- Multi-layer perceptrons (MLP) constitute the most famous and most employed type of neural networks [1].
- At first, they were introduced in the context of classification, and it was soon recognized that they had the property of universal approximation capability.
- The introduction of the back-propagation algorithm made them flexible enough for many applications.





• Their basic structure for prediction is:





• The output of the ith neuron in the intermediate (hidden) layer is:

$$y_i = f\left(\theta_j + \sum_{j=1}^p w_{ij} x_{n-j}\right)$$

- with *f*(.) a sigmoidal function such as *f*(u) = tanh(*au*).
- The final output is a weighted sum of the outputs in the hidden layer.





• As for the polynomial predictor, one considers a NAR formulation:

$$x_n = g(x_{n-1}, \dots, x_{n-p}) + \mathcal{E}_n$$

and one estimates g(.) in the least-square sense using the data at hand. That is, one uses the universal approximation capability of MLPs.

• Of course, the input vector to predict  $x_n$  is

 $\boldsymbol{x}_{n} = [x_{n-1}, x_{n-2}, \dots, x_{n-p}].$ 



- It is possible to show that, when *p* increases, the size of the MLP increases more slowly than that of a polynomial predictor. But:
  - It is almost impossible to relate the values of the MLP parameters to the characteristic of the predicted signal (black box effect).
  - Convergence of the back-propagation algorithm is sometimes problematic.
  - Adaptive prediction is not very efficient.





• <u>Example</u>: prediction of RR intervals (p = 3, 5 neurons in the hidden layer). ESR: - 16 dB







• Generalization: prediction on another part of the signal. ESR = -13 dB







- Radial basis functions (RBF) constitute another type of neural networks, issued from interpolation theory.
- By some aspects, RBF combine the advantages of polynomial models and MLP:
  - Universal approximation property,
  - Estimation of parameters is simple,
  - There is a physical interpretation of these parameters.





- Let us suppose we have N pairs {x<sub>i</sub>, y<sub>i</sub>=g(x<sub>i</sub>)}, i = 1, ..., N, with x<sub>i</sub> p-dimensional vectors, y<sub>i</sub> scalars, and g(.) a continuous function.
- the *interpolation* of g(.) from these N pairs is an *ill-posed* one, because there is usually no information on g(.) between these pairs.
- A sensible approach, called *regularization*, has been developed to tackle this type of problem.





In this approach, one hypothesizes that g(.) should not oscillate erraticaly, and should present some degree of smoothness. This translates into defining an interpolating function G(.) minimizing a composite criterion:

$$C(G) = \sum_{i=1}^{N} [y_i - G(\mathbf{x}_i)]^2 + \rho \| \mathbf{P}G \|^2$$

with P a differential operator.





- Parameter  $\rho$  balances the relative influence of the two terms. The first one deals with the fidelity with respect to the data, the second with interpolation smoothness.
- It is possible to show that, if P is an infinite sum of differential operators of increasing degree, and rotation and translation invariant, then *G*(.) is:

$$G(\mathbf{x}) = \sum_{i=1}^{N} w_i \phi(\|\mathbf{x} - \mathbf{x}_i\|) \quad \text{with } \phi(s) = \exp\left(-\frac{s^2}{2\beta^2}\right)$$



• Parameter  $\beta$  must be specified with respect to the regularization parameter  $\rho$ . If there is a true interpolation then one must have:

$$G(\mathbf{x}_i) = y_i, i = 1, ..., N$$

and one must solve the following linear system:

$$\begin{bmatrix} \phi(\|\boldsymbol{x}_1 - \boldsymbol{x}_1\|) & \cdots & \phi(\|\boldsymbol{x}_1 - \boldsymbol{x}_N\|) \\ \vdots & \ddots & \vdots \\ \phi(\|\boldsymbol{x}_N - \boldsymbol{x}_1\|) & \cdots & \phi(\|\boldsymbol{x}_N - \boldsymbol{x}_N\|) \end{bmatrix} \begin{bmatrix} w_1 \\ \vdots \\ w_N \end{bmatrix} = \begin{bmatrix} y_1 \\ \vdots \\ y_N \end{bmatrix}$$



Thus a matrix equation:

$$\mathbf{A}\boldsymbol{w}=\boldsymbol{y}$$

- It is possible to show that if all vectors {x<sub>i</sub>} are different, matrix A is always invertible, and it will always be possible to determine the coefficients {w<sub>i</sub>} of w.
- The value of β defines the behavior of G(.) between the {x<sub>i</sub>}. The larger it is, the smoother G(.) is.







True function (not known).







- This approach of interpolation presents some limitations:
  - The fact that the interpolating function must include all pairs  $\{x_i, y_i = g(x_i)\}$  makes the approach highly sensitive to noise (erroneous values) and thus prone to overfitting.
  - When the number of pairs increases G(.) becomes soon complex.



ÉCOLE POLYTECHNIQUE FÉDÉRALE DE LAUSANNE • The idea that has been proposed to create a new type of neural networks is quite simple. One considers a slightly different formulation:

$$G(\mathbf{x}) = \sum_{j=1}^{M} w_j \phi \left( \|\mathbf{x} - \mathbf{c}_j\| \right) \quad \text{avec } \phi(s) = \exp \left( -\frac{s^2}{2\beta^2} \right)$$

with M < N, and vectors  $\{c_j\}$  called *centers*, which are not constrained to be a subset of the  $\{x_i\}$ . Function G(.) will now *approximate* g(.), typically in the leastsquare sense.





- To sum up, the RBF network approximates locally *g*(.) in the neighborhood of each center, and merges these local approximations to create a global one.
- The nice feature of RBF networks is that, once the centers and β have been defined, *determination of coefficients* {w<sub>j</sub>} *is simple, since* G(.) *depends linearly on them.*





• If least-square estimation is used, one must solve:

$$\begin{bmatrix} \phi(\|\mathbf{x}_1 - \mathbf{c}_1\|) & \cdots & \phi(\|\mathbf{x}_1 - \mathbf{c}_M\|) \\ \phi(\|\mathbf{x}_2 - \mathbf{c}_1\|) & \cdots & \phi(\|\mathbf{x}_2 - \mathbf{c}_M\|) \\ \vdots & & \vdots \\ \phi(\|\mathbf{x}_N - \mathbf{c}_1\|) & \cdots & \phi(\|\mathbf{x}_N - \mathbf{c}_M\|) \end{bmatrix} \begin{bmatrix} w_1 \\ \vdots \\ w_M \end{bmatrix} = \begin{bmatrix} y_1 \\ \vdots \\ y_N \end{bmatrix} + \begin{bmatrix} e_1 \\ \vdots \\ e_N \end{bmatrix}$$

that is  $\mathbf{A}\mathbf{w} = \mathbf{y} + e$ . A is now an  $N \times M$  matrix.

• One may add a column of "1" to A if y is significantly non zero-mean.





- When minimizing ||Aw y||, one performs of course an orthogonal projection of vector y on the subspace generated by the columns of A.
- Since matrix A may be ill-conditioned it is better to use a robust estimation scheme, typically SVD.
- But of course centers should not be too close to each other.





- Note first there is no really optimal way to select the centers. Also, we will come back later to the problem of determining the *number* of centers.
- One of the firts methods proposed consisted in selecting centers randomly (uniformly), in the hypercube defined by the {x<sub>i</sub>}. This is not very efficient, because the spatial distribution of the {x<sub>i</sub>} is not taken into account.





• A second approach consists in selecting randomly the centers among the  $\{x_i\}$ . This is already better, since statistically those centers are representative of the spatial distribution of the  $\{x_i\}$ . That is, there are more centers in the regions of space where the  $\{x_i\}$  are clustered. However, the constraint remains that the centers are a subset of the  $\{x_i\}$ .





- A solution yielding better performance is to select centers using *learning vector quantization* (LVQ), which is used in other contexts too to find representatives of vector sets. LVQ works as follows:
- 1. The initial centers are chosen  $\{c_j\}$  at random in the hypercube defined by the  $\{x_i\}$ .





2. Vectors  $\{x_i\}$  are presented, generally for several epochs. At iteration *k*, the following correction si applied to center  $c_j$ :

$$\mathbf{c}_{j}(k+1) = \mathbf{c}_{j}(k) + \alpha(k) [\mathbf{x}(k) - \mathbf{c}_{j}(k)]$$
  
if  $\|\mathbf{x}(k) - \mathbf{c}_{j}(k)\| < \|\mathbf{x}(k) - \mathbf{c}_{m}(k)\|, \ m \neq j$ 

with x(k) the vector presented at iteration k. the net effect is to draw  $c_j$  closer to x(k), and thus possibly to a cluster.





Parameter  $\alpha(k)$  must decrease with *k*, generally using:

$$\alpha(k) = \alpha_0 \left( 1 - \frac{k}{K} \right)$$

- *K* total number of iterations, and  $\alpha_0$  a constant (typically  $\alpha_0 = 0.05$ ).
- It is to be noted that LVQ is sensitive to the initial center selection.



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• <u>Example:</u> LVQ for M = 5 centers, vectors  $\{x_i\}$  drawn from two bi-dimensional Gaussian pdf.





- Another scheme, less sensitive to initial center choice, is *Lloyd-Max vector quantization algorithm*.
- Centers {c<sub>j</sub>} (sometimes called in this context *centroids*) are chosen such that they minimize a distortion criterion:

$$D = \sum_{j=1}^{M} \sum_{\mathbf{x}_i \in V_j} \|\mathbf{x}_i - \mathbf{c}_j\|^2$$





- Subsets V<sub>j</sub> contain vectors x<sub>j</sub> such that c<sub>j</sub> is the center closest to them. So to speak, c<sub>j</sub> must be a good representative of V<sub>j</sub>. Lloyd-Max algorithm works as follows
- Initial random selection of {*c<sub>j</sub>*} among the {*x<sub>i</sub>*}
  Update:

$$\boldsymbol{c}_j \rightarrow \boldsymbol{c}_j = \operatorname{mean}(\boldsymbol{x}_i \in V_j)$$

3. If distortion *D* stabilizes, stop, otherwise back to 2.



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<u>Example</u>: Lloyd-Max for *M* = 5 centers, vectors {*x<sub>i</sub>*} drawn from two bi-dimensional Gaussian pdf.

- data
- centroids







- Once the centers {c<sub>j</sub>} have been chosen, parameter β can be determined.
- This is the same as for interpolation: too large a β means too strong an interaction (overlap) between the local approximations. Too small a β means a "bumpy" approximation.
- A good empirical rule is:

$$\beta = \frac{d}{\sqrt{M}}$$
 with  $d = \max \|\mathbf{c}_i - \mathbf{c}_j\|$ ,  $1 \le i, j \le M$ 



- Nothing imposes to use the same value for  $\beta$  for each RBF.
- It is even intuitevly appealing that  $\beta$  be smaller in regions with many centers, since the spatial influence of the RBFs corresponding to these centers should be smaller
- A good empirical rule for  $\beta_i$  corresponding to center  $c_i$ is:







• As for MLP, one considers a NAR formulation:

$$x_n = g(x_{n-1}, \dots, x_{n-p}) + \mathcal{E}_n$$

- and one estimates g(.) in the least-square sense using the data at hand. That is, one uses the universal approximation capability of RBF networks.
- Of course, the input vector to predict  $x_n$  is

$$\boldsymbol{x}_{n} = [x_{n-1}, x_{n-2}, \dots, x_{n-p}].$$



• <u>Example:</u> prediction of RR intervals (p = 3, 20RBF, multiple  $\beta$ ). ESR: - 10 dB







• Generalization: prediction on another part of the signal. ESR = -4 dB





- Center selection, which conditions the choice of parameter  $\beta$  also, has an important role in RBF network performance.
- LVQ and Lloyd-Max algorithms are intuitively appealing. There are however other interesting schemes [3].
- Again, we note that solving Aw = y + e in the least squares sense means performing an *orthogonal projection of y* on the linear subspace generated by the *columns of A*.





- One can apply the same procedure as for term selection in polynomial prediction, that is a Gram-Schmidt orthogonalization on selected columns (*orthogonal least squares*, OLS).
- At each iteration, one adds in A the column yielding maximum erro reduction. This column is defined by a pair (c<sub>j</sub>, β<sub>j</sub>), and center c<sub>j</sub> is selected among the vectors {x<sub>i</sub>}.





- One proceeds as follows:
- 1) The first center is selected by examining all the  $\{x_i\}$ , and keeping the smallest least squares error.
- 2) Successive centers are chosen among the remaining  $\{x_i\}$ , with an orthogonalization of  $[\phi(x_1), \ldots, \phi(x_N)]^{\mathsf{T}}$  with respect to the terms already selected. In this way the center giving maximum error reduction is spotted.





• <u>Example</u>: prediction of RR intervals (p = 3, 5 RBF, multiple  $\beta$ ). ESR: - 11 dB

Signal Prediction

• samples corresponding to selected *x*<sub>i</sub>





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LTS





LTS













$$MDL = N \cdot \ln \sigma_e^2 + M \cdot (\ln N + B \cdot \ln 2)$$

## B: Number of bits to encode one center

















- It is of course possible to derive an adaptive version of RBF network prediction.
- In what follows, reference to time index *k* will not be explicit.
- w(i) ith component of coefficient vector w
- $\phi(i)$  ith RBF
- **\overline RBF** vector
- instantaneous square error is thus

$$\varepsilon^2 = [y - \mathbf{G}(\mathbf{x})]^2 = [y - \mathbf{w}^{\mathsf{T}} \mathbf{\phi}]^2$$



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• In LMS, the updates are:

$$w \to w - \mu_1 \frac{\partial \varepsilon^2}{\partial w} = w - 2\mu_1 \varepsilon \frac{\partial \varepsilon}{\partial w} = w + 2\mu_1 \varepsilon \frac{\partial G(x)}{\partial w}$$
$$= w + 2\mu_1 \varepsilon \phi$$

$$\boldsymbol{c}_{j} \rightarrow \boldsymbol{c}_{j} + 2\mu_{2}\varepsilon \frac{\partial G(\boldsymbol{x})}{\partial \boldsymbol{c}_{j}} = \boldsymbol{c}_{j} + 2\mu_{2}\varepsilon \boldsymbol{w}(j)\boldsymbol{\phi}(j)(\boldsymbol{x} - \boldsymbol{c}_{j})$$

$$\beta \rightarrow \beta + 2\mu_3 \varepsilon \frac{\partial G(\mathbf{x})}{\partial \beta} = \beta + \frac{2\mu_3 \varepsilon}{\beta^3} \sum_{i=1}^M \|\mathbf{x} - \mathbf{c}_i\|^2 \phi(i)$$



 Example: prediction of RR intervals (p = 3, 5 RBF, unique β). ESR: - 8 dB







- Remarks:
  - Better thsn batch algorithm, but not impressive
  - Tuning of adaptation parameters is uneasy (except for *w* for which it is possible to normalize by  $\phi^{T}\phi$ )
- But maybe the gradient approach is not suited to center and β parameter update. For instance, nothing guarantees that the mean sugare error has a unique minimum.





- Why not *decouple* the update of *w*, for which LMS should work well, from that of the centers and β?
- One might use an LVQ-style update for the centers, then modify  $\beta$  by using the empirical rule based on the distance between centers.
- But it is necessary to define a link between the two update schemes.





1. One first computes the modification on  $c_i$ , the center closest to x with LVQ (but  $\alpha$  must remain constant to preserve adaptability):

 $c_i \rightarrow c_i + \Delta c_i = c_i + \alpha (x - c_i)$ 

- 2. If the maximum distance between centers changes,  $\beta$  should become  $\beta + \Delta\beta$ , by using the empirical rule for unique  $\beta$ .
- The output changes from G(x) to  $G(x) + \Delta G(x)$ .



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- But:  $\Delta G(x) \approx (\Delta c_i)^{\mathsf{T}} \frac{\partial G(x)}{\partial c_i} + \Delta \beta \frac{\partial G(x)}{\partial \beta}$
- Thus:

LTS

$$\frac{\partial (G(\boldsymbol{x}) + \Delta G(\boldsymbol{x}))}{\partial \boldsymbol{w}} \approx \frac{\partial G(\boldsymbol{x})}{\partial \boldsymbol{w}} + (\Delta \boldsymbol{c}_i)^{\mathsf{T}} \frac{\partial G(\boldsymbol{x})}{\partial \boldsymbol{w} \partial \boldsymbol{c}_i} + \Delta \beta \frac{\partial G(\boldsymbol{x})}{\partial \boldsymbol{w} \partial \beta}$$

• And the instantaneous square error becomes:

$$\varepsilon^2 = [y - G(\mathbf{x}) - \Delta G(\mathbf{x})]^2$$

The coefficient vector *w* is updated with:

$$w \rightarrow w - \mu_1 \frac{\partial \varepsilon^2}{\partial w} = w + 2\mu\varepsilon \frac{\partial (G(x) + \Delta G(x))}{\partial w}$$







• <u>Example:</u> prediction of RR intervals (p = 3, 5RBF, unique  $\beta$ ). ESR: - 10.7 dB







- Remarques:
  - Prediction performance improves in the nonstationary intervals.
  - It is a lot easier to tune  $\mu$  for *w* and  $\alpha$  for the centers.
  - This approach is easily modified for the multiple  $\beta$  case.





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