- Some criticisms may be addressed to the standard SETAR model:
 - Although there are situations where an abrupt change in regime takes place, nature usually does not produce discontinuties.
 - It may seem a bit limited that the change in regime be linked to only one lagged sample.
- This has triggered the development of variants to the SETAR which incorporate one (or both) remarks.



- A smoother transition between regimes can be obtained by replacing the indicator function by a continuous function $\phi(.)$ changing gradually from 0 to 1.
- One gets a Smooth Transition AR (STAR) model [1]. With z_n the transition variable, a popular choice for $\phi(.)$ is the logistic function:

$$\phi(z_n;\gamma,r) = \frac{1}{1 + \exp[-\gamma(z_n - r)]}$$



- A 2-regime model can be described by:
- $x_n = (a_{10} + a_{11}x_{n-1} + \dots + a_{1p_1}x_{n-p_1})(1 \phi(x_n; \gamma; r))$
- + $(a_{20} + a_{21}x_{n-1} + \dots + a_{2p}x_{n-p})\phi(x_n;\gamma;r) + \varepsilon_n$

with ε_n is an i.i.d. sequence.

• The parameter γ determines the steepness of the logistic function. The larger it is, the closer $\phi(.)$ is to as step function. When $\gamma = 0$, $\phi(.)$ is a constant.



- Since the STAR model is based on a continuous function, least squares estimation is a maximum likelihood one if the residuals are supposed to be Gaussian.
- Estimation of the sub-models coefficients is easy if γ and r are fixed, but full estimation requires either the use of an optimization algorithm or an exhaustive grid search.



• The idea is quite simple: it consists in defining a partition of the state space itself instead of a partition of the real line for a single threshold variable as for the SETAR:

$$x_{n} = \sum_{k=1}^{K} \{a_{k0} + a_{k1}x_{n-1} + \dots + a_{kp_{k}}x_{n-p_{k}} + \sigma_{k}\varepsilon_{n}\}I(x_{n-1})$$

with $x_{n-1} = [x_{n-1}, ..., x_{n-p}]^T$, *p* usually taken smaller than or equal to $\max[p_k]$.

 $\in A_k$





• The easiest way to define the partition $\{A_k\}$ is through a set of centers $\{c_k\}$ and the use of the Euclidian distance, that is:

$$\boldsymbol{x}_{n-1} \in A_k \Leftrightarrow || \boldsymbol{x}_{n-1} - \boldsymbol{c}_k || < || \boldsymbol{x}_{n-1} - \boldsymbol{c}_m ||,$$

• This corresponds to segmenting the state space by hyperplanes. Of course the main question is how to define the centers $\{c_k\}$.

 $m \neq k$



- Three main avenues:
 - The centers are selected to represent the spatial distribution of the $\{x_{n-1}\}$, with an algorithm such as the Lloyd-Max one.
 - A sequential segmentation of the state space.
 - A heuristic selection (by a genetic algorithm for instance) of the centers among the $\{x_{n-1}\}$.



- Once the centers are selected, least squares estimation of the AR coefficients for each sub-model is simple.
- Here again, a modified MDL selection criterion can be employed. Quantization of the center coordinates (as for the RBF networks) must be incorporated to prevent selection of a limited number of centers only.



• Concerning stability, the sufficient condition of *drift back to the center* is satisfied if all AR sub-models used when $||\mathbf{x}_n||$ becomes large are stable. For instance:





- A very interesting alternative has been proposed in [2] under the name *competitive* local linear modeling.
- The opposite course of action is taken: first, the AR sub-models are estimated and second, the partition is performed.
- In this way, it may be that more complex partitions than center-based ones are obtained.





Sub-model estimation

- 1) First, a set of AR sub-models $\boldsymbol{a}_k = [a_{k0}, ..., a_{kp}]^{\mathsf{T}}$ is initialized randomly.
- 2) A vector $\mathbf{x}_{n-1} = [1, x_{n-1}, ..., x_{n-p}]^{\mathsf{T}}$ is chosen at random in the set of available data.
- 3) The AR model a_i giving minimum prediction error e_n with:

$$e_n = x_n - a_i^{\mathsf{T}} x_{n-1}$$

is found.



4) This model is updated using normalized LMS (NLMS):

$$a_i \leftarrow a_i + \beta e_n x_{n-1} / (x_{n-1}^T x_{n-1})$$

5) Back to 2) until the error variance stabilizes or becomes smaller than some threshold. Partitioning

Some classifier (MLP, k-nearest neighbors) is trained to establish the correspondence between each x_{n-1} and the appropriate model.



• The exponential AR (EXPAR) model has been introduced in [3]. It is described by:

$$x_n = \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^{p} [a_i + b_i \exp(-\alpha x_{n-1}^2)] x_{n-i} + \frac{1}{2} \sum_{i=1}^$$

• The motivation was to define a continuous NAR model able to produce limit cycles such as observed in the Lynx data.





- It is shown in [3] that sufficient conditions for the existence of a limit cycle (no excitation) are:
- All the poles of the AR model defined by the Ι. coefficients a_1, \ldots, a_p , i.e. for x_{n-1} very large, should be inside the unit circle.
- 2. Some of the poles of the AR model defined by the coefficients a_1+b_1, \ldots, a_p+b_p , i.e. for $x_{n-1}=0$, should be outside the unit circle.
- One of the two conditions below should hold: 3. $(1-\Sigma a_i)/\Sigma b_i > 1$ or $(1-\Sigma a_i)/\Sigma b_i < 0$



- Note that condition 1) is a stability condition of *drift* back to the center.
- Example: $x_n = [1.95 + 0.23 \exp(-x_{n-1}^2)]x_{n-1}$ $-[0.96 + 0.24 \exp(-x_{n-1}^2)]x_{n-2}$





- Least squares estimation of the coefficients $\{a_i, b_i\}$, once the scale parameter α is fixed, is quite easy.
- Concerning α , an optimization procedure can be used. Otherwise, an exhaustive grid search in a suitable range of values can be performed. Typically, $exp(-\alpha x_{n-1}^2)$ should be in the interval [0.25 0.75] for all values of x_{n-1} , which gives bounds on α .



• As a matter of fact the EXPAR model can be seen as a special (simple) case of the state dependent AR model introduced by Priestley [4]. This model is described by:

$$x_n = a_0(x_{n-1}) + a_1(x_{n-1})x_{n-1} + \dots + a_p(x_{n-1})x_{n-1} +$$

which expresses nonlinearity as a dependence of the AR coefficients on the state vector $\mathbf{x}_{n-1} = [x_{n-1}, \dots, x_{n-p}]^{\mathsf{T}}.$

 $X_{n-p} + \mathcal{E}_n$



- But one can observe that the state dependence in the EXPAR model can be seen as a very simple, mono-dimensional RBF network with only one center at 0.
- Also, if there is indeed a direct dependence of the AR coefficients on the state, then it could be well approximated by an RBF network, since it possesses the property of universal approximation.



• These remarks lead to the definition of the RBF-AR model, in which the state dependence of the AR coefficients is expressed by:

$$a_{i}(\boldsymbol{x}_{n-1}) = w_{i0} + \sum_{k=1}^{K} w_{ik} \exp\left(-\frac{\|\boldsymbol{x}_{n-1} - \boldsymbol{c}_{k}\|^{2}}{2\beta^{2}}\right)$$

• Note that, although it is not mandatory, one uses generally the same centers for all the coefficient expansions.



- The centers and width parameter β can be determined using heuristics as such as the ones presented for the RBF networks (it is also possible to have multiple width parameters).
- Once this is done, estimation of the coefficients $\{w_{ki}\}$ becomes a classical linear least squares problem.
- The MDL approach beased on center quantization can also be employed.



• But the terms in the RBF-AR model can also be regrouped differently:

$$x_{n} = w_{00} + \sum_{k=1}^{K} w_{0k} \phi_{k}(\mathbf{x}_{n-1}) + \left[w_{10} + \sum_{k=1}^{K} w_{1k} \phi_{k}(\mathbf{x}_{n-1}) + \left[w_{p0} + \sum_{k=1}^{K} w_{pk} \phi(\mathbf{x}_{n-1}) \right] \right] x_{n-p}$$

$$= w_{00} + \sum_{i=1}^{p} w_{i0} x_{n-i} + \sum_{k=1}^{K} \phi_{k}(\mathbf{x}_{n-1}) \left[w_{0k} + \sum_{i=1}^{K} w_{i0} x_{n-i} + \sum_{k=1}^{K} \phi_{k}(\mathbf{x}_{n-1}) \right] x_{n-p}$$

 \mathbf{x}_{n-1}) $|x_{n-1}|$





i.e., the RBF-AR model is the sum of a linear AR model plus a sum weighted by RBF of linear AR models.





- The RBF-AR model is thus a soft-transition version of the piecewise AR model. A particular sub-model will have more influence if is closer to the corresponding center.
- Due to the exponential decrease of the RBF with the norm of x_{n-1} , stability of the RBF-AR model depends only on the "background" sub-model (with weight 1).



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