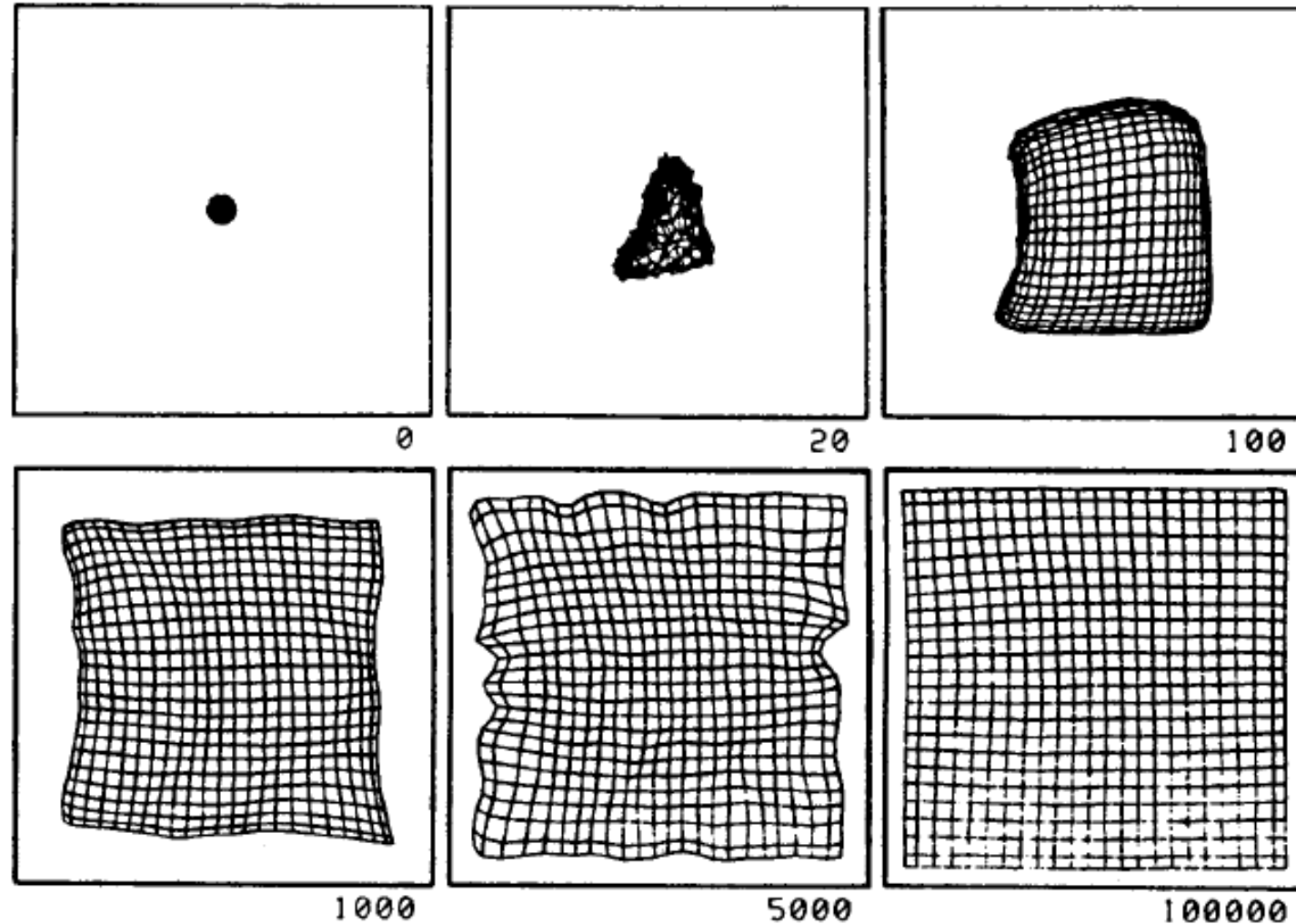
Covering a triangle by 1D SOM.



*T. Kohonen: Self Organizing Maps*

# More Examples contd.

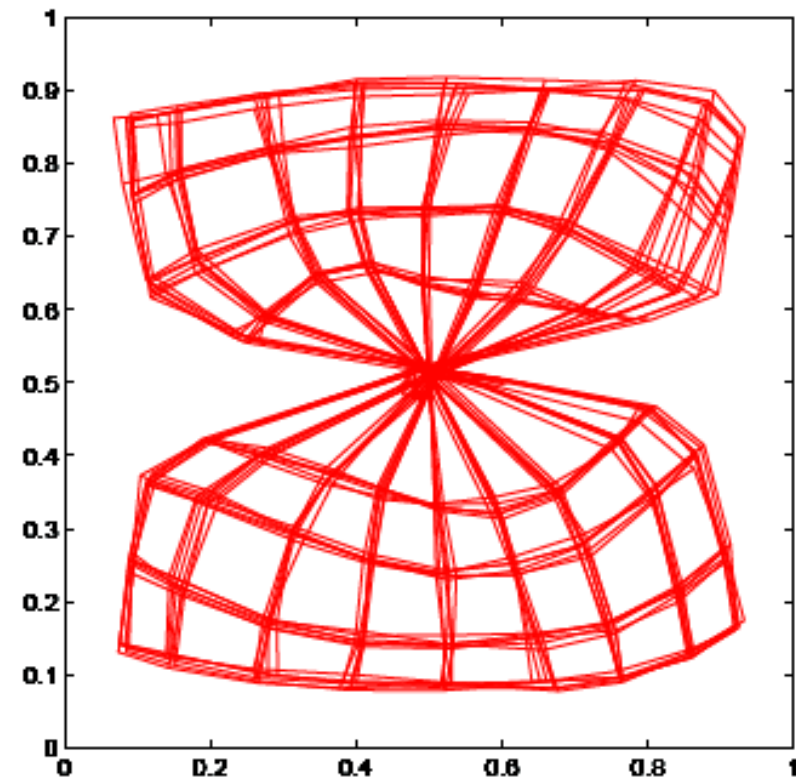
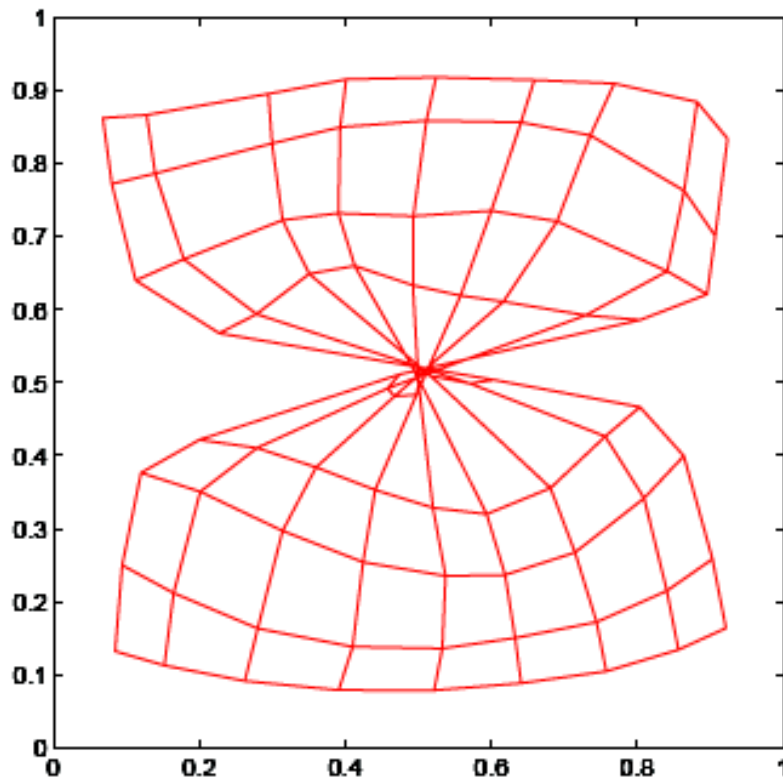
Covering a square by 2D SOM.



*T. Kohonen: Self Organizing Maps*

# Possible Problem: Knots

- This problem is not likely to be corrected by further learning if the *plasticity* is low:



Rojas: *Neural Networks - A Systematic Introduction*

# What is the Cause?

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- There are many:
  - Random initialization of weights → we are unable to change bad initial position/orientation of vectors.
  - Choice of a neighbourhood function.
  - Scheduling of neighbourhood modification in time.
  - Input data of course...

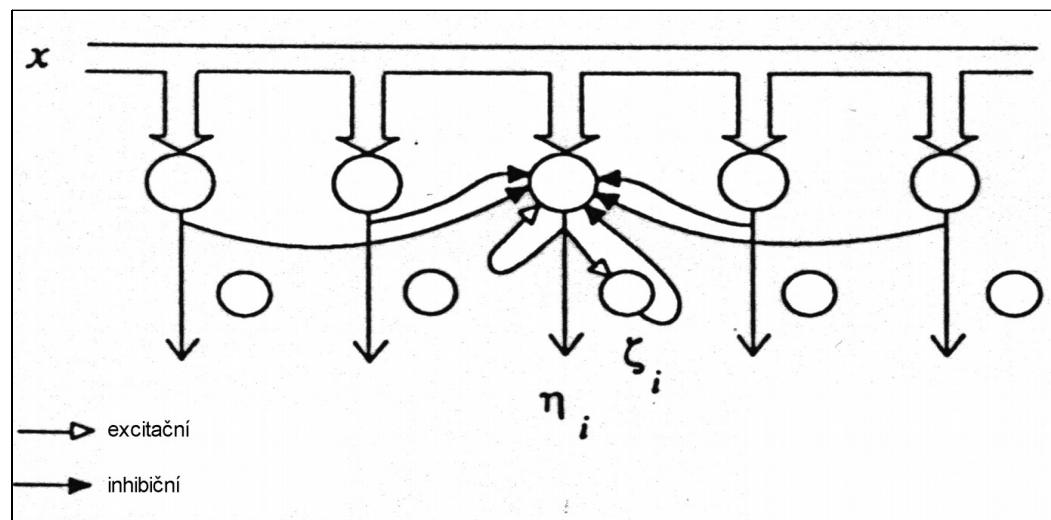
# What Can Help?

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- Same weights for all neurons initially → each neuron has a same chance to represent a pattern.
- Add random noise to input patterns at start.
- Lateral inhibition...

# Lateral Inhibition

- When choosing the BMU we do not pick isolated winner.
- The choice does not depend on an activation of a single neuron but also on activity of its neighbours...





# Lateral Inhibition II

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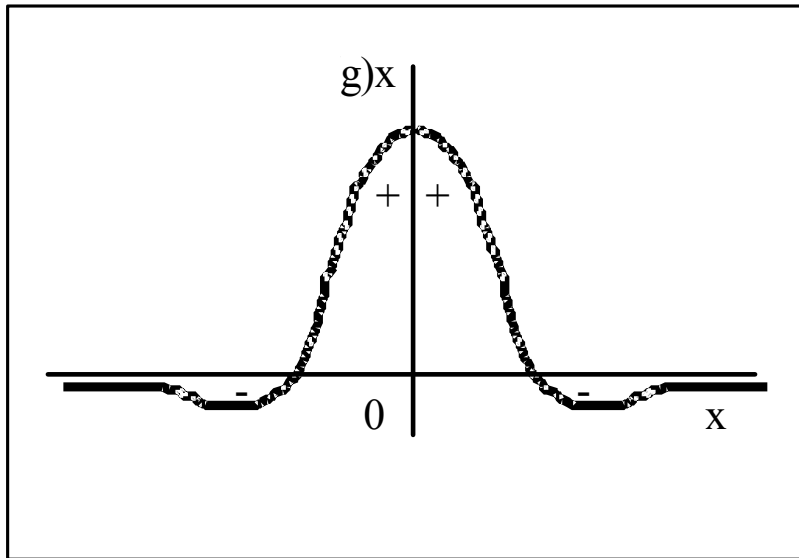
$$I_j = I_j^l + I_j^f = d_j + \sum_k g_{jk} I_k$$

Diagram illustrating the equation for the j-th neuron response,  $I_j$ , which is the sum of its local response,  $I_j^l$ , and its neighbourhood response,  $I_j^f$ . The equation is expanded as  $I_j = d_j + \sum_k g_{jk} I_k$ .

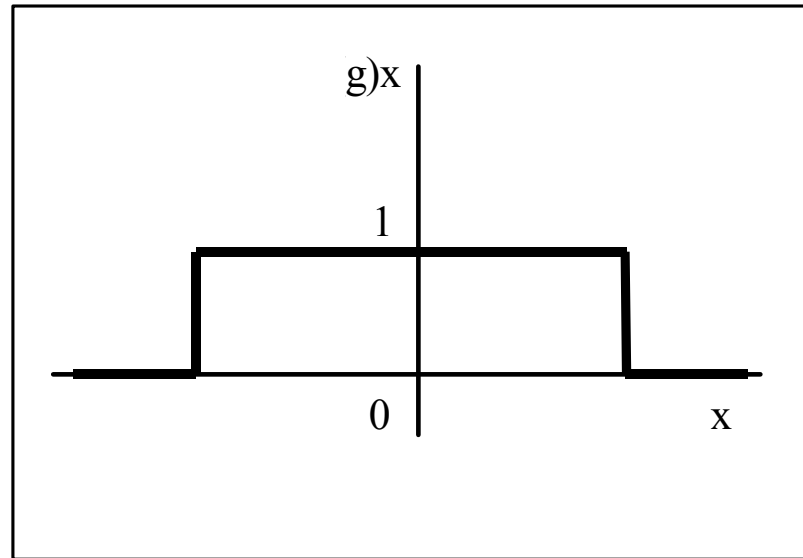
Labels and arrows pointing to the equation components:

- $I_j$ : j-th neuron response
- $I_j^l$ : local response
- $I_j^f$ : neighbourhood response
- $d_j$ : distance from input vector
- $\sum_k$ : neighbours
- $g_{jk}$ : lateral inhibition interaction

# Lateral Inhibition Functions



biological



simplified

# SOM Visualization

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- How to visualize representatives?
- Weight dimension = input vector dimension.
- How to show in 2D?
  - U-matrix,
  - P-matrix,
  - PCA (linear projection),
  - Sammon's projection (non-linear).

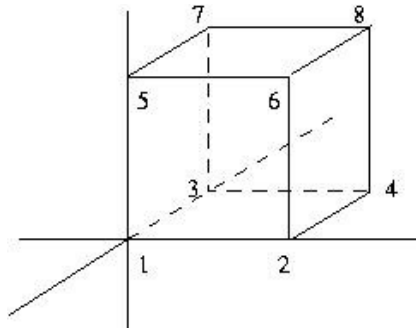
# U-matrix (UMAT)

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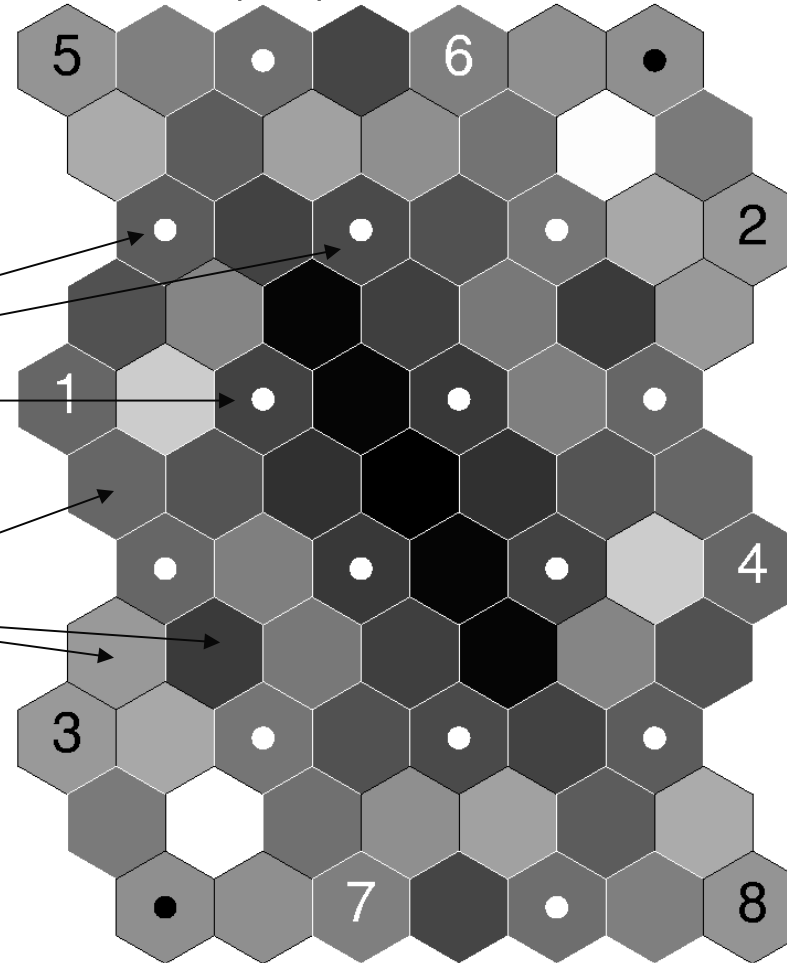
- Visualizes distances between neurons:
  - Dark coloring between neurons → large distance.
  - Light → close in input space.
- Dark gaps separate clusters.
- Neuron colour reflects the distance of its weight vector to all other weight vectors, again:
  - dark → large distance,
  - light → close distance.

# U-matrix Example

data



cube.cod - Dim: 3, Size: 4\*6 units, gaussian neighborhood



neurons

distance between adjacent neurons

# P-matrix (Pareto Density Estimation)

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- Shows the number of input space vectors which belong to a sphere centered in the neuron's weight vector.
- Visualizes data density.
- Neurons with high value belong to “dense” areas of input space.
- Neurons with low value are “lonesome”.
- Valleys separate clusters (“plateaus”).



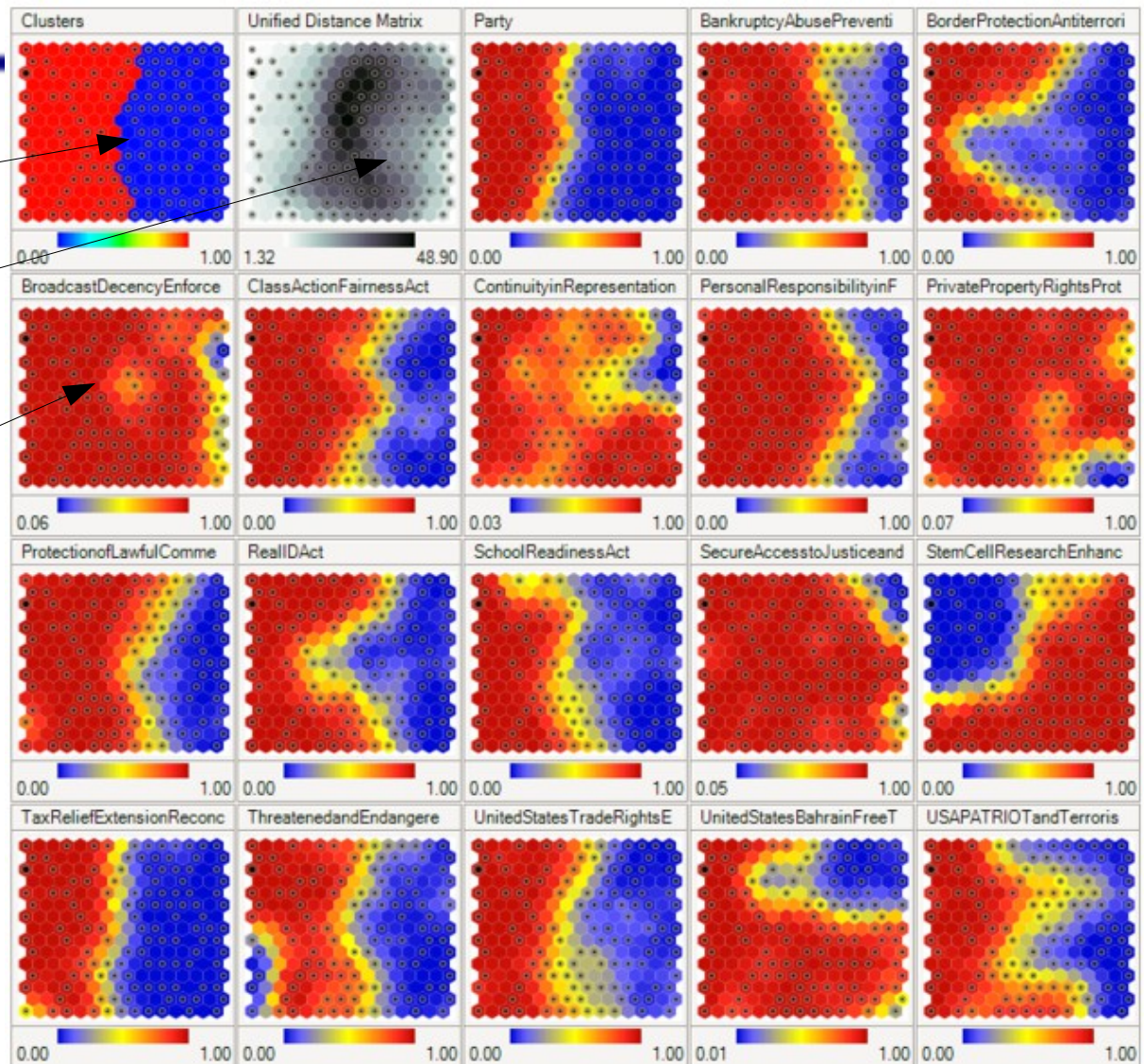
# Feature Plots

clustering

UMAP

**feature plot**  
shows a value  
of a single  
component (feature)  
of a weight vector

can be used to  
check if two  
components  
correlate



[http://en.wikipedia.org/wiki/Self-organizing\\_map](http://en.wikipedia.org/wiki/Self-organizing_map)



# Drawbacks of UMAT, PMAT

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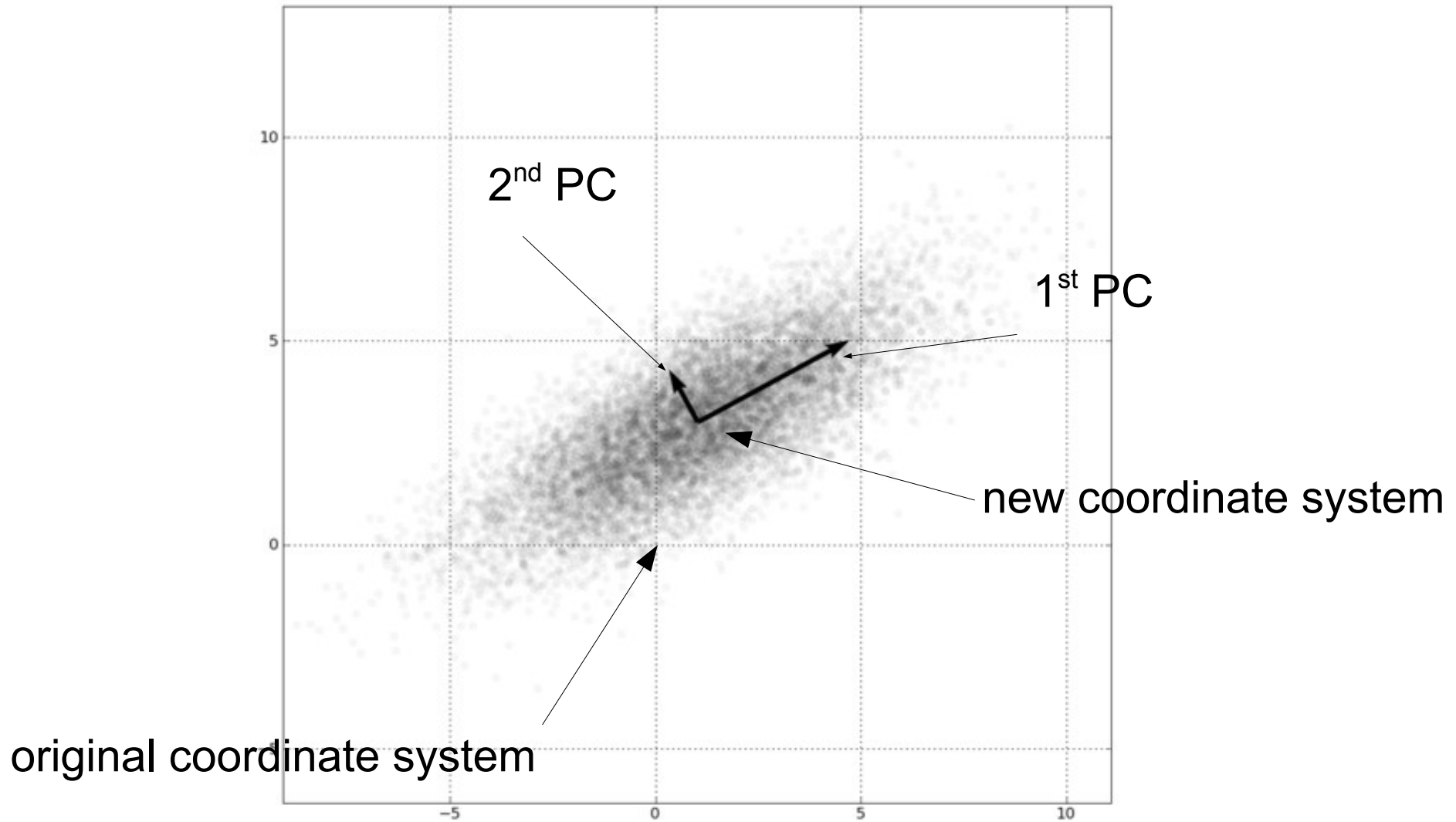
- Only distances between neighbours.
- New learning on the same data may give different results: (i.e. 90 degrees rotation)
- Not intuitive.
- **How can we show high-dimensional data in 2D(3D) keeping notion of original distances?**

# PCA

---

- Principal Component Analysis.
- Linear transformation to a new coordinate system such that:
  - 1<sup>st</sup> coordinate (principal component) → greatest variance by any projection of the data
  - 2<sup>nd</sup> coordinate → 2<sup>nd</sup> greatest variance
  - etc.
- Dimension reduction → use only  $N$  first coordinates, **throw the rest away...**

# Principal Components Example



[http://en.wikipedia.org/wiki/Principal\\_component\\_analysis](http://en.wikipedia.org/wiki/Principal_component_analysis)

# Sammon's Projection

- Non-linear reduction of higher-dimensional space to lower-dimensional space.
- Tries to preserve distances.

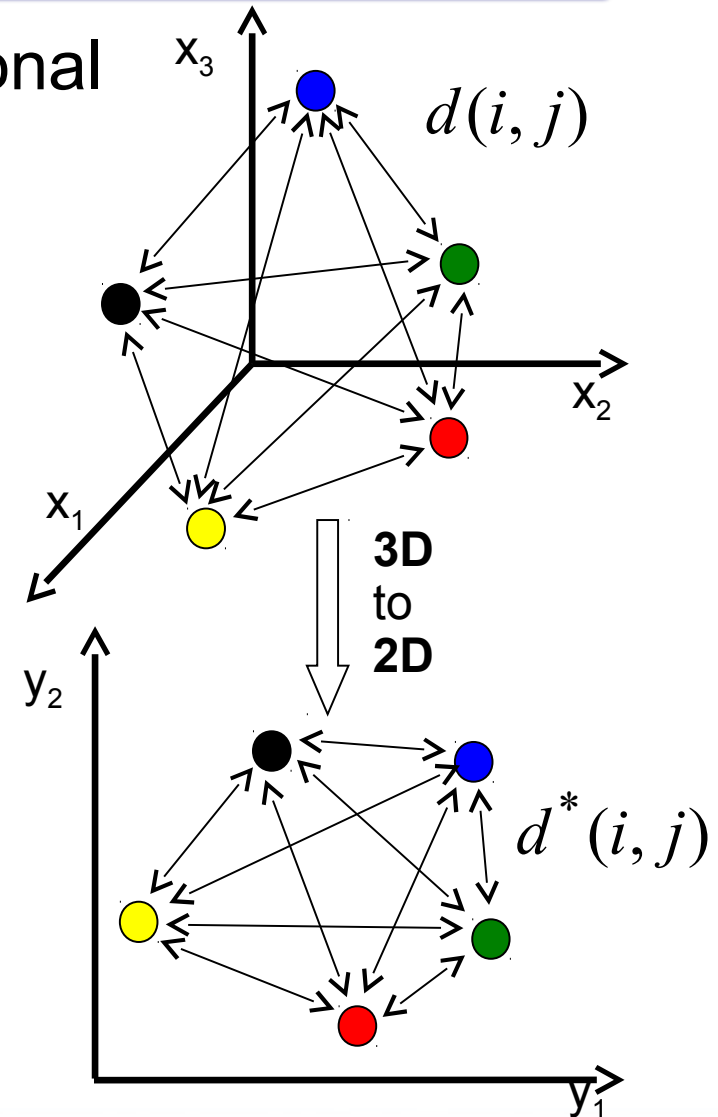
**energy function** →  
low for similar distances  
in both spaces.

distance in  
high dim. space

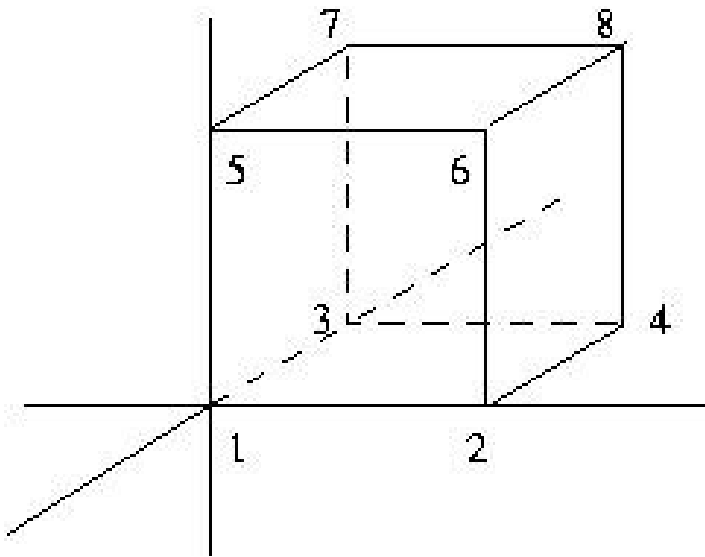
distance in  
low dim. space

$$E = \frac{1}{\sum_{i=1}^{N-1} \sum_{j=i+1}^N d(i, j)} \sum_{i=1}^{N-1} \sum_{j=i+1}^N \frac{(d(i, j) - d^*(i, j))^2}{d(i, j)}$$

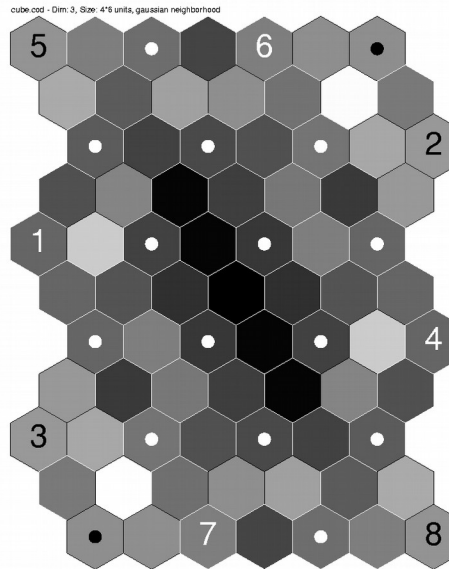
- Energy function is a subject to minimization (originally using gradient descent).



# Standard SOM Visualizations

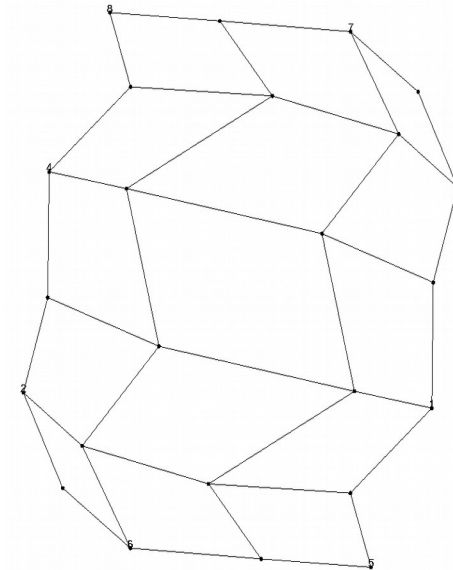


**UMAT**



**Sammon**

neuron weights projected to 2D,  
neighbours connected



# SOM Applications

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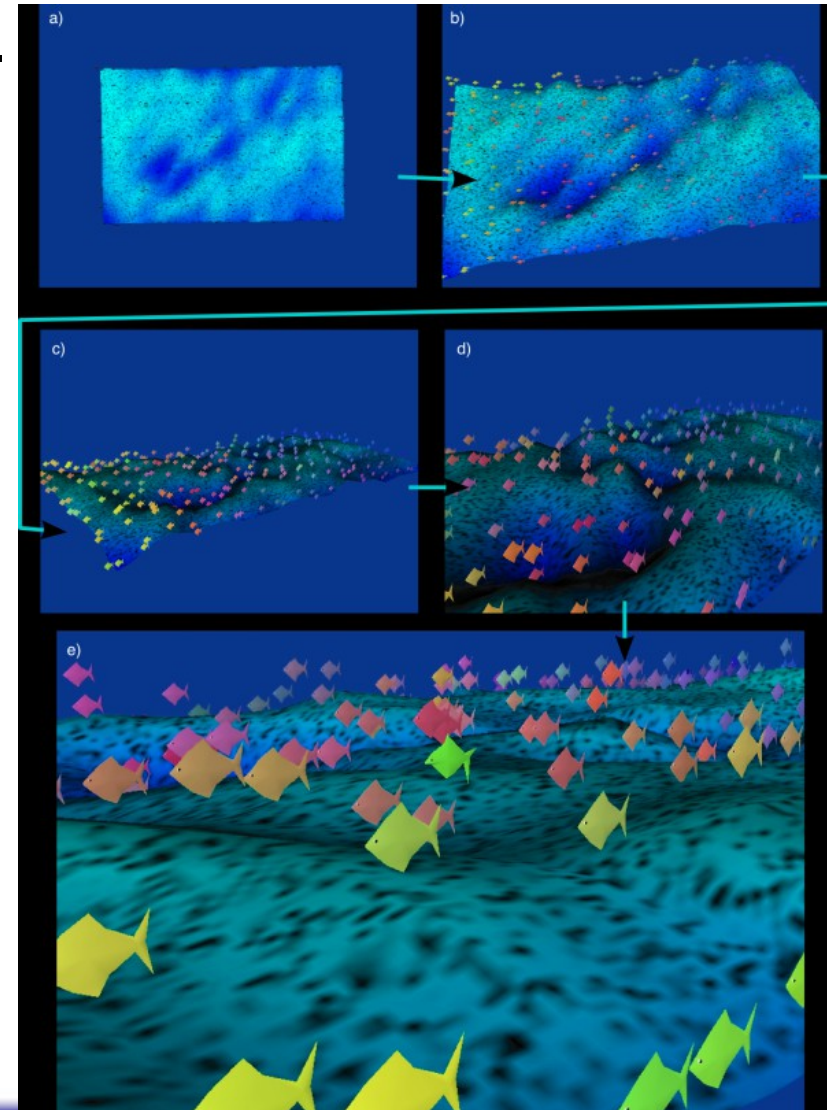
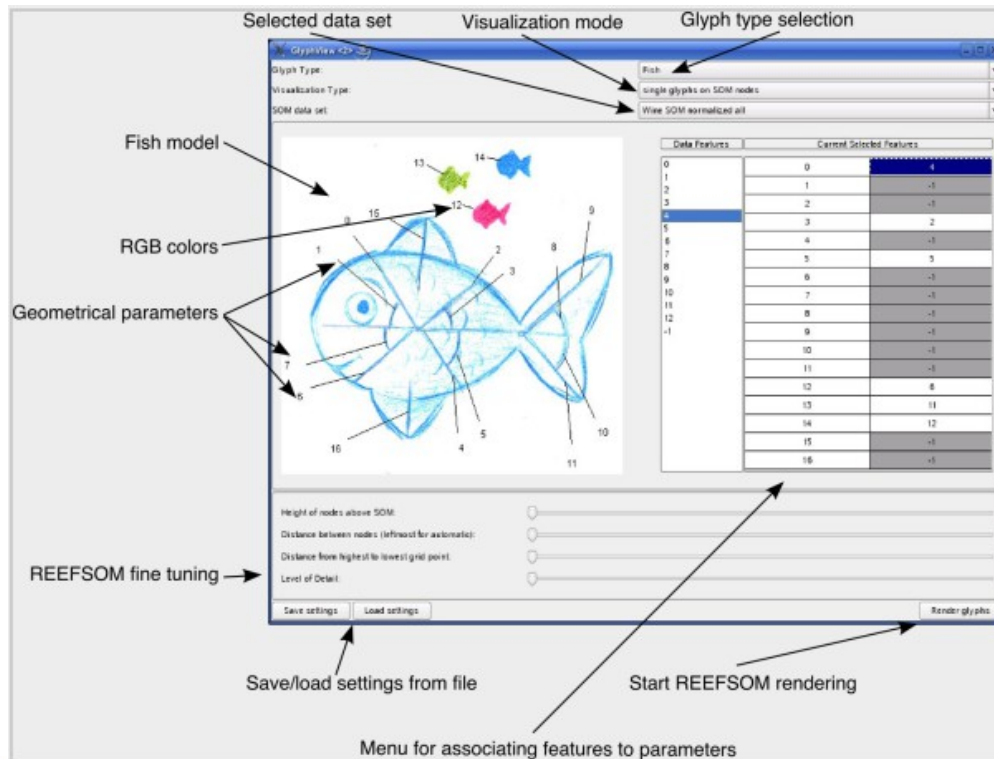
- Detection of similar images.
- <http://www.generation5.org/content/2007/kohonenImage.asp>





# ReefSOM

- SOM visualization for non-experts.
- UMAT + glyphs.
- <http://www.brains-minds-media.org/archive/305>





# SOM Evaluation

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- VQ – vector quantization, more input vectors mapped into a single neuron → **quantization error or distortion.**
- Compression of an input space dimension.
- Preserves data topology – neighbour vectors (from an input space) are mapped to neighbour neurons (in the mesh) → **topographic error.**

# SOM Quantization Error & Distortion

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- Quantization Error → average distance between input vector and its BMU (computed over all input vectors).
  - precision of mapping.

- Distortion → count with neighbours:

$$E = \sum_{i \in N} \sum_{j \in I} \eta_{i, bmu(j)} \|w(i) - x(j)\|^2$$

neurons

input  
vectors

Energy function again!

# Topographic Error of SOM

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- # of input vectors, for which the winner (BMU) and the second best neuron are not adjacent in the mesh.

# Next Lecture

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- ANNs + EAs = Neuro-evolution.