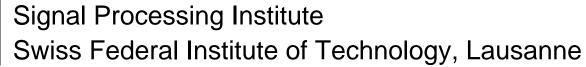
- ARMA models have enjoyed (and still enjoy) a wide popularity.
- Recent developments on ARMA models concern their extension to long-range dependence with fractionally integrated ARMA (FARIMA) models, and multivariate ARMA models.
- However, it was observed early on that some effects such as the *regime* one (different behaviors of residuals for different ranges of signal samples) cannot be dealt with using linear models.





- There are two main possibilities to model these types of effects:
  - Use non-Gaussian probability density functions (pdf)
  - Use non-linear models
- The first approach is usually intractable (it is usually hard to define the appropriate pdf), and most research has been centered on the second approach.







We will focus now on mimimum mean square error (MMSE) prediction. That is, if the sample x<sub>n+m</sub>, m ≥ 0, is to be predicted using the vector X<sub>n-1</sub> = [x<sub>n-1</sub>, x<sub>n-2</sub>,..., x<sub>1</sub>]<sup>T</sup> of past samples, one looks for a function f(X<sub>n-1</sub>) that minimizes:

$$\mathsf{E}[\{x_{n+m} - f(X_{n-1})\}^2]$$

• The MMSE predictor is the mean of  $x_{n+m}$ conditioned on the past, i.e.  $f(X_{n-1}) = \mathsf{E}[x_{n+m}|X_{n-1}]$ .



• Demonstration: for any predictor function g(.):

$$\mathsf{E}[\{x_{n+m} - g(X_{n-1})\}^2] = \mathsf{E}[\{x_{n+m} - \mathsf{E}[x_{n+m}|X_{n-1}]\}^2] + \mathsf{E}[\{\mathsf{E}[x_{n+m}|X_{n-1}] - g(X_{n-1})\}^2] + 2C$$

with

$$C = \mathsf{E}[\{x_{n+m} - \mathsf{E}[x_{n+m}|X_{n-1}]\}\{\mathsf{E}[x_{n+m}|X_{n-1}] - g(X_{n-1})\}]$$
$$= \mathsf{E}\big(\mathsf{E}[\{x_{n+m} - \mathsf{E}[x_{n+m}|X_{n-1}]\}\{\mathsf{E}[x_{n+m}|X_{n-1}] - g(X_{n-1})\}]|X_{n-1}\big)$$

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$$C = \mathsf{E}\big(\{\mathsf{E}[x_{n+m}|X_{n-1}] - g(X_{n-1})\}\mathsf{E}\{x_{n+m} - \mathsf{E}[x_{n+m}|X_{n-1}]|X_{n-1}\}\big)$$
$$= \mathsf{E}\big(\{\mathsf{E}[x_{n+m}|X_{n-1}] - g(X_{n-1})\}\{\mathsf{E}[x_{n+m}|X_{n-1}] - \mathsf{E}[x_{n+m}|X_{n-1}]\}\big)$$
$$= 0.$$

Thus 
$$\mathsf{E}[\{x_{n+m} - g(X_{n-1})\}^2] \ge \mathsf{E}[\{x_{n+m} - \mathsf{E}[x_{n+m}|X_{n-1}]\}^2]$$

• With  $\varepsilon_{n+m} = x_{n+m} - \mathsf{E}[x_{n+m}|X_{n-1}]$ , one can show that:  $\mathsf{E}[\varepsilon_{n+m}|X_{n-1}] = 0$   $\operatorname{Cov}(\varepsilon_s, \varepsilon_t) = 0$ 

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• In the mean square sense, a sample can be written as:

$$x_{n+m} = \mathsf{E}[x_{n+m}|X_{n-1}]\} + \mathcal{E}_{n+m}$$

that is, as the sum of the component predictible from the past, and the non-predictible part (innovation).

It can be proven [1] that, if the signal samples
 {x<sub>n</sub>} are Gaussian, then the MMSE predictor is linear.





• In the light of the discussion above, we will focus now on nonlinear predictors/models described by:

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

- with g(.) and  $\sigma(.)$  well behaved functions and  $\{\varepsilon_n\}$  an independent identically distributed sequence with unit variance.
- In almost all cases, σ(.) will be constant, except when we consider models with conditional heteroscedasticity.





- The models with  $\sigma(.)$  constant are called nonlinear autoregressive models (NAR).
- As for the linear AR models (which constitute a subset of NAR), one tries to have as small an order *p* as possible.
- Of course, what is desirable is that g(x<sub>n-1</sub>,..., x<sub>n-p</sub>) approximates E[x<sub>n</sub>|X<sub>n-1</sub>]} as well as possible. The choice of g(.) may be suggested by some a priori knowledge about the dynamics, or because it is in a set of functions with universal approximation capability.





- <u>Minimum variance criterion</u>
- Let us examine the case of a single model parameter  $\theta$  and an estimator  $\hat{\theta}$ . A natural optimality criterion for the estimator is the mean square error (MSE):

mse
$$(\hat{\theta}) = \mathsf{E}[(\hat{\theta} - \theta)^2] = \mathsf{E}\{[(\hat{\theta} - \mathsf{E}(\hat{\theta})) + (\mathsf{E}(\hat{\theta}) - \theta)]^2\}$$
  
= var $(\hat{\theta}) + [\mathsf{E}(\hat{\theta}) - \theta]^2 = var(\hat{\theta}) + b^2(\hat{\theta})$   
with  $b(\hat{\theta}) = \mathsf{E}(\hat{\theta}) - \theta$  the estimator bias

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- Unfortunately, the minimum MSE estimator cannot be obtained in most cases. A feasible approach consists in constraining the bias to be zero and find the estimator with the minimum variance. This estimator is the minimum variance unbiased (MVU) estimator.
- Note that the variance of the MVU estimator should be the smallest for *all possible values* of *θ*.





- There is no general procedure to find the MVU estimator. One possible way to find it is establish the Cramer-Rao lower bound (CRLB).
- The CRLB is the lower bound on the variance of *any* unbiased estimator. If indeed an estimator variance reaches this bound, then it is the MVU one.
- It may happen that no estimator reaches this bound, but that the MVU still exists.





- CRLB for a parameter vector  $\boldsymbol{\theta}$
- The covariance matrix of any unbiased estimator satisfies:  $C_{\hat{\theta}} - I^{-1}(\theta) \ge 0$ Meaning that the matrix is positive semidefinite. The Fisher information matrix  $I(\theta)$  is given by:  $\begin{bmatrix} \partial^2 \ln p(\mathbf{r}; \theta) \end{bmatrix}$

$$\mathbf{I}(\boldsymbol{\theta})_{ij} = -\mathsf{E}\left[\frac{\partial^2 \ln p(\boldsymbol{x};\boldsymbol{\theta})}{\partial \theta_i \partial \theta_j}\right]$$

With  $p(x; \theta)$  the probability density function of the data x parameterized by  $\theta$ .



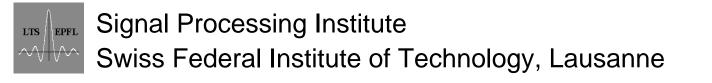
• An unbiased estimator that reaches the CRLB, that is:  $C_{\hat{\theta}} = I^{-1}(\theta)$ 

can be found if and only if

$$\frac{\partial \ln p(\boldsymbol{x};\boldsymbol{\theta})}{\partial \boldsymbol{\theta}} = \mathbf{I}(\boldsymbol{\theta})(h(\boldsymbol{x}) - \boldsymbol{\theta})$$

for some multidimensional function *h*. In that case the MVU estimator is:  $\hat{\theta} = h(x)$ 

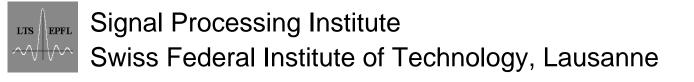
Unfortunately, this approach is easy to apply only for linear models and data with Gaussian statistics.





- <u>Maximum likelihood (ML) estimation</u>
- The ML estimator is the value of  $\theta$  maximizing the likelihood  $p(x;\theta)$ , where x is now the vector of observed data samples.
- It is not optimal in general, but asymptotically is the MVU estimator.
- Under some conditions, asymptotically:

 $\hat{\theta} \approx N(\boldsymbol{\theta}, \mathbf{I}^{-1}(\boldsymbol{\theta}))$ 





 Least squares (LS) estimation
 For NAR models, and N samples available, the LS estimator is the value of minimizing:

lse = 
$$\sum_{n=1}^{N} [x_n - g(x_{n-1}, \dots, x_{n-p}; \theta)]^2$$

• The estimation performance depends on the distribution of the modeling errors. If this distribution is Gaussian and the errors are uncorrelated, then LS is equvalent to ML.





- Conditions have been derived in the literature
   [2] on the ergodicity of (vector) Markov
   chains {Y<sub>n</sub>}.
- A sufficient condition for ergodicity is the existence of positive constants  $\alpha$ ,  $\beta$ , and  $\gamma$  such that:

$$E(||\boldsymbol{Y}_{n}|| - ||\boldsymbol{Y}_{n-1}|| \mid \boldsymbol{Y}_{n-1} = \boldsymbol{y}) \leq -\beta, ||\boldsymbol{y}|| > \alpha \quad (1)$$
$$E(||\boldsymbol{Y}_{n}|| - ||\boldsymbol{Y}_{n-1}|| \mid \boldsymbol{Y}_{n-1} = \boldsymbol{y}) \leq \gamma, ||\boldsymbol{y}|| \leq \alpha \quad (2)$$

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- Condition (2) is not very stringent. It just imposes that the increase in norm inside a ball of radius *α* is bounded.
- Condition (1) expresses that there must be some mechanism for *drift back to the center*, i.e. that if the norm of the instance of the Markov chain becomes large, then the norms of successive instances must decrease. In this way, ergodicity (existence of an equilibrium pdf) is guaranteed.





• A stronger condition for geometrical ergodicity (exponentially fast convergence to the equilibrium pdf starting from arbitrary initial conditions) is obtained by replacing condition (1) with:

 $E(r||Y_n|| - ||Y_{n-1}|| | Y_{n-1} = y) \le -\beta, ||y|| > \alpha$  (1b) with *r* a constant > 1.





 Now {x<sub>n</sub>} in the NAR model is not Markovian. A Markov chain is obtained by using a state space representation:

$$\boldsymbol{X}_n = \boldsymbol{G}(\boldsymbol{X}_{n-1}) + \boldsymbol{E}_n$$

with

$$X_{n-1} = [x_{n-1}, x_{n-2}, \dots, x_{n-p}]^{\mathsf{T}}$$

$$G(X_{n-1}) = [g(x_{n-1}, \dots, x_{n-p}), x_{n-1}, \dots, x_{n-p+1}]^{\mathsf{T}}$$

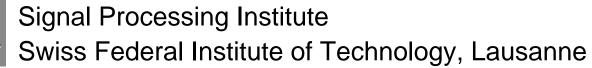
$$E_n = [\sigma(x_{n-1}, \dots, x_{n-p}) \mathcal{E}_n, 0, \dots, 0]^{\mathsf{T}}$$

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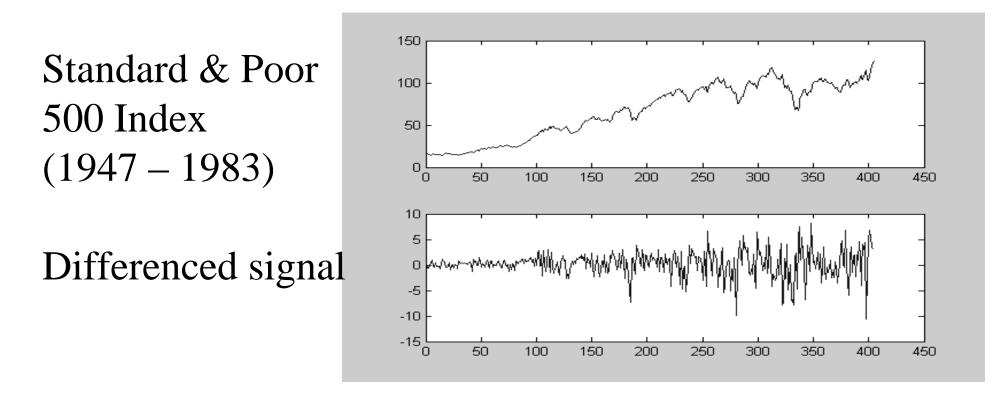
- If the mean does not carry information by itself, it is often judicious to remove it from the data. If it is not removed, one should include a constant term in the NAR model to take it into account.
- Also, deterministic trends (ramp) and oscillations (for instance seasonal effects) should be removed.
- Stochastic trends (unit root) should also be dealt with. Statistical tests (such as the Augmented Dickey-Fuller one) test for the presence of such a trend.







By all means, the presence of a unit root is usually visible in the signal, and differencing  $y_n = x_n - x_{n-1}$ , usually solves this problem.





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- Obviously, the first test is to check whether the signal samples are Gaussian or not, using the Kolmogorov-Smirnov or the Chi-square test for instance.
- Likelihood-ratio (LR) tests have been developed to test the significance of a nonlinear model with respect to a linear one. Unfortunately, LR tests have to be tailored to the nonlinear models used, and the test statistics may be hard to evaluate [3].



- A test for nonlinearity based on higher-order statistics has also been proposed in [4]. It uses 3rd and 4th order cumulants and test statistics are derived.
- A test based on time irreversibility (linear signals are time reversible) has also been proposed. The measure of time irreversibility is given by:

$$\phi_{rev}(\tau) = \frac{1}{N-\tau} \sum_{n=\tau+1}^{N} (x_n - x_{n-\tau})^3$$

## For a suitable lag $\tau$ .

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It is also possible to select the best polynomial model on the signal with respect to a model selection criterion such as MDL.
The linear AR models constitute a subset of the set of polynomial models. If only linear terms ar retained, then there is non nonlinearity in the signal.





- In order to assess the significance of a test, a powerful approach, *surrogate analysis*, has recently been introduced [5].
- The idea is the following: suppose one has measured some feature *m* on the signal at hand. One generates synthetic (surrogate) signals sharing some properties of the original signal (sample pdf and 2nd-order statistics), *but not the hypothetized nonlinear relationship between samples*.





Then one can compute a significance S:

$$S = \frac{|m - \langle m \rangle_{surr}|}{\sigma_{surr}}$$

Where  $\langle m \rangle_{surr}$  is the mean of the distribution of the feature for the surrogates and  $\sigma_{surr}$  its standard deviation.

Assuming *m* is Gaussian a value S = 2.6corresponds to a significance level of 0.01 for the value of *m* obtained on the original signal.





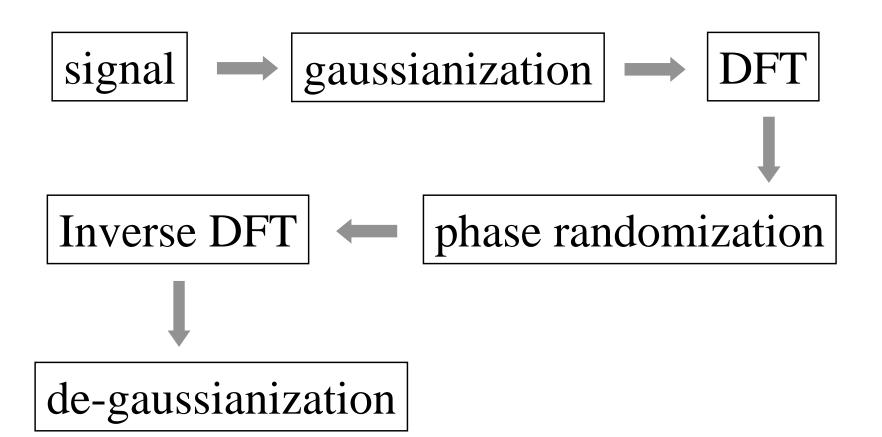
Since the assumption that *m* is Gaussian may be bold, one can also use a rank-order test [5].

• To build these surrogates, one uses the fact that linear relationships between samples imply only 2nd-order statistics, i.e. the autocorrelation function, which is even and doest not carry any phase information.





• Principle of surrogate generation:





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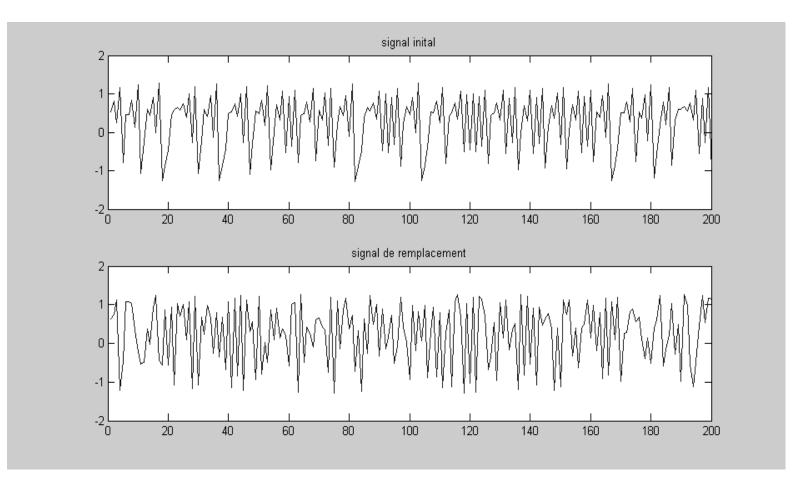
- To "Gaussianize" the samples, one feeds them through an instantaneous nonlinearity which is the distribution of the samples.
- Phase randomization on the discrete Fourier transform (phases uniformly drawn between 0 and 2π), destroys any potential nonlinear structure.
- De-Gaussianization consist in applying the inverse of the instantaneous linear transform.

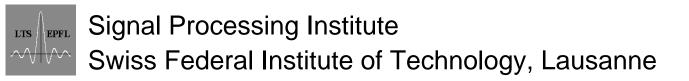






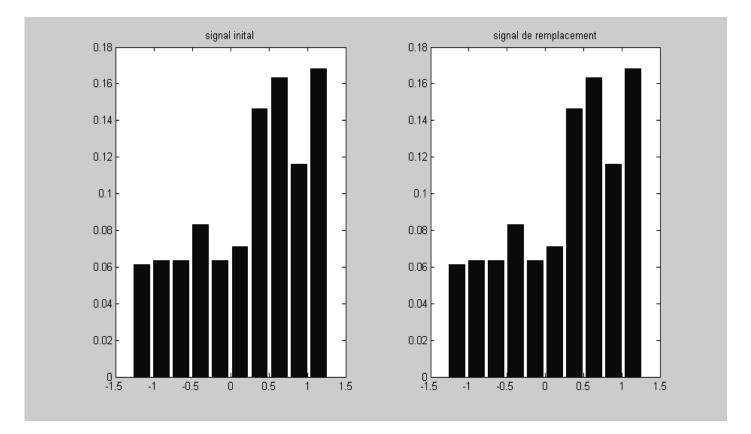
## • Example: surrogate for a chaotic signal







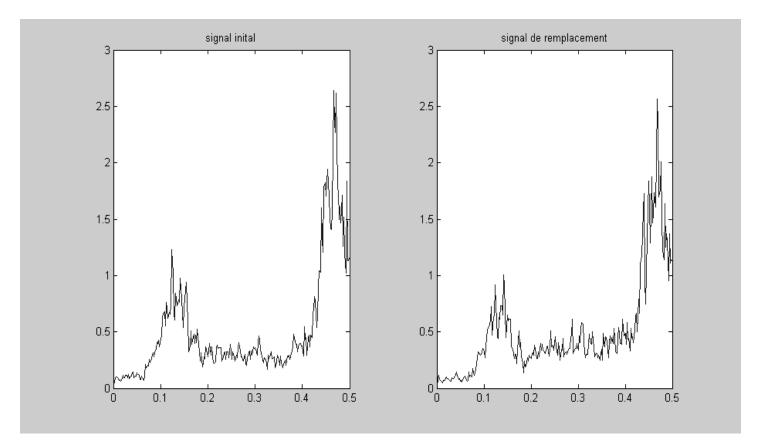
## Estimated probability density functions:

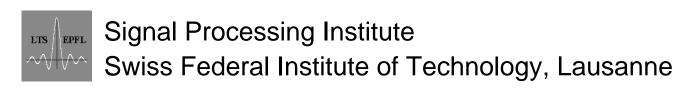






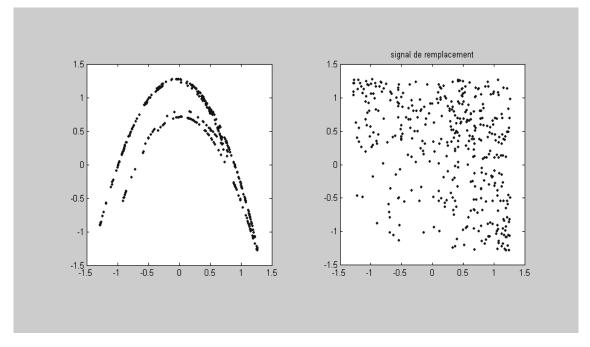
#### Estimated power spectra







#### But in the state space...



# The structure present in the initial signal has been destroyed.

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