

# Adaptive Singular Value QRST Cancellation for the Analysis of Short Single Lead Atrial Fibrillation Electrocardiograms

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## Abstract

*Atrial fibrillation (AF) is the most commonly diagnosed sustained supraventricular arrhythmia in clinical practice and is characterized by uncoordinated atrial activation. The proper analysis and characterization of AF from surface ECG recordings requires to cancel out ventricular activity (i.e. QRS-T complexes). Some powerful methods exploit the spatial diversity of multi-lead ECG, however, their performance is seriously reduced in single-lead environments. For this latter case, techniques based on averaged beat subtraction (ABS) are the most widely used. However, these methods are very sensitive to QRS-T wave variations, thus, a high quality QRS-T cancellation template may be difficult to obtain when only short length recordings are available. To overcome these difficulties, a new QRS-T cancellation method based on singular value decomposition (SVD) of each single beat is presented. This methodology was tested and validated using a significative database with simulated and real AF recordings. The results showed that SVD is able to obtain a very accurate ventricular activity (VA) representation, thus providing high quality atrial activity (AA) extraction in short and single-lead AF recordings. Therefore, the inherent limitations in ABS techniques due to variations in the QRS-T shape could be avoided.*

## 1. Introduction

Atrial fibrillation (AF) is the most commonly diagnosed sustained arrhythmia in clinical practice and affects up to 1% of the general population [1]. Considering its prevalence with age, this arrhythmia affects up to 15% of the population older than 80 and has an incidence that doubles with each advancing decade [2, 3]. During AF, the coordinated activation of the atria is replaced by an uncoordinated atrial activation that leads to the deterioration of atrial mechanical function with a consequent increased risk of stroke and mortality[2].

The surface ECG provides a noninvasive way to study

AF mechanisms and to investigate the effects of remodelling and treatment on AF [4]. Some of the advantages of using the surface ECG include the ability to record data for a long period of time, the minimal cost and risks involved compared with invasive electrophysiological studies, and its reflection of the global activity in the atria and ventriculus during AF [4]. However, the proper analysis and characterization of AF from ECG recordings requires the extraction or cancellation of the signal components associated with ventricular activity (VA), that is, the QRS complex and the T-wave (QRS-T)[5]. Unfortunately, a number of facts hinder this operation. First, atrial activity (AA) presents in the surface ECG much lower amplitude than its ventricular counterpart. Additionally, both phenomena possess spectral distributions that notably overlap, rendering linear filtering solutions unsuccessful [6].

To date, several ways to extract the AA from surface ECG recordings have been presented. The most powerful techniques are those that exploit the spatial diversity of the multilead ECG [7, 8], such as the method that solves the blind source separation problem [6]. However, the performance of these techniques is seriously reduced when they analyze the early stages of AF, i. e., paroxysmal AF, because the recordings are usually obtained from a Holter system with no more than two or three electrodes. Such a reduced number of leads is not sufficient to exploit the spatial information of the ECG, rendering those techniques based on averaged beat subtraction (ABS) the main alternatives. However, these techniques are very sensitive to QRST wave variations, and the estimated AA can be affected by some QRS-T residues, which may be important owing to the low AA amplitude [9]. Additionally, in clinical practice the ECG consists of 10 seconds in length recordings and, therefore, high quality QRS-T cancellation template may be difficult to obtain [10].

Thereby, this work presents a new cancellation method based on singular value decomposition (SVD) of each single beat capable to cancel out the QRS-T complexes in short duration and single-lead AF electrocardiograms. Thus, the proposed SVD-based methodology tries to ex-

tract the basis signal corresponding to the ventricular activity by exploiting the mutual information contained on each ECG beat.

## 2. Materials

### 2.1. Database

The presented methodology was validated using a database composed of twenty 15 seconds-length simulated AF recordings and twenty 15 seconds-length AF ECGs. Simulated recordings allowed to compare the estimated and original AA, as it was known a priori. Synthesized AF signals were created from the combination of AA and VA, which were synthesized separately. The AA was generated from the smooth concatenation of successive TQ segments extracted from AF ECGs. The VA was synthesized from normal sinus rhythm ECGs, after P-wave cancellation [9].

In addition, to evaluate the suitability of the algorithm to be applied over real scenarios, which was the final purpose, real AF recordings were used. These signals were obtained from Holter systems of two leads (V1 and II), digitized at a sampling frequency of 128 Hz and a resolution of 12 bits. Lead V1 was the input signal of the AA estimation approach, as it was the signal with higher AA content.

### 2.2. Data preprocessing

The real ECG recordings were preprocessed in order to reduce noise, nuisance interferences and improve later analysis. Firstly, baseline wander was removed making use of bidirectional high pass filtering with 0.5 Hz cut-off frequency [11]. Secondly, high frequency noise was reduced with an eight order bidirectional IIR Chebyshev low pass filtering, whose cut-off frequency was 70 Hz [12]. Finally, powerline interference was removed through adaptive notch filtering, which preserves the ECG spectral information [13]. Moreover, these signals were upsampled to 1 kHz in order to improve time alignment accuracy for QRS-T complex subtraction.

## 3. Methods

### 3.1. AA extraction algorithm

The ECG signal presents a high degree of temporal redundancy which could be exploited in order to cancel VA. Indeed, the QRS-T waveform usually exhibits a recurrent pattern, although different QRS-T morphologies as well as minor variations in the QRS-T waveform may occur [9]. Thus, all R waves were firstly detected making use of the Pan & Tompkins technique [14]. Next, the  $i$ -th QRS-T complex start point was defined as  $s_i = r_i - 0.3 \cdot RR_{min}$ ,

being  $r_i$  the R peak wave event and  $RR_{min}$  the minimum R-R interval found in the ECG. The  $i$ -th QRS-T complex end point was selected as  $e_i = r_i + 0.7 \cdot RR_{min}$  [15]. Each QRS-T complex was assumed to be represented by a column vector of the matrix  $\mathbf{X} \in \mathfrak{R}^{L \times N}$ :

$$\mathbf{X} = [\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N] \quad (1)$$

where  $\mathbf{x}_i$  contains  $L$  samples of  $i$ -th complex, and  $N$  is the complex number in the analyzed ECG. Note that all beats were temporally aligned using its R peak timing.

The Singular Value Decomposition (SVD) of the matrix  $\mathbf{X}$  can be expressed as:

$$\mathbf{X} = \mathbf{U}\mathbf{S}\mathbf{V}^T \quad (2)$$

where  $\mathbf{U} \in \mathfrak{R}^{L \times N}$  is a unitary matrix so that  $\mathbf{U}\mathbf{U}^T = \mathbf{I}$ ,  $\mathbf{S} \in \mathfrak{R}^{N \times N}$  is a diagonal matrix, and  $\mathbf{V} \in \mathfrak{R}^{N \times N}$  fulfills  $\mathbf{V}\mathbf{V}^T = \mathbf{I}$ . The matrix  $\mathbf{U}$  contains the  $N$  normalized principal components of  $\mathbf{X}$ , so that the columns of  $\mathbf{U} = [\mathbf{u}_1 \dots, \mathbf{u}_N]$  are the eigenvectors of  $\mathbf{X}$ , and their cross-correlations are nulls. The matrix  $\mathbf{S}$  contains the amplitude coefficients corresponding to the  $N$  principal components of  $\mathbf{X}$ . These coefficients are called eigenvalues or singular values and are sorted in descending order. Thus, the  $N$  non-normalized principal components can be obtained as the columns of the matrix  $\mathbf{P} = \mathbf{U}\mathbf{S}$ , and can be interpreted as follows:

- The most significant component is related to the main QRS-T waveform.
- Subsequently, there are several components related to AA.
- The remaining components correspond to noise.

Atrial activity can be viewed as being uncoupled to VA. Thus, each observed beat can be modelled as a sum of atrial activity ( $\mathbf{X}_{AA}$ ) and ventricular activity ( $\mathbf{X}_{VA}$ ) [7]:

$$\mathbf{X} = \mathbf{X}_{AA} + \mathbf{X}_{VA} \quad (3)$$

Hence, the first principal component, which is called  $\mathbf{t}$ , can be used as QRS-T template to cancel out VA. However, since considerable R peak amplitude differences between each individual QRS-T complex and template were found,  $\mathbf{t}$  amplitude was individually adapted to each beat:

$$\mathbf{t}_i = \frac{QR_i}{QR_t} \cdot \mathbf{t} \quad (4)$$

where  $QR_i$  and  $QR_t$  are the distances between the  $Q$  and  $R$  points of the  $i$ -th complex and template, respectively. Thus, the AA estimation ( $\hat{\mathbf{X}}_{AA}$ ) was obtained as:

$$\hat{\mathbf{X}}_{AA} = \mathbf{X} - \mathbf{T} \quad (5)$$

being  $\mathbf{T}$  the matrix constituted by the column vectors  $\mathbf{t}_i$ :

$$\mathbf{T} = [\mathbf{t}_1, \mathbf{t}_2, \dots, \mathbf{t}_N] \quad (6)$$

### 3.2. Performance assessment

The AA estimation performance in simulated recordings was computed by comparing the estimated and original AA in terms of Pearson correlation index. This coefficient measures the similarities between two signals, and becomes 1 in the case of perfect matching and 0 in the case of completely different and non-dependent signals.

An additional performance parameter was also introduced to evaluate AA estimation in real recordings. Thus, ventricular activity reduction (VAR) was defined as the ratio between the spectral power concentrations of the original ECG and obtained AA in 0.5-2.5 Hz band:

$$VAR = \sum_{f=0.5Hz}^{2.5Hz} \frac{P_x(f)}{P_{x_{AA}}(f)} \quad (7)$$

where  $P_x$  and  $P_{x_{AA}}(f)$  are the ECG and estimated AA power spectrums, respectively.

## 4. Results

The proposed methodology was firstly applied to the simulated AF signals. The Pearson correlation index obtained for each recording is detailed in Table 1, being  $0.92 \pm 0.07$  in average. The estimated AA quality is illustrated in Figure 1, which shows the ECG, original and estimated AA corresponding to a typical recordings (P8). As can be appreciated, the estimated AA nearly matches the original AA, without any QRS-T residue.

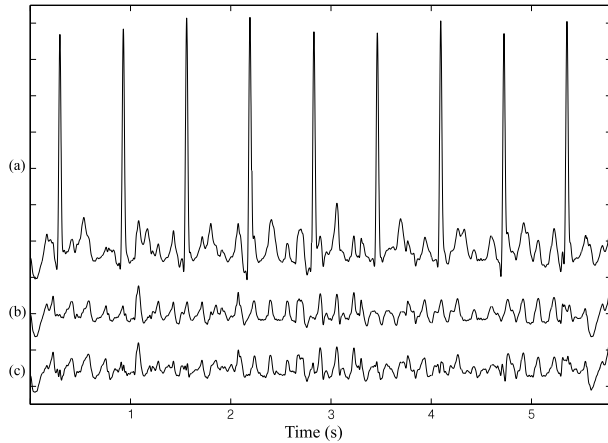


Figure 1. (a) Simulated 6 seconds-length ECG signal. (b) Real AA. (c) Estimated AA.

Secondly, the same methodology was applied to the recordings obtained from real AF patients. In this case, the VAR parameter was used to evaluate the proposed method performance. The VAR value obtained for each patient is shown in Table 2, being  $12.14 \pm 1.99$  in average. Taking

into account that AA presents much less amplitude than VA, these results indicate that the proposed approach is able to estimate AA free from QRS-T residues. Analyzing real AF recordings, the AA extraction quality is illustrated in Figure 2. Again, it can be observed that AA can be extracted without QRS-T residues.

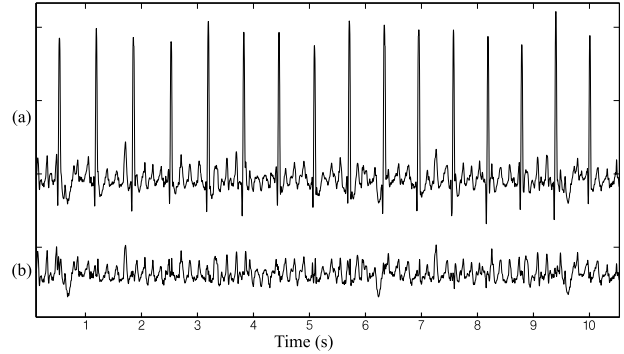


Figure 2. (a) Real 11 seconds-length ECG signal. (b) Estimated AA.

## 5. Discussion and conclusions

The AA estimation in short AF episodes requires the implementation of QRS-T cancellation techniques for single-lead ECGs, rendering ABS techniques the unique existing solutions. In this work, an alternative method based on SVD concepts has been presented. The proposed methodology tries to overcome the limitations of the ABS strategies, which are highly sensitive to QRS-T morphology variations.

The results have shown that SVD is able to obtain a very accurate representation of VA, thus providing high quality AA extraction in short and single-lead AF recordings. Thereby, it can be concluded that a high accuracy QRS-T template is obtained, since dynamics in the QRS-T waveform are considered. Thus, the inherent limitations in ABS techniques could be avoided.

To date, other two methods have been proposed to extract AA from short duration and single-lead AF recordings. Thus, Castells et. al. [9] presented a new method based on PCA concepts, however, the correlation index between estimated and original AA was  $0.774 \pm 0.106$  in average. Recently, Lemay et. al. [10] has presented other method, in which cancellation during the JQ intervals is carried out using dominant T and U wave methodology and AA estimation during the QRS intervals is performed making use of a weighted sum of sinusoids. Because of the estimated and real AA during the QRS interval could be very different when AA is very irregular, it can be considered that this new proposed method can obtain a more accurate AA estimation.

Table 1. Correlation values obtained for the simulated AF ECGs.

<b>S1</b>	<b>S2</b>	<b>S3</b>	<b>S4</b>	<b>S5</b>	<b>S6</b>	<b>S7</b>	<b>S8</b>	<b>S9</b>	<b>S10</b>
0.91	0.87	0.91	0.88	0.95	0.89	0.87	0.91	0.89	0.84
<b>S11</b>	<b>S12</b>	<b>S13</b>	<b>S14</b>	<b>S15</b>	<b>S16</b>	<b>S17</b>	<b>S18</b>	<b>S19</b>	<b>S20</b>
0.82	0.84	0.93	0.94	0.90	0.88	0.80	0.82	0.94	0.92

Table 2. VAR values of the estimated AA from real AF recordings.

<b>R1</b>	<b>R2</b>	<b>R3</b>	<b>R4</b>	<b>R5</b>	<b>R6</b>	<b>R7</b>	<b>R8</b>	<b>R9</b>	<b>R10</b>
12.58	9.32	13.42	15.24	10.61	13.71	14.50	8.81	9.11	13.14
<b>R11</b>	<b>R12</b>	<b>R13</b>	<b>R14</b>	<b>R15</b>	<b>R16</b>	<b>R17</b>	<b>R18</b>	<b>R19</b>	<b>R20</b>
11.20	13.38	13.63	13.42	14.58	13.33	14.38	9.59	11.96	11.68

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