

Average, synchronize and analyze recurrent Bioelectrical Waves

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Goal : Use formal approaches for very practical problems

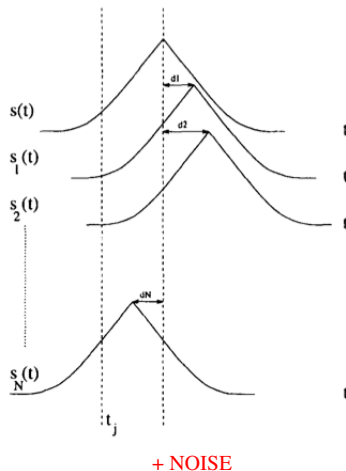
Bioelectrical signals are :

- ✓ ● repetitive (ECG, EMG, ...)
- ✓ ● spontaneously or evoked (ECG, EP, ...)
- with small variations and noisy

From the Signal Processing/Biological knowledge :

- How to get rid of the noise presence ($s(t)$?)
- How to measure variability?
- How to interpret results?
- How to model the observations?

⇒ improve the Bio/DSP skills



Goal : characterize Waves or variabilities (simple model)

From the data set :

- delays ?
- amplitudes (small, large) ?

⇒ averaging and synchronization

- estimation theory
- probabilistic approach

Averaging

Let's assume the observation model (normal law $\mathcal{N}(0, \sigma)$ for the white noise) with unknown signal s

$$x_i(n) = s(n) + w_i(n) \quad \text{digital signal} \quad (1)$$

define the vector $\mathbf{x}_i = [x_i(0) \ x_i(1) \ \cdots \ x_i(N-1)]^T$, then the PDF of the observation is :

$$p(x_i(n); s(n)) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2\sigma^2} (x_i(n) - s(n))^2} \Rightarrow \quad (2)$$

$$p(\mathbf{x}_i; \mathbf{s}) = \prod_{n=0}^{N-1} p(x_i(n); s(n)) = \frac{1}{(2\pi\sigma^2)^{N/2}} e^{-\frac{1}{2\sigma^2} \sum_{n=0}^{N-1} (x_i(n) - s(n))^2} \quad (3)$$

$$p(\mathbf{x}_i; \mathbf{s}) = \frac{1}{(2\pi\sigma^2)^{N/2}} e^{-\frac{1}{2\sigma^2} (\mathbf{x}_i - \mathbf{s})^T (\mathbf{x}_i - \mathbf{s})} \quad (4)$$

If the observations are independent the global PDF is :

$$p(\mathbf{x}_1; \mathbf{x}_2; \cdots; \mathbf{x}_I; \mathbf{s}) = \prod_{i=1}^I p(\mathbf{x}_i; \mathbf{s}) = M e^{-\frac{1}{2\sigma^2} \sum_{i=1}^I (\mathbf{x}_i - \mathbf{s})^T (\mathbf{x}_i - \mathbf{s})} \quad (5)$$

This PDF can be viewed as a Likelihood function \Rightarrow MLE

Averaging

An estimation $\hat{\mathbf{s}}$ of \mathbf{s} can be computed from the minimization (see MLE Theory)

$$\hat{\mathbf{s}} = \arg \min_{\mathbf{s}} \sum_{i=1}^I (\mathbf{x}_i - \mathbf{s})^T (\mathbf{x}_i - \mathbf{s}) = \arg \min_{\mathbf{s}} \sum_{i=1}^I (\mathbf{x}_i^T \mathbf{x}_i - 2\mathbf{x}_i^T \mathbf{s} + \mathbf{s}^T \mathbf{s}) = \arg \min_{\mathbf{s}} J \quad (6)$$

Equivalent to :

$$\frac{\partial J}{\partial \mathbf{s}} = 0 \Leftrightarrow -2 \sum_{i=1}^I \mathbf{x}_i + 2 \sum_{i=1}^I \mathbf{s} = 0 \Rightarrow \hat{\mathbf{s}} = \frac{1}{I} \sum_{i=1}^I \mathbf{x}_i \quad (7)$$

That is simply the sample mean (approximated mathematical expectation) !!!

What happens with this estimation if a **magnitude factor** is present (physiological variability)?

Weighted Averaging

Let's assume the observation model (normal law $\mathcal{N}(0, \sigma)$ for the white noise) with unknown signal s and a_i 's :

$$x_i(n) = a_i s(n) + w_i(n) \quad (8)$$

define the vector $\mathbf{x}_i = [x_i(0) \ x_i(1) \ \cdots \ x_i(N-1)]^T$ and \mathbf{w}_i , the observation is replaced by :

$$\mathbf{x}_i = a_i \mathbf{s} + \mathbf{w}_i \quad (9)$$

Then the criteria J to be minimized is :

$$\hat{\mathbf{s}}, \hat{\mathbf{a}} = \arg \min_{\mathbf{s}, \mathbf{a}} \sum_{i=1}^I (\mathbf{x}_i - a_i \mathbf{s})^T (\mathbf{x}_i - a_i \mathbf{s}) = \arg \min_{\mathbf{s}, \mathbf{a}} \sum_{i=1}^I \|\mathbf{x}_i - a_i \mathbf{s}\|_2^2 \quad (10)$$

$$\hat{\mathbf{s}}, \hat{\mathbf{a}} = \arg \min_{\mathbf{s}, \mathbf{a}} \|\mathbf{X} - \mathbf{s} \mathbf{a}^T\|_F^2 \quad (11)$$

with \mathbf{X} the matrix formed by the \mathbf{x}_i 's, and $\|\cdot\|_F^2$ the squared Frobenius norm.

The solution of this minimization is obtained by using two partial derivatives $\frac{\partial J}{\partial \mathbf{s}}$ and $\frac{\partial J}{\partial \mathbf{a}}$.

$$\frac{\partial J}{\partial \mathbf{s}} = \frac{\partial}{\partial \mathbf{s}} \frac{1}{I} \sum_{i=1}^I (\mathbf{x}_i^T \mathbf{x}_i - 2a_i \mathbf{x}_i^T \mathbf{s} + a_i^2 \mathbf{s}^T \mathbf{s}) = \sum_{i=1}^I (-2a_i \mathbf{x}_i + 2a_i^2 \mathbf{s}) \quad (12)$$

$$\frac{\partial J}{\partial \mathbf{s}} = 0 \Rightarrow \hat{\mathbf{s}} = \frac{1}{\sum_i a_i^2} \sum_{i=1}^I a_i \mathbf{x}_i = \text{weighted averaging} \quad \checkmark \quad (13)$$

but $\mathbf{x}_i = a_i \mathbf{s} = \frac{a_i}{\alpha} \mathbf{s} \alpha = \tilde{a}_i \tilde{\mathbf{s}}$, not unique solution ! \Rightarrow impose $\sum a_i^2 = \mathbf{a}^T \mathbf{a} = 1$ (good for Lagrange multiplier)

Weighted Averaging

But a_i 's are unknown ...

Let's compute the second partial derivative (turns out to be several derivatives regards a_i 's) :

$$\frac{\partial J}{\partial a_i} = \frac{\partial}{\partial a_i} \frac{1}{I} \sum_{i=1}^I (\mathbf{x}_i^T \mathbf{x}_i - 2a_i \mathbf{x}_i^T \mathbf{s} + a_i^2 \mathbf{s}^T \mathbf{s}) = \sum_{i=1}^I (-2\mathbf{x}_i^T \mathbf{s} + 2a_i \mathbf{s}^T \mathbf{s}) \quad (14)$$

$$\frac{\partial J}{\partial a_i} = 0 \Rightarrow \hat{a}_i = \frac{\mathbf{x}_i^T \mathbf{s}}{\mathbf{s}^T \mathbf{s}} \quad (15)$$

In fact, since \mathbf{s} and \mathbf{a} are unknown results (13) and (15) cannot be computed. To perform the global minimization result (13) should be replaced in (9) and the partial derivatives with respect to the a_i 's computed to get the estimations of the magnitude factors \Rightarrow highly non linear ... instead use an Alternated Least Square algo :

INIT

$$(1) \mathbf{s} = \frac{1}{I} \sum_{i=1}^I \mathbf{x}_i ; (2) \mathbf{a}_i = \frac{\mathbf{x}_i^T \mathbf{s}}{\mathbf{s}^T \mathbf{s}} \text{ for } i = 1 \dots I \quad (16)$$

$$\text{normalize } \mathbf{a}'_i \mathbf{s} \text{ by } \sqrt{\sum \mathbf{a}_i^2} \quad (17)$$

DO

$$(1) \mathbf{s} = \frac{1}{\sum_i \mathbf{a}_i^2} \sum_{i=1}^I \mathbf{a}_i \mathbf{x}_i ; (2) \mathbf{a}_i = \frac{\mathbf{x}_i^T \mathbf{s}}{\mathbf{s}^T \mathbf{s}} \text{ for } i = 1 \dots I \quad (18)$$

$$\text{normalize } \mathbf{a}'_i \mathbf{s} \text{ by } \sqrt{\sum \mathbf{a}_i^2} \quad (19)$$

WHILE CONVERGENCE (e.g. Frobenius norm)

Weighted Averaging-PCA

If we impose the solution $\mathbf{s} = \mathbf{X}\mathbf{m}$ (weighted averaging of the observations or linear combination) and maximize its energy $C = \mathbf{s}^T \mathbf{s} (\approx E[s^2])$ subject to the constraint $\mathbf{m}^T \mathbf{m} = 1$. Then,

$$C = \mathbf{m}^T \mathbf{X}^T \mathbf{X} \mathbf{m} = \mathbf{m}^T \mathbf{R}_x \mathbf{m} \quad (20)$$

Where \mathbf{R}_x is the approximated Covariance matrix of the observations ($\mathbf{R}_x = E[\mathbf{x}\mathbf{x}^T]$)

Maximization of C can be accomplished by means of the Lagrange's multiplier technique by defining the new criterion :

$$J = J(\mathbf{m}) = \mathbf{m}^T \mathbf{R}_x \mathbf{m} - \lambda (\mathbf{m}^T \mathbf{m} - 1) \quad (21)$$

Then the derivation of J with respect to \mathbf{m} is :

$$\frac{\partial J}{\partial \mathbf{m}} = 2\mathbf{R}_x \mathbf{m} - 2\lambda \mathbf{m} = \mathbf{0} \Rightarrow \mathbf{R}_x \mathbf{m} = \lambda \mathbf{m} \quad (22)$$

This corresponds to a eigenvalues/eigenvectors decomposition of the matrix \mathbf{R}_x , where \mathbf{m} is an eigenvector and λ the respective eigenvalue. How to select the correct eigenvalue/eigenvector ?

Replace \mathbf{m} by an eigenvector \mathbf{v} of \mathbf{R}_x in J , it gives $J = \lambda$. Then J is max when choosing $\mathbf{m} = \mathbf{v}$ the eigenvector corresponding to the largest λ .

Weighted Averaging-Generalization

How to relate this estimation to the solution of the eigenvalue/eigenvector decomposition ?

The criteria to be minimized is in fact :

$$J = \|\mathbf{X} - \mathbf{s}\mathbf{a}^T\|_F^2 = tr(\mathbf{X} - \mathbf{s}\mathbf{a}^T)^T(\mathbf{X} - \mathbf{s}\mathbf{a}^T) = tr(\mathbf{X}^T\mathbf{X} - \mathbf{X}^T\mathbf{s}\mathbf{a}^T - \mathbf{a}\mathbf{s}^T\mathbf{X} + \mathbf{a}\mathbf{s}^T\mathbf{s}\mathbf{a}^T) \quad (23)$$

$$J = tr(\mathbf{R}) - tr(\mathbf{X}^T\mathbf{s}\mathbf{a}^T) - tr(\mathbf{a}\mathbf{s}^T\mathbf{X}) + tr(\mathbf{a}\mathbf{s}^T\mathbf{s}\mathbf{a}^T) = tr(\mathbf{R}) - 2tr(\mathbf{X}^T\mathbf{s}\mathbf{a}^T) + tr(\mathbf{a}\mathbf{s}^T\mathbf{s}\mathbf{a}^T) \geq 0 \quad (24)$$

Where \mathbf{R}_x is the approximated Covariance matrix of the observations ($\mathbf{R}_x = E[\mathbf{x}\mathbf{x}^T]$).

We aim to minimize J (≥ 0), since $tr(\mathbf{R}) \geq 0$ then this minimization is equivalent to :

$$\max_{\mathbf{s}, \mathbf{a}} \{2tr(\mathbf{X}^T\mathbf{s}\mathbf{a}^T) - tr(\mathbf{a}\mathbf{s}^T\mathbf{s}\mathbf{a}^T)\} = \max_{\mathbf{s}, \mathbf{a}} J' \quad (25)$$

Imposing the solution $\mathbf{s} = \mathbf{X}\mathbf{m}$ (\mathbf{s} is a linear combination of the observations, see before), then we get :

$$\max_{\mathbf{s}, \mathbf{a}} \{2tr(\mathbf{X}^T\mathbf{X}\mathbf{m}\mathbf{a}^T) - tr(\mathbf{a}\mathbf{m}^T\mathbf{X}^T\mathbf{X}\mathbf{m}\mathbf{a}^T)\} = \max_{\mathbf{s}, \mathbf{a}} \{2tr(\mathbf{R}\mathbf{m}\mathbf{a}^T) - tr(\mathbf{a}\mathbf{m}^T\mathbf{R}\mathbf{m}\mathbf{a}^T)\} \quad (26)$$

Weighted Averaging-Generalization

The derivation with respect to \mathbf{a} gives :

$$\frac{\partial J'}{\partial \mathbf{a}} = 2\mathbf{R}\mathbf{m} - 2\mathbf{m}^T \mathbf{R}\mathbf{m}\mathbf{a} = \mathbf{0} \Rightarrow \mathbf{R}\mathbf{m} = \mathbf{m}^T \mathbf{R}\mathbf{m}\mathbf{a} \quad (27)$$

If \mathbf{m} is selected as an eigenvector \mathbf{v} of \mathbf{R} (constant with respect to the maximization), then (27) turns out to be $\lambda \mathbf{v} = \mathbf{v}^T \lambda \mathbf{v}\mathbf{a} = \lambda \mathbf{a}$, then $\mathbf{a} = \mathbf{v}$. This solution is valid because it has been seen that the optimal \mathbf{s} is $\mathbf{s} = \frac{\mathbf{X}\mathbf{a}}{\mathbf{a}^T \mathbf{a}}$, corresponding to the form $\mathbf{s} = \mathbf{X}\mathbf{m}$.

Which eigenvector should be chosen (J' has to be maximized) ?

Replace these solutions in J' we get :

$$J' = 2tr(\mathbf{R}\mathbf{v}\mathbf{v}^T) - tr(\mathbf{v}\mathbf{v}^T \mathbf{R}\mathbf{v}\mathbf{v}^T) = 2tr(\lambda \mathbf{v}\mathbf{v}^T) - tr(\lambda \mathbf{v}\mathbf{v}^T \mathbf{v}\mathbf{v}^T) = \lambda tr(\mathbf{v}\mathbf{v}^T) = \lambda \mathbf{v}^T \mathbf{v} = \lambda \quad (28)$$

Then it is maximized when choosing the eigenvector \mathbf{v}_{max} corresponding to the largest eigenvalue λ_{max} !

The optimal solution consists in selecting $\mathbf{s} = \mathbf{X}\mathbf{v}_{max}$.

With respect to eigenvalue decomposition, it can be interpreted as the maximized weighted averaging

$\mathbf{s} = \mathbf{X}\mathbf{m}$ subject to $\mathbf{m}^T \mathbf{m} = 1$.

Averaging

The model can be extended to :

$$x_i(n) = s(n - d_i) + w_i(n) \quad (29)$$

The ensemble mean is expressed as :

$$\hat{s}(n) = \frac{1}{I} \sum_{i=1}^I x_i(n) \quad (30)$$

when the number of observed signals is large enough and that the variable d is random, this relation can be expressed as :

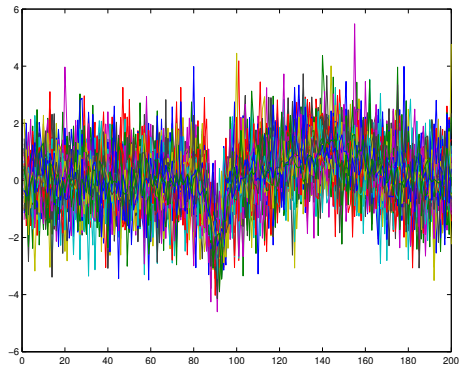
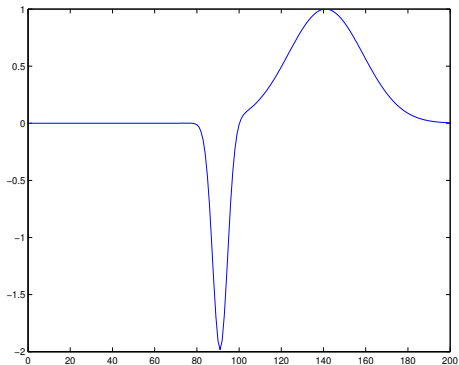
$$\hat{s}(n) = \int s(t - a) p_D(a) da \quad (31)$$

with $p_D(a)$ the probability densities of the delays. Then, the mean is the convolution of the pdf and the signal s

- If the pdf is a gaussian, s is low pass filtered
- What happens if the noise is not gaussian (laplacian ?) \Rightarrow the median replace the mean
- However, this quantity may be use to address the quality of a synchronization procedure.

Simulations

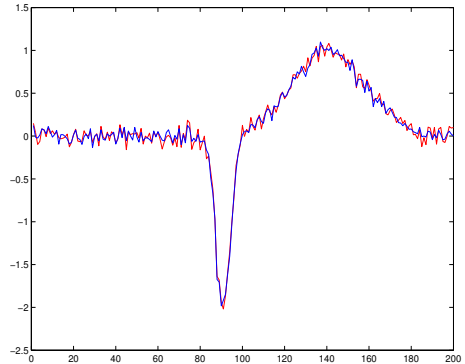
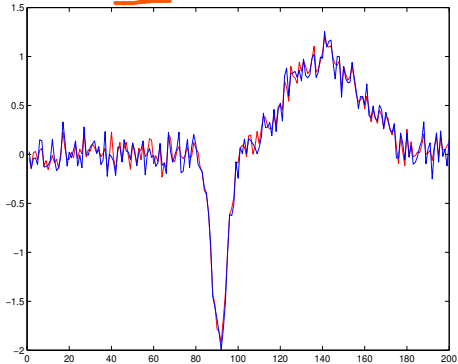
One pure observation, 100 noisy sweeps



Simulations

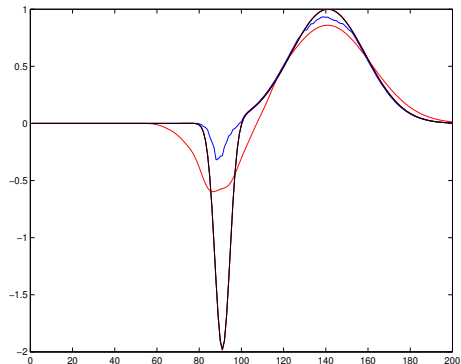
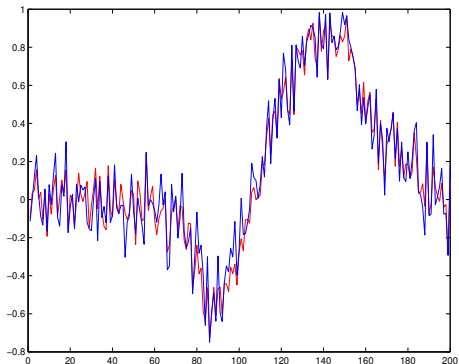
Gaussian noise (Left) and Laplacian noise (Right)

Mean and Median



Simulations

Delayed signal ($\sigma_d = 10$) with and without noise
 Mean, Median and Original



Alignment quality

Assuming that delayed signals are observed as following :

$$x_i(t) = w_i s(t - d_i) + n_i(t) \quad (32)$$

The ensemble mean is expressed as :

$$m_1(t) = \frac{1}{K} \sum_{i=1}^K x_i(t) \quad (33)$$

when the number of observed signal is large enough and that the random variables w and d are independant, this relation can be expressed as :

$$m_1(t) = \int \int w s(t - a) p_D(a) p_W(w) da dw \quad (34)$$

with $p_D(a)$ and $p_W(w)$ the probability densities of the delays and the weighting factor. Then, the mean is :

$$m_1(t) = \bar{w} \int s(t - a) p_D(a) da \quad (35)$$

Alignment quality

When calculating the energy of the mean signal V as :

$$V = \int m_1^2(t) dt \quad (36)$$

and assuming that in each observation x_i the signal $s(t)$ is completely observed, the integral is :

$$V = \bar{w}^2 \iint p_D(a) p_D(b) R_{ss}(a-b) da db \quad (37)$$

where $R_{ss}(\tau)$ is the temporal cross-correlation function defined as :

$$R_{ss}(\tau) = \int s(t) s(t+\tau) dt \quad (38)$$

Using the definition of the characteristic function $\phi_D(u)$ such as :

$$p_D(a) = \int e^{-jua} \phi_D(u) du \quad (39)$$

and $\hat{R}_{ss}(u)$ the Fourier transform of $R_{ss}(t)$, it can be shown that (37) is :

$$\bar{w}^2 \iint p_D(b) \phi_D(v) \hat{R}_{ss}(v) e^{-jvb} dv db \quad (40)$$

which can be reduced to :

$$\bar{w}^2 \int \phi_D(v) \phi_D(-v) \hat{R}_{ss}(v) dv \quad (41)$$

Alignment quality

The probability density being real, the property $\phi_D^*(u) = \phi_D(-u)$ is verified, giving :

$$V = \bar{w}^2 \int |\phi_D(u)|^2 \hat{R}_{ss}(v) dv \quad (42)$$

The cross-correlation being semi-definite positive, its Fourier transform $\hat{R}_{ss}(u)$ is then non negative and even. This implies that :

$$V = 2\bar{w}^2 \int_0^\infty |\phi_D(v)|^2 \hat{R}_{ss}(v) dv \geq 0 \quad (43)$$

Assuming the normal law $\mathcal{N}(m, \sigma)$ for $p_D(a)$, its characteristic function is :

$$\phi_D(v) = e^{jmv - (1/2)\sigma^2 v^2} \quad (44)$$

which implies for V :

$$V = 2\bar{w}^2 \int_0^\infty e^{-\sigma^2 v^2} \hat{R}_{ss}(v) dv \quad (45)$$

Alignment quality

The derivative of V regards to σ gives :

$$\frac{d}{d\sigma} V = -4\bar{w}^2 \sigma \int_0^\infty v^2 e^{-\sigma^2 v^2} \hat{R}_{ss}(v) dv \leq 0 \quad (46)$$

Then if the jitter variance decreases the criteria V increases. Considering that after the alignment process the probability density of the residual delays could change its shape, if the function $|\phi_D(v)|^2$ can be approximated by a gaussian law such that :

$$|\phi_D(v)|^2 \approx e^{-\sigma^2 v^2} \text{ pour } v \in \Omega \quad (47)$$

with Ω the support of $\hat{R}_{ss}(v)$, then **when V increases it means that the variance, i.e. the alignment error, has been reduced.**

Goal : Use formal approaches for very practical problems

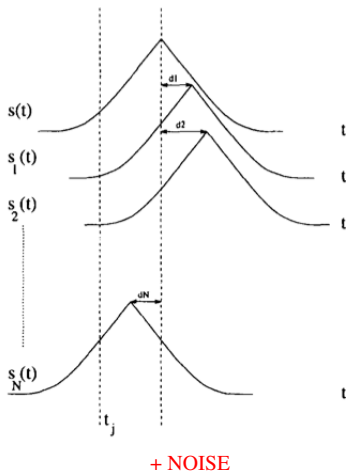
Bioelectrical signals are :

- repetitive (ECG, EMG, ...)
- spontaneously or evoked (ECG, EP, ...)
- with small variations and noisy

From the Signal Processing/Biological knowledge :

- How to get rid of the noise presence : average
- **How to measure variability (d_i 's)?**
- How to interpret results ?
- How to model the observations ?

⇒



Mean -> Variance

- the delays are artifacts and need to be canceled ?
- the delays convey physiological informations ?

⇒ Each individual delay is estimated (TDE) or its statistics (Variance) ?

Define the simple model :

$$x_i(t) = s(t - d_i) + n_i(t) \quad (48)$$

The variance is defined by (without noise) :

$$V_1(t) = \int s^2(t - a) p_D(a) da - m^2(t) \quad (49)$$

Assuming that delays are small (versus the derivatives of s), we have the Taylor expansion :

$$s(t - a) \approx s(t) - as'(t) \quad (50)$$

Then,

$$V_1(t) = \int (s^2(t) + a^2 s'^2(t) - 2s(t)s'(t)) p_D(a) da - m^2(t) \quad (51)$$

$$= s^2(t) + s'^2(t) \int a^2 p_D(a) - 2s(t)s'(t) \int a p_D(a) - m^2(t) \quad (52)$$

Mean -> Variance

Assuming that $\bar{a} = 0$, $m(t) = \int s(t-a)p_D(a)da = \int (s(t) - as'(t))p_D(a)da = s(t) \int p_D(a)da - s'(t).0 = s(t)$
 Then,

$$V_1(t) = \sigma_d^2 s'^2(t) \quad (53)$$

Introducing the noise and a random magnitude w with a unit mean, corresponding to the model :

$$x_i(t) = w_i s(t - d_i) + n_i(t) \quad (54)$$

It can be shown that similarly we finally get :

$$V_1(t) = s^2(t) \sigma_w^2 + s'^2(t) \sigma_d^2 (1 + \sigma_w^2) + \sigma_n^2(t) \quad (55)$$

Then, the variance of the observations is function of the second order statistics of the magnitude, the delay and the noise.

If $s(t)$ is known (or approximated), the quantities can be estimated (Least squares)

A real example : EP

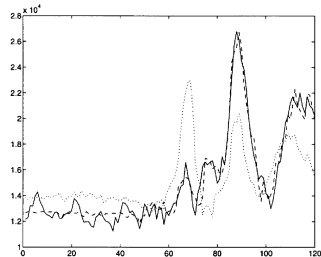
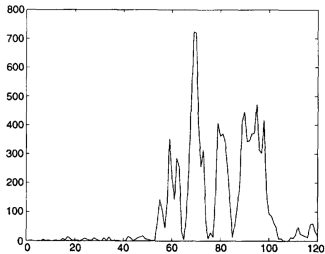
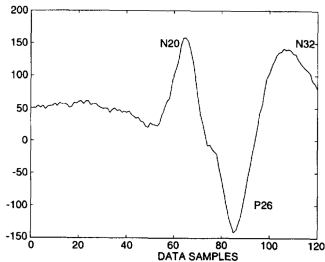


Table 1
Characteristics of the three waves

Wave	N20	P26	N32
$\hat{\sigma}_d$ (ms)	0.17	2.06	2.16
$\hat{\sigma}_p$	0.17	0.49	0.00

Time Delay Estimation

Using the simple model :

$$x_i(n) = s(n - d_i) + w_i(n) \quad (56)$$

and using the MLE (see previous development), the delays \mathbf{d} are estimated by maximizing the criteria :

$$\hat{\mathbf{d}} = \arg \max_{\mathbf{d}} \sum_n \sum_i \sum_{k>i} x_k(n + d_k) x_i(n + d_i) \quad (57)$$

Not proven to be the best estimator. It is if it attains the Cramer-Rao Lower Bound (CRLB).

Assuming this model and gaussian noise, the CRLB is :

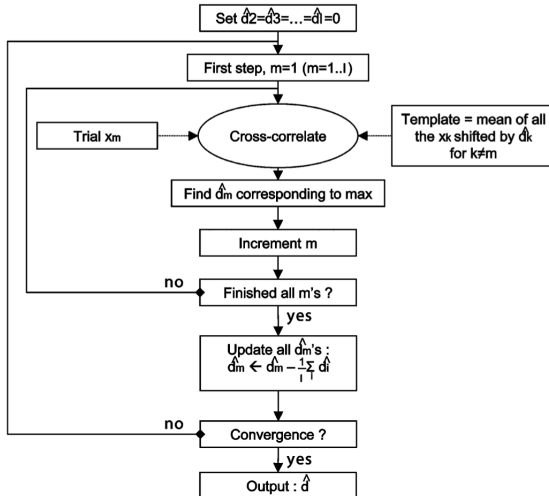
$$\text{var}(\hat{d}_i) \geq \frac{2\sigma_w^2}{\mathbf{s}'^T \mathbf{s}'} \quad (58)$$

Rem : if \mathbf{s} is known $\text{var}(\hat{d}_i) \geq \frac{\sigma_w^2}{\mathbf{s}'^T \mathbf{s}'}$

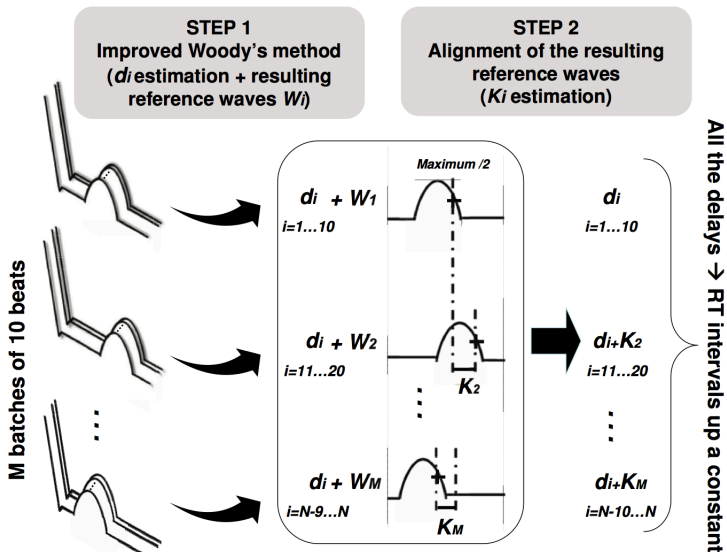
Use an iterative (delays estimated iteratively) scheme instead of brute force optimization

Application to ECG intervals modeling (QT/RR)

Flowchart



Batch TDE



Goal : characterize Waves variations (complex model)

From the data set :

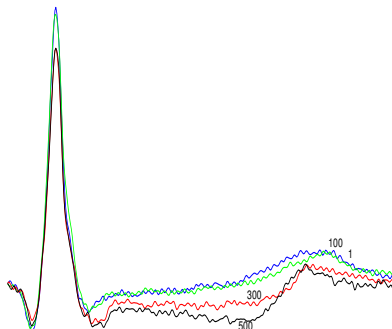
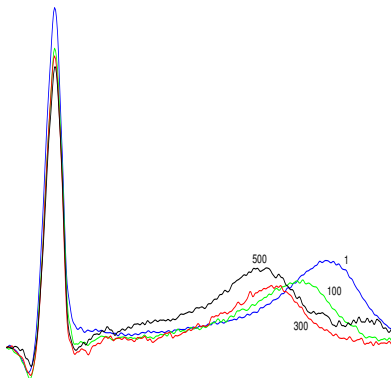
- delays ? \Rightarrow change of the conduction velocity
- amplitudes ? \Rightarrow change of the number of depolarized cells
- $x_i(t) = k_i s(\Phi(t)) + n_i(t)$
- scales (narrow, wide) ? \Rightarrow change of the synchronization of the cells
Repol./Depolarization
- mean shape (\Rightarrow notion of shape distance) ?

\Rightarrow unique framework

An example : characterize T Waves variations

One healthy and one ischemic subjects during exercise test. *ECG LEAD V5*

- Constant cycling speed with 50W increased by 25W every 2 minutes
- Healthy : RR 950ms (1) - RR 575 ms (500)
- Ischemic : RR 750ms (1) - RR 570 ms (500)



Model and properties

- based on the simple observation model (slightly different than before)

$$x_i(t) = k_i s\left(\frac{t - d_i}{\alpha_i}\right) + n_i(t) \text{ with } \alpha_i > 0; k_i > 0$$

k_i, α_i, d_i : amplitude coefficient, scaling factor, delay

- the normalized integrals are defined as (assuming positivity)

$$S(t) = \left(\int_0^t s(u) du \right) / \left(\int_0^T s(u) du \right)$$

$$X_i(t) = \left(\int_0^t x_i(u) du \right) / \left(\int_0^T x_i(u) du \right)$$

- for any value of t we get (increasing functions)

$$y = S(t) = X_i(t_i) \Leftrightarrow t = S^{-1}(y) \text{ with } t_i = \psi_i(t)$$

- providing the key relation

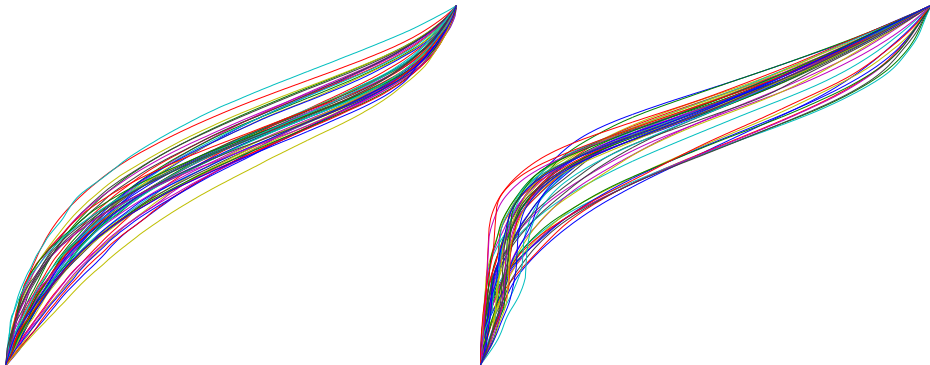
$$t_i = \alpha_i S^{-1}(y) + d_i$$

Model and properties

- in the discrete case $\mathbf{t}_i = [X_i^{-1}(0) X_i^{-1}(\delta_y) \cdots X_i^{-1}(1)]$
- the key relation is

$$\mathbf{t}_i = \alpha_i \mathbf{t} + d_i \mathbf{1} \quad (59)$$

- \Rightarrow linear function of the parameters (not the case in time)
- example on 50 ($i = 1 : 10 : 500$) roughly segmented T waves (\mathbf{t}_i 's)



Model and properties

- in the discrete case $\mathbf{t}_i = [X_i^{-1}(0) X_i^{-1}(\delta_y) \cdots X_i^{-1}(1)]$
- the key relation is

$$\mathbf{t}_i = \alpha_i \mathbf{t} + d_i \mathbf{1}$$

- \implies linear function of the parameters (not the case in time)

α_i 's, d_i 's, \mathbf{t} are estimated by computing $\mathbf{T} = [\mathbf{t}_1 \cdots \mathbf{t}_N] = \mathbf{V}\Sigma\mathbf{U}'$ (SVD)

k_i 's are then calculated by using the expression $\int_0^T x_i(u) du / \alpha_i$

Parameters estimation

- $\alpha_i, d_i, \mathbf{t}$ are of interest
- two stages estimation (\mathbf{t}) & (α_i, d_i) by zeroing the mean of each \mathbf{t}_i
- first estimation solve

$$\check{\mathbf{t}} = \arg \min_{\mathbf{t}} \left(\sum_i \|\mathbf{t}_i - \alpha_i \mathbf{t}\|^2 \right) \text{ subj } \mathbf{t}^T \mathbf{t} = 1 \quad (60)$$

equivalent to

$$\check{\mathbf{t}} = \arg \max_{\mathbf{t}} \mathbf{t}^T \mathbf{R} \mathbf{t} \quad (61)$$

where \mathbf{R} stands for the correlation matrix of the observations \mathbf{t}_i 's.

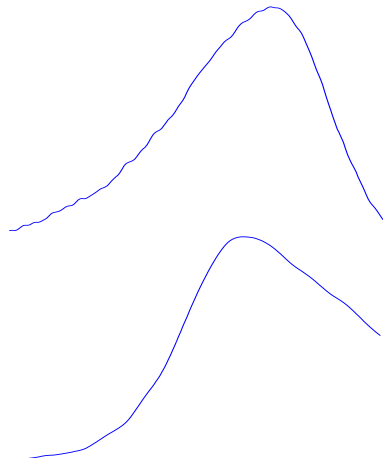
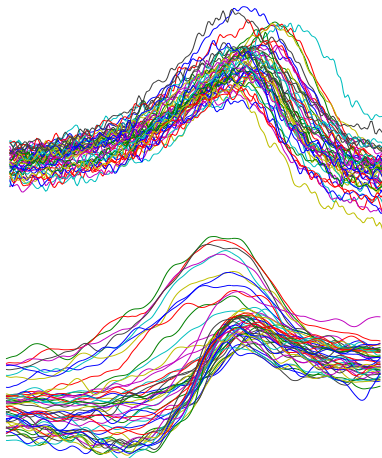
$\check{\mathbf{t}}$ is then the first column of \mathbf{V} such that $\mathbf{T} = [\mathbf{t}_1 \cdots \mathbf{t}_N] = \mathbf{V}\mathbf{\Sigma}\mathbf{U}'$ (SVD)

The α_i 's are computed from the SVD decomposition.

The mean m_i of each \mathbf{t}_i is in fact the center of gravity of $x_i(t)$ and is related to m_1 by $m_i = \alpha_i m + d_i$

Parameters estimation

- \check{t} plays the role of the "mean shape" (scale and shift invariant)
- \check{t} is in fact a weighted (> 0) average of increasing functions
 - ▶ qualified as inverse normalized integral
 - ▶ its derivative provides the positive temporal mean shape



Model and properties

- The first observation x_1 is taken as the reference \implies correction of the parameters

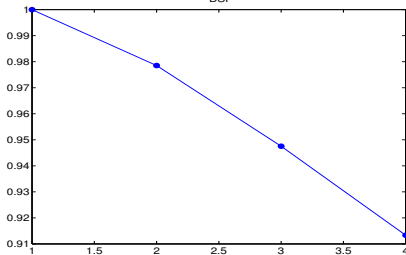
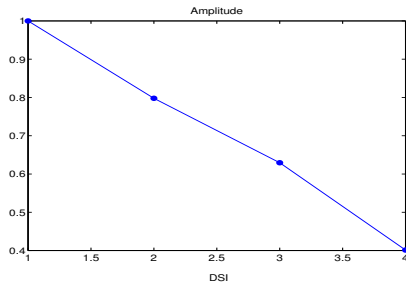
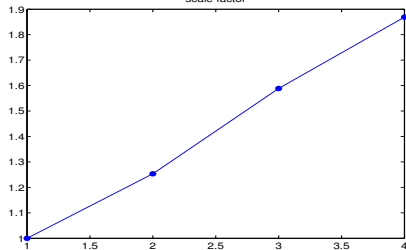
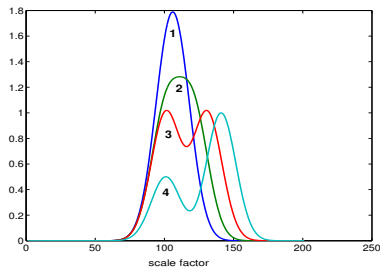
$$\alpha_i / \alpha_1, k_i / k_1, d_i - d_1$$

- Difference Shape Index (DSI) is performed on each couple of observations $((1;2), (1;3), \dots)$ by using previous SVD decomposition on $2 \times N$ \mathbf{T}_i matrices.

For each matrix \mathbf{T}_i the 2 singular values $\lambda_{1,i}$ and $\lambda_{2,i}$ (with $\lambda_{1,i} > \lambda_{2,i}$) allows :

$$0.5 < DSI_i = \frac{\lambda_{1,i}}{\lambda_{1,i} + \lambda_{2,i}} < 1$$

Simulation : different shapes

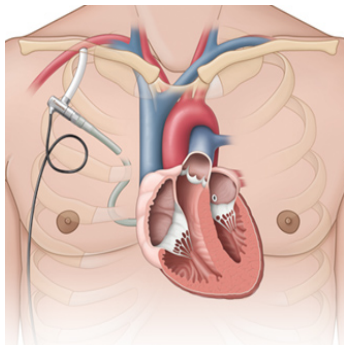


Goal and tools

Develop non invasive new strategies for ventricular continuous monitoring during continuous-flow Left Ventricular Assist Device (LVAD) therapy by exploring possible relationship between the **ventricular volume** and the **electrical myocardial activity**.

Assumption

LVAD \Rightarrow mechanical unloading of LV \Rightarrow changes of ventricular electrical activity



Changes of the ventricle blood volume \Rightarrow Influence on ECG :

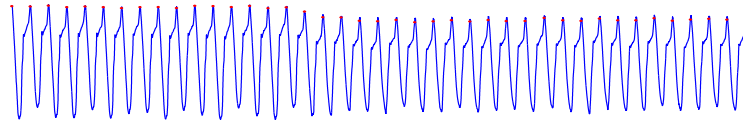
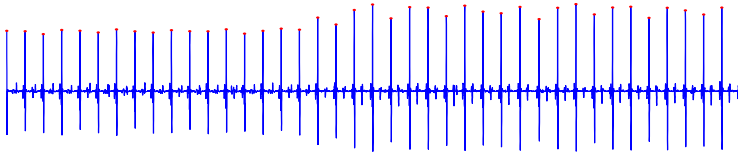
- Position of the heart in the chest
- Brody effect
- Change in thickness of the ventricular muscle wall
- **Effect of stretch of the muscle fibers on the AP (depolarization) [Franz]**

Initial Approach

R wave peak (RWP) values are compared to the corresponding (in time) Left Ventricular Volume (LVV) during pump speed changes.

Other ECG features

For each QRS : Amplitude, Difference Shape Index, scale factor, delay ?



Pump OFF

Pump ON

Experimental data collection

- 6 Pigs underwent LVAD implantation : (1-5) Gyro Centrifugal Pump 2 and (6) Circulite Synergy Micropump
 - (1-5) 2 females and 3 males 47 ± 8 Kg, (6) male 80 Kg.
 - For Gyro 1300-1700 rpm and for Circulite 12000-18000 rpm
 - ECG (DIII), systemic arterial pressure and left ventricular volume and pressure (transducer catheter) were recorded
-
- The R wave peak magnitude (RWP) and QRS complex extracted from each cycle.
 - LVV_{RWP} , measured at the time of RWP.
 - To reduce the effect of ventilation (pigs ventilated), 20 consecutive RWP and LVV_{RWP} are averaged over time : RWP_p and LVV_{RWP_p}

All Pigs

Correlation coefficients between left ventricular volume LVV_{RWP_p} and :

- R wave peak RWP_p
- Difference Shape Index (DSI)
- amplitude coefficient (AMP)
- Scale factor (SF) for the 6 pigs.
- p-value $\dagger < 0.001$, $\S NS$

LVV_{RWP_p} vs	1	2	3	4	5	6
RWP_p	-0.94 [†]	-0.96 [†]	-0.86 [†]	-0.84 [†]	-0.86 [†]	-0.84 [†]
DSI	-0.02 [§]	0.05 [§]	0.43 [†]	0.01 [§]	-0.09 [§]	-0.27 [§]
AMP	-0.54 [†]	-0.89 [†]	0.71 [†]	-0.48 [†]	-0.46 [†]	-0.89 [†]
SF	0.03 [§]	0.44 [†]	-0.83 [†]	-0.12 [§]	0.08 [§]	-0.29 [§]

Fig (6)

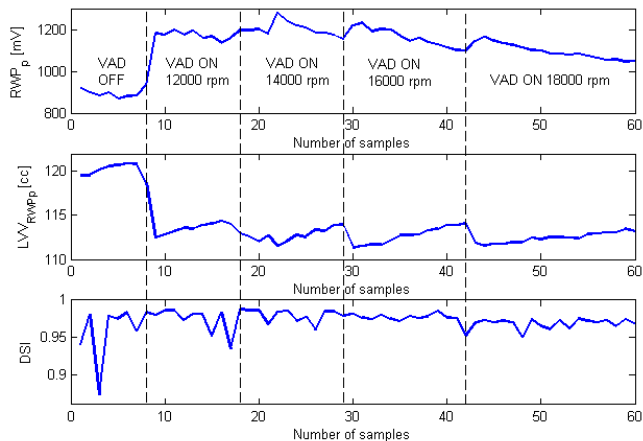


Figure – RWP_p , LVV_{RWP_p} and DSI profiles for pig (6) before VAD activation and during VAD speed change.

Conclusion

- Only RWP is consistently correlated with LVV
- Results are in agreement with Franz's work
- Relationship with other LVV features ?
- Possible non-invasive monitoring of the LVV
⇒ setting of the pump speed and prevent suction

Summary

- T or R waves analysis can be addressed as a shape analysis
- Delays, scales, mean shape estimated in the inverse normalized integrals domain
- Dynamic Time Warping as an alternative to find $\Phi(t)$