Order Statistics Correlation Coefficient as a Novel Association Measurement With Applications to Biosignal Analysis

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Abstract—In this paper, we propose a novel correlation coefficient based on order statistics and rearrangement inequality. The proposed coefficient represents a compromise between the Pearson's linear coefficient and the two rank-based coefficients, namely Spearman's rho and Kendall's tau. Theoretical derivations show that our coefficient possesses the same basic properties as the three classical coefficients. Experimental studies based on four models and six biosignals show that our coefficient performs better than the two rank-based coefficients when measuring linear associations; whereas it is well able to detect monotone nonlinear associations like the two rank-based coefficients. Extensive statistical analyses also suggest that our new coefficient has superior anti-noise robustness, small biasedness, high sensitivity to changes in association, accurate time-delay detection ability, fast computational speed, and robustness under monotone nonlinear transformations.

Index Terms—Atrial fibrillation, atrial flutter, concomitant, Kendall's tau, nonlinear association measure, order statistics, Pearson's coefficient, rearrangement inequality, Spearman's rho.

I. INTRODUCTION

THERE has been great interest in measuring the association between two time series, with application in many areas including biosignal analysis. Association can be considered as the strength of relationship between the two time series. A measure of association should be large and positive if there is a high probability that large (small) values of one time series are associated with large (small) values of another. On the other hand, if the direction is inverse, namely, large (small) values of one time series occur in conjunction with small (large) values of another time series, the measure should be large and negative [1]. A multitude of methods have been used in the literature of biosignal processing for many years to measure the association between two time series. Among these measures the Pearson's linear correlation coefficient [2]–[4], Spearman's

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rank-based coefficient (*Spearman's rho*) [5] and Kendall's concordance coefficient (*Kendall's tau*) [5] are perhaps the most widely used [6]. The linear correlation coefficient is appropriate mainly for indicating linear associations [1], while the other two measures are invariant under linear or nonlinear increasing monotone transformations [5]. Some authors have employed the average amount of mutual information (AAMI) to measure the association between two biosignals [7], [8], others used nonlinear regression coefficient (denoted by h^2) [9] or contingency table based methods (Cramer V) for such purpose [10].

There are many advantages and disadvantages to the measures mentioned before. Linear correlation coefficient is very fast, however, it will yield misleading results if nonlinearity is involved in the system [10]. On the other hand, the two rank correlation coefficients, Spearman's rho and Kendall's tau, are not as powerful and as fast as Pearson's coefficient when measuring linear associations between biosignals; nevertheless they are independent of increasing nonlinear transformations which makes them suitable for many nonlinear cases [1], [5], [6]. Despite their robustness for nonlinear association measurements, the values of h^2 and V are between 0 and 1, meaning their inability of distinguishing positive associations from negative associations. Furthermore, the computational load of AAMI and h^2 are rather heavy, which makes them inappropriate in cases when high computational speed is mandatory.

To overcome the problems of the existing measures of association in different *a priori* unknown situations, we propose a novel measure called order statistic correlation coefficient which possesses the following advantages: 1) it can discriminate positive associations from negative associations (admissible range [-1,1]); 2) its time complexity is of order $O(N\log(N))$, a little slower than Pearson's coefficient but much faster than Kendall's tau, the nonlinear regression coefficient h^2 , and AAMI; 3) it has small biasedness in both linear and nonlinear scenarios; 4) it is sensitive to changes of degree of association; and 5) it possesses certain robustness under increasing nonlinear transformations.

In Section II, we will give the definition and properties of our new order statistic coefficient as well as the other three classical indices of association. Section III depicts the models and performance evaluation strategy we use in this study. In Section IV, we present the simulated signals and the associated results of four models used in our investigation. Section V is devoted to discussions and interpretations of our new method. Finally, in Section VI, we draw our conclusions on the novel order statistics correlation coefficient.

II. ORDER STATISTICS CORRELATION COEFFICIENT

A. Definition and Properties

Let (x_i, y_i) , $i = 1, \ldots, N$, be two time series of length N. Rearranging pairwise the two time series with respect to the magnitudes of x, we get two new series denoted by $(x_{(i)}, y_{[i]})$, where $x_{(1)} \leq \cdots \leq x_{(N)}$ are called the *order statistics* of x and $y_{[1]}, \ldots, y_{[N]}$ the associated *concomitants* [11]–[13]. Reversing the roles of x and y, we also define the order statistics of y and the corresponding concomitants which are denoted by $y_{(1)}, \ldots, y_{(N)}$ and $x_{[1]}, \ldots, x_{[N]}$, respectively. As proposed by Xu et al. [14], the order statistics correlation coefficient can be defined as

$$r_X(x,y) \stackrel{\Delta}{=} \frac{\sum_{i=1}^{N} (x_{(i)} - x_{(N-i+1)}) y_{[i]}}{\sum_{i=1}^{N} (x_{(i)} - x_{(N-i+1)}) y_{(i)}}.$$
 (1)

Theorem 1: The order statistics correlation coefficient has the basic properties of a correlation coefficient, as follows:

- 1) $-1 \le r_X \le 1$;
- 2) $r_X(x, y)$ attains +1(-1) when x and y are in strict increasing (decreasing) relationship;
- 3) $r_X(x', y') = r_X(x, y)$ for $x' = k_x x + \text{const}_x$ and $y' = k_y y + \text{const}_y$, where $k_x > 0$ and $k_y > 0$;
- 4) if x and y are mutually independent and each is independent identically distributed (IID), the expectation $E\{r_X(x,y)\}=0$ when $N\to\infty$.
- According to the rearrangement inequality [15], it follows that:

$$\sum_{i=1}^{N} x_{(N-i+1)} y_{(i)} \le \sum_{i=1}^{N} x_{(i)} y_{[i]} \le \sum_{i=1}^{N} x_{(i)} y_{(i)}$$
 (2)

and

$$\sum_{i=1}^{N} x_{(N-i+1)} y_{(i)} \le \sum_{i=1}^{N} x_{(N-i+1)} y_{[i]} \le \sum_{i=1}^{N} x_{(i)} y_{(i)}.$$
 (3)

Subtracting (3) by (4) and dividing the difference by $\sum (x_{(i)} - x_{(N-i+1)})y_{(i)}$, we have $-1 \le r_X \le 1$, hence the result.

- 2) Assume $y_i = \phi(x_i)$, i = 1, ..., N. If $\phi(\bullet)$ is a strict increasing function, we have $y_{[i]} = y_{(i)}$ for all i. Substituting this into (1), we have $r_X = 1$; and similarly $r_X = -1$ if $\phi(\bullet)$ is a strict decreasing function.
- 3) Substituting x' and y' into (1), we have

$$r_X(x',y') = \frac{\sum \left(x'_{(i)} - x'_{(N-i+1)}\right) y'_{[i]}}{\sum \left(x'_{(i)} - x'_{(N-i+1)}\right) y'_{(i)}}$$

$$= \frac{k_x k_y \sum \left(x_{(i)} - x_{(N-i+1)}\right) y_{[i]} + A}{k_x k_y \sum \left(x_{(i)} - x_{(N-i+1)}\right) y_{(i)} + A}$$
(4)

where $A = k_x \operatorname{const}_y \sum [x_{(i)} - x_{(N-i+1)}] = 0$. Hence, we have $r_X(x', y') = r_X(x, y)$.

4) Denote the numerator and denominator of (1) by U and V, respectively. An application of the Delta method [16] yields

$$E(r_X) = \frac{E(U)}{E(V)} + O(N^{-1}).$$
 (5)

In order to prove that $E(r_X) = 0$ (with N large), it is adequate to show that E(U) is null when the assumptions are satisfied. Imposing the independence assumption of x and y, we have

$$E(U) = \sum \left[E\left(x_{(i)}\right) - E\left(x_{(N-i+1)}\right) \right] E\left(y_{[i]}\right). \quad (6)$$

It is known [6] that when y is IID, the probability density function (PDF) of $y_{[i]}$ is

$$g_{[i]}(y) = \int_{-\infty}^{+\infty} f(y|x) f_{(i)}(x) dx$$
 (7)

where $g_{[i]}(y)$ denotes the PDF of $y_{[i]}$, f(y|x) the conditional PDF of y given x and $f_{(i)}(x)$ the PDF of $x_{(i)}$. The conditional PDF f(y|x) degenerates to f(y) if x and y are independent. Then, we have

$$E(y_{[i]}) = \int yg_{[i]}(y)dy$$

$$= \int y \int f(y|x)f_{(i)}(x)dxdy$$

$$= \int yf(y)dy \int f_{(i)}(x)dx$$

$$= \int yf(y)dy$$

$$= E(y).$$
(8)

Substituting (8) into (6), we have E(U)=0 and, thus, $E(r_X)=0$ to the order of $O(N^{-1})$.

B. Estimation of Correlation Coefficient in Normal Case

It will be shown in the following theorem that for samples from a bivariate normal population with correlation coefficient ρ , r_X is an asymptotically unbiased estimation of ρ .

Theorem 2: If (x_i, y_i) , $i = 1, \ldots, N$ is a pair of IID time series from a bivariate normal distribution with correlation coefficient ρ , then $\lim_{N \to \infty} E\{r_X(x,y)\} = \rho$.

Proof: Without loss of generality, we assume that both x

Proof: Without loss of generality, we assume that both x and y have zero mean and unity variance. The order statistics $x_{(i)}$ and $y_{(i)}$ can be expressed as

$$x_{(i)} = \mu_i + \varepsilon_i$$

$$y_{(i)} = \mu_i + \delta_i$$
 (9)

where $\mu_i = E(x_{(i)}) = E(y_{(i)})$ and $E(\varepsilon_i) = E(\delta_i) = 0$. It is obvious that ε_i and δ_i have identical distributions, and hence we have

$$E\left(\varepsilon_i^2\right) = E\left(\delta_i^2\right). \tag{10}$$

The symmetry of the normal distribution yields [17], [18]

$$\mu_{i} = -\mu_{N-i+1}$$

$$E\left(\varepsilon_{i}^{2}\right) = E\left(\varepsilon_{N-i+1}^{2}\right). \tag{11}$$

It follows that the concomitants associated with $x_{(i)}$ can be written as [11]

$$y_{[i]} = \rho x_{(i)} + z_i \tag{12}$$

where $x_{(i)}$ and z_i are mutually independent, the latter being IID normal with mean zero and variance $1 - \rho^2$. Given (9)–(12), the numerator in (1) can be expressed as

$$U = 2\sum x_i y_i - \sum (\rho \mu_i + \rho \varepsilon_i + z_i)(\varepsilon_i + \varepsilon_{N-i+1}). \quad (13)$$

From (11), we have

$$\sum \mu_i(\varepsilon_i + \varepsilon_{N-i+1}) = -\sum \mu_{N-i+1}(\varepsilon_i + \varepsilon_{N-i+1}). \quad (14)$$

Replacing N - i + 1 by j in (14) leads to

$$\sum \mu_i(\varepsilon_i + \varepsilon_{N-i+1}) = -\sum \mu_j(\varepsilon_j + \varepsilon_{N-j+1}) = 0. \quad (15)$$

Hence, (13) can be further simplified by

$$U = 2\sum x_i y_i - \sum (\rho \varepsilon_i + z_i)(\varepsilon_i + \varepsilon_{N-i+1}). \tag{16}$$

Taking expectations on both sides of (16) and applying the mutual independence facts mentioned before, we have

$$E(U) = 2N\rho \left\{ 1 - \frac{\sum E\left(\varepsilon_i^2\right) + \sum E(\varepsilon_i \varepsilon_{N-i+1})}{2N} \right\}. \quad (17)$$

An application of Cauchy–Schwarz inequality [15] and the fact $E(\varepsilon_i \varepsilon_{N-i+1}) > 0$ [11] to (14) yields

$$\begin{cases} 2N\rho > E(U) > 2N\rho \left[1 - \frac{\sum E\left(\varepsilon_{i}^{2}\right)}{N}\right], & \text{when } \rho \geq 0\\ 2N\rho < E(U) < 2N\rho \left[1 - \frac{\sum E\left(\varepsilon_{i}^{2}\right)}{N}\right], & \text{when } \rho < 0. \end{cases}$$

$$\tag{18}$$

It can be easily verified that the following two identities hold for any two real numbers a and b:

$$ab = \frac{1}{2} \left[a^2 + b^2 - (a - b)^2 \right]$$
$$-ab = \frac{1}{2} \left[a^2 + b^2 - (a + b)^2 \right]. \tag{19}$$

Substituting (9)–(12) and (16) into the denominator in (1), we arrive at

$$V = \sum (x_i^2 + y_i^2) - \frac{1}{2} \sum (\varepsilon_i - \delta_i)^2 - \frac{1}{2} \sum (\varepsilon_i + \delta_{N-i+1})^2.$$
(20)

Applying elementary inequalities and taking expectation on both sides of (17), we have

$$2N\left[1 - \frac{2\sum E\left(\varepsilon_i^2\right)}{N}\right] \le E(V) < 2N. \tag{21}$$

It is implied in [18] that

$$\lim_{N \to \infty} \frac{\sum E\left(\varepsilon_i^2\right)}{N} = 0. \tag{22}$$

Substituting (15), (18), and (19) into (5) and letting N tend to infinity, we have $\lim_{N \to \infty} E\{r_X(x,y)\} = \rho$, hence, the result.

According to Theorem 2, we can estimate ρ using order statistics correlation coefficient by the following estimator:

$$\hat{\rho}_X = r_X. \tag{23}$$

C. Comparison With Three Classical Correlation Coefficients

As defined in Section II-A, $x_{(1)},\ldots,x_{(N)}$ are the order statistics of the time series x_1,\ldots,x_N . Suppose x_j is at the kth position in the sorted series $x_{(1)},\ldots,x_{(N)}$, the number $1 \leq k \leq N$ is termed the rank of x_j and is denoted by $p_j(=k)$. Similarly, we can get the rank of y_j denoted by q_j . Such operation of obtaining the ranks of all elements in a series is called ranking [5]. Let (x_i,y_i) and (x_j,y_j) with $i=1,\ldots,N$ and $j=i+1,\ldots,N$ be two data-pairs from the original time series. If p_j-p_i and q_j-q_i have the same sign, we say that the two data-pairs are concordant, otherwise, we say that they are discordant [5]. Let P stand for the number of concordant pairs and Q the number of discordant pairs, it follows that P+Q=N(N-1)/2. Let $\overline{x}, \overline{y}, \overline{p}$, and \overline{q} be the arithmetic averages of x, y, p, and q, respectively, Pearson's correlation coefficient (r_P) , Spearman's rho (r_S) , and Kendall's tau (r_K) are defined as follows [1], [5]:

$$r_P(x,y) \stackrel{\triangle}{=} \frac{\sum_{i=1}^{N} (x_i - \overline{x})(y_i - \overline{y})}{\sqrt{\sum_{i=1}^{N} (x_i - \overline{x})^2 \sum_{i=1}^{N} (y_i - \overline{y})^2}}$$
(24)

$$r_S(x,y) \stackrel{\Delta}{=} 1 - \frac{6\sum_{i=1}^{N} (p_i - q_i)^2}{N^3 N}$$
 (25)

$$r_K(x,y) \stackrel{\Delta}{=} \frac{2(P-Q)}{N(N-1)}.$$
 (26)

If x and y are bivariate Gaussian with correlation coefficient ρ , three reasonable estimators of ρ can be constructed as follows [5]:

$$\hat{\rho}_P = r_P$$

$$\hat{\rho}_S = 2\sin\left(\frac{1}{6}\pi r_S\right)$$

$$\hat{\rho}_K = \sin\left(\frac{1}{2}\pi r_K\right).$$
(27)

For brevity, we will drop the circumflex throughout and use ρ_{ξ} , $\xi = X, P, S, K$ to denote the four estimators. It is easy to verify that 1) ρ_{ξ} are monotonic with r_{ξ} ; 2) three critical values $\{-1,0,+1\}$ are invariant under the transformations ρ_{ξ} ; and $3)\rho_{\xi}$ is also limited within [-1,+1].

III. MODELS OF ASSOCIATION AND PERFORMANCE EVALUATION

In this section, we propose three linear models and one non-linear model to model the linear and nonlinear association between two time series. Several indices will also be proposed to evaluate the performance of our order statistics correlation coefficient in comparison with the other classical correlation coefficients, in terms of their abilities to estimate the associations between time series. In each model, a time series x(i) is derived from a pure signal s(i), and another signal y(i) is obtained as a combination of the transformed pure signal and a white noise, n(i). In all these models, the time index i runs from 1 to 1000.

A. Models of Association

1) Linear Model 1 (LM1): LM1 is constructed as

$$x(i) = s(i)$$

$$y(i) = s(i) + \alpha \cdot n(i)$$
(28)

where $\alpha \in [0,1]$ is increased from 0 to 1 with a step $\Delta \alpha = 0.1$ to control the signal-to-noise ratio (SNR). With increasing α , the association between x and y becomes smaller and smaller, which means that r_{ξ} (ρ_{ξ}) should have a decreasing relationship with α . For a fixed α , the greater the magnitude of $E(\rho_{\xi})$, the better its performance in the context of noise robustness.

2) Linear Model 2 (LM2): LM2 is a regression model of the form [14]

$$x(i) = s(i)$$

$$y(i) = \rho \cdot s(i) + \sqrt{1 - \rho^2} \cdot n(i)$$
(29)

where $\rho \in [-1,1]$ with a step $\Delta \rho = 0.01$ characterizing the linear association. It follows by straightforward calculation that $E(\rho_P) = \rho$ for any distribution of s(i). Unfortunately, the property of unbiasedness does not hold for the other three estimators ρ_X , ρ_S , and ρ_K except for the bivariate normal case. The aim of this model is to compare the biasedness of these three biased estimators as well as their power to discriminate different ρ 's.

3) Linear Model 3 (LM3): LM3 is similar to LM1 except for a time delay $\Delta=30$ introduced in channel y, as follows:

$$x(i) = s(i)$$

$$y(i) = s(i - \Delta) + \alpha \cdot n(i).$$
 (30)

4) Nonlinear Model (NM): NM is a nonlinear model used to study the effect of nonlinear transformations to the signals on the four coefficients, as follows [14]:

$$x(i) = T_x \left[\beta \cdot s(i) \right]$$

$$y(i) = T_y \left[\beta \left\{ \rho \cdot s(i) + \sqrt{1 - \rho^2} \cdot n(i) \right\} \right]$$
(31)

where $T_x[\bullet]$ and $T_y[\bullet]$ are two increasing nonlinear functions. The parameter $\beta=2,4,6,8,10$ is used to control the extent of nonlinearity (greater value of β corresponding to stronger nonlinearity), while ρ has the same meaning as in LM2.

B. Performance Evaluation

Several methods are used to evaluate the performance of r_{ξ} under each of the four models previously mentioned.

- 1) Noise Robustness: Under LM1, we compare the decreasing rates of $E(\rho_{\mathcal{E}})$ with the increase of α .
- 2) Attenuation Measurement: We propose two indices called absolute attenuation (AAT) and relative attenuation (RAT) to measure the extent of biasedness, defined as

$$AAT_{\xi} = \frac{\int_{-1}^{1} \left| \overline{\rho}_{\xi}(\rho) - \rho \right| d\rho}{\int_{-1}^{1} \int_{-1}^{1} d\rho d\rho_{\xi}} = \frac{1}{4} \int_{-1}^{1} \left| \overline{\rho}_{\xi}(\rho) - \rho \right| d\rho \quad (32)$$

and

$$RAT_{\xi} = \frac{\int_{-1}^{1} \left| \overline{\rho}_{\xi}(\rho) - \overline{\rho}_{\xi}^{(L)}(\rho) \right| d\rho}{\int_{-1}^{1} \int_{-1}^{1} d\rho d\rho_{\xi}} = \frac{1}{4} \int_{-1}^{1} \left| \overline{\rho}_{\xi}(\rho) - \overline{\rho}_{\xi}^{(L)}(\rho) \right| d\rho$$
(33)

where $\overline{\rho}_{\xi}^{(L)}(\rho)$ is the mean of ρ_{ξ} under LM2. AAT is to measure the extent of biasedness under LM2, while RAT is to measure the biasedness caused by the nonlinearity involved in NM. For Pearson's coefficient, we have $\overline{\rho}_P=\rho$, thus $\mathrm{AAT}_P=0$. For the other three coefficients, AAT relates positively to their biasedness under LM2. It is clear from (30) that under LM2 $\mathrm{RAT}_{\xi}=0$ and under NM, a smaller RAT means lesser effect of the nonlinearity and hence better robustness.

3) Sensitivity to Changes in ρ : We employ another index called sensitivity ratio (SR) [4] to test the sensitivity of ρ_{ξ} to changes in ρ . For this purpose the Fisher's z-transformation of ρ_{ξ} , denoted as z_{ξ}

$$z_{\xi} = \tanh^{-1} \rho_{\xi} = \frac{1}{2} \log_e \frac{1 + \rho_{\xi}}{1 - \rho_{\xi}}.$$
 (34)

After such transformation, which maps [-1, +1] to $(-\infty, +\infty)$, the resultant z_{ξ} follows approximately normal distributions with constant variances (i.e., independent of the means) [2], [4]. Given two distinct ρ_1 and ρ_2 ($\rho_2 > \rho_1$), we have two sets of coefficients $\rho_{\xi 1}$ and $\rho_{\xi 2}$, and their respective Fisher's z-transformation, $z_{\xi 1}$ and $z_{\xi 2}$. SR is then defined as

$$SR_{\xi} = \frac{\overline{z}_{\xi 2} - \overline{z}_{\xi 1}}{\sqrt{v_{z1}^2 + v_{z2}^2}}$$
 (35)

where \overline{z}_{ξ} and v_{ξ} denote the mean and standard deviation of z_{ξ} , respectively. SR measures the ability of ρ_{ξ} to detect the changes of underlying ρ . A greater value of SR indicates better discrimination sensitivity. SR is computed for the results of the linear model LM2 and the nonlinear model NM.

4) Time Complexity Measurement: We analyze the time complexities of r_{ξ} in the language of big Oh. We also estimate the relationship between computational loads of r_{ξ} versus the length of signal N from 100 to 1000 with a step $\Delta N = 100$.

IV. COMPARISON ON RESULTS FOR SIMULATED AND REAL BIOSIGNALS

Signals derived from biological processes fall into two main categories: deterministic and stochastic signals. The former are those that can be described by explicit mathematical relationships; whereas the latter can be described only in statistical terms. The deterministic group is further subdivided into periodic, semi-periodic, and transient signals; the stochastic group is subdivided into stationary and nonstationary signals [19], [20]. Recently, there has been growing evidence indicating that many biosignals exhibit long range power-law correlations [21]. A stochastic process is said to have long range correlation if its autocorrelation function $R(k) \sim k^{2H-2}$ as $k \to \infty$, where 0 < H < 1 is the Hurst parameter. The corresponding power spectral density is proportional to $f^{-(2H-1)}$ [22]. In order to evaluate the feasibility of r_X in association studies, several simulated and real biosignals with respect to six types of previously mentioned biosignals are employed for investigation. For notational convenience, the six signals are denoted uniformly as s_{ζ} , $\zeta = p, h, t, a, e, l$, which represent periodic, semi-periodic, transient, stationary, nonstationary, and long-range-correlated signals. A number of 1000 independent white Gaussian noise ($\mu = 0$ and $\sigma^2 = 1$) are generated with a sampling rate of 1000 Hz to serve as noise in the linear and nonlinear models. Due to the 1000 noise involved, each r_{ε} becomes a random variable and has a distribution, which allows us to perform statistical analysis. Under each model, two channels of signals x and y are generated from s_{ζ} and the 1000 episodes of white noise. Four sets of correlation coefficients between x and y are then computed for comparative study.

A. Simulated and Real Biosignals

As remarked before, the following six representative biosignals are included in our study:

- 1) sin wave $s_P(i)$ of frequency 5 Hz emulating periodic biosignals;
- 2) real bipolar intra-atrial flutter signal $s_h(i)$ recorded during electrophysiological procedure [23];
- 3) atrial action potential waveform $s_t(i)$ generated from a mathematical model [24];
- 4) episode of alpha wave $s_a(i)$ simulated from a random Gaussian noise filtered by a band-pass Butterworth filter with passband 8 to 12 Hz [25];
- 5) second of real EEG signal $s_e(i)$ (sampling rate 256 Hz) from a dataset provided by University of Tuebingen for BCI Competition 2003 [26], [27];
- 6) segmentation $s_l(i)$ of an artificial time series $s_{lf}(i)$ exhibiting long range correlation with Hurst parameter H = 0.9 [21], [28], [29].

Fig. 1 illustrates the six biosignals. All the first four signals contain 1000 samples (N=1000). The EEG signal $s_e(i)$ is up-sampled from 256 to 1000 Hz by linear interpolation. As for the long-range-correlated signal $s_{lf}(i)$ containing 2^{17} samples, we use the first 1000 samples as $s_l(i)$ for association analysis. The whole time series $s_{lf}(i)$ is used when we demonstrate the capability of r_X for estimating the Hurst parameter H. After these manipulations, all the six original biosignals s_ζ , $\zeta=p,h,t,a,e,l$ can be considered of duration 1 s. Without loss of generality (property c), s_ζ are normalized to have mean zero and variance unity before feeding them into the four models described in Section III-B.

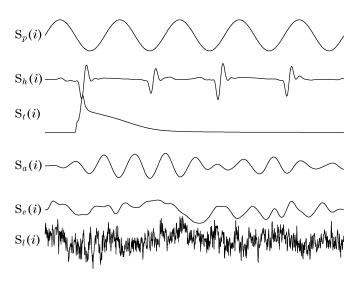


Fig. 1. Illustration of the four simulated signals and two real signals.

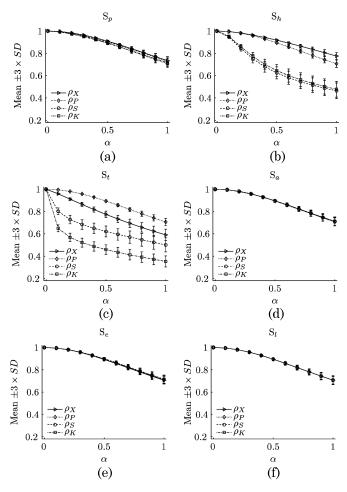


Fig. 2. Noise robustness comparison of simulation results under LM1.

B. Comparative Study Under Linear Model LM1

The results under LM1 are shown in Fig. 2. It is clear that $\overline{\rho}_{\xi}(\alpha)$ drops with increasing of α . However, the decreasing rates are quite different. Fig. 2(a)–(c) illustrate that the means of ρ_X and ρ_P descend more slowly than those of ρ_S and ρ_K , suggesting the superiority of the former two coefficients when deterministic signals s_p , s_h , and s_t are fed into LM1. Furthermore,

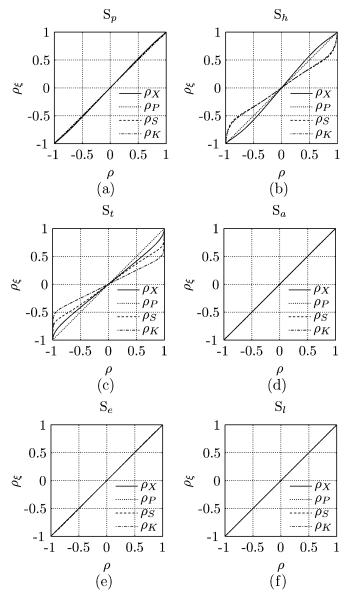


Fig. 3. Relations between $\overline{\rho}_{\xi}$ and ρ under LM2.

 ρ_X outperforms ρ_P in cases with respect to s_p and s_h . On the other hand, the immaterial differences observed in Fig. 2(d)–(f) indicating the equivalence of the four methods when the inputs are stochastic signals s_a , s_e , and s_l . The overall noise robustness performance thus can be ordered as r_X , $r_P > r_K$, and r_S .

C. Comparative Study Under Linear Model LM2

The relationships between $\overline{\rho}_{\xi}$ and ρ for the six original signals s_{ζ} are shown in Fig. 3. It is easily observed that 1) $-1 \leq \rho_{\xi} \leq 1$ (Property 1); 2) $\rho_{\xi} = \pm 1 (r_{\xi} = \pm 1)$ as $\rho = \pm 1$, respectively (Property 2); 3) $\overline{\rho}_{\xi} = 0$ ($\overline{r}_{\xi} = 0$) as $\rho = 0$ (Property 4); and 4) $\overline{\rho}_{\xi}$ (\overline{r}_{ξ}) is an increasing function of ρ . Noticing that the closer the distance of $\overline{\rho}_{\xi}$ to the diagonal line, the smaller the associated biasedness, we also observe that when the model inputs are deterministic signals excepting s_{p} , the unbiasedness performance can be ordered as $r_{P} > r_{X} > r_{K} > r_{S}$; whereas for the stochastic signals, there is no substantial difference among the four methods. This phenomenon is quantitatively highlighted in

TABLE I AAT Comparison Results Among Four Methods

	AAT_{X} (%)	AAT_{P} (%)	AAT_{S} (%)	$\overline{AAT_{K}\left(\% ight)}$
$\overline{S_p}$	0.88	0.07	0.55	0.33
$S_{\!h}$	2.34	0.02	7.53	7.12
S_{t}	3.75	0.01	6.95	12.12
$S_{\!a}$	0.09	0.01	0.13	0.13
$S_{\!\!e}$	0.04	0.02	0.23	0.16
S_l	0.02	0.01	0.03	0.03

Table I which summarizes the AAT_{ξ} with respect to the six original signals s_{ζ} . As for the performance of detecting changes in the underlying ρ , we tabulate the sensitivity ratios (SR) in Table II, showing that the capability of discriminating changes in ρ can be ordered as $r_P > r_X > r_K > r_S$, the same as that of unbiasedness performance.

D. Comparative Study Under Linear Model LM3

Under this model, r_{ξ} is computed as a function of time-shift κ , say, which varies from -100 to 100 ms. For each α and each episode n(i) of 1000 white noises, $r_{\varepsilon}(\alpha, \kappa)$ is calculated and the time-shift with respect to the maximum of $r_{\xi}(\alpha, \kappa)$ is the estimate of the time-delay Δ and denoted by κ_{Δ} . Limited by the length of this paper, we only present the results with respect to s_e here. Fig. 4(a) shows four typical waveforms of $r_{\varepsilon}(\alpha,\kappa)$ in the presence of a 50% SNR ($\alpha=1$). All the four coefficients can correctly detect the time-delay between x and y giving $\kappa_{\Delta} = 30$ ms which equals the true time-delay Δ . In Fig. 4(b), we present the statistical results of κ_{Δ} versus the underlying α from 0 to 1 with $\Delta \alpha = 0.1$. The levels of rectangular bars represent the means $\overline{\kappa}_{\Delta}$ and the error bars represent $3 \times v_{\kappa_{\Delta}}$ with $v_{\kappa_{\Delta}}$ denoting the standard deviation of κ_{Δ} . It can be observed from Fig. 4(b) that $\overline{\kappa}_{\Delta}$ slightly increases with increase of noise levels and so does the standard deviation $v_{\kappa \wedge}$ for all four r_{ξ} . The performances of r_X and r_P are better than those of two rank-based methods r_S and r_K in the sense that the former two coefficients have smaller deviations. However, the performance of time-delay detection is not further compared since the maximal error is only 2 ms in all cases for all the four methods. In other words, we do not consider that there are significant differences between the four methods in the aspect of detecting time delays.

E. Comparative Study Under Nonlinear Model NM

The nonlinear model NM is constructed on the linear model LM2 by introducing two increasing nonlinear transformations $T_x[\bullet] = \operatorname{sgn}(\bullet) \cdot (\bullet)^2$ and $T_y[\bullet] = \exp(\bullet)$. Besides the association parameter ρ carrying the same meaning as in LM2, we employ another parameter $\beta = 2, 4, 6, 8, 10$ to control the extent of nonlinearity. It is noteworthy that $T_x[\bullet]$ and $T_y[\bullet]$ have no effect on the rank-based measures r_S and r_K because rankings are invariant under strictly increasing transformations.

Fig. 5 shows the relationships between $\overline{\rho}_{\xi}$ and ρ with respect to the six biosignals s_{ζ} with nonlinearity parameter $\beta=2$. We observe that 1) for periodic signal s_{p} , ρ_{X} performs comparably

,	$ ho_1$	0.0	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8
/	o_2	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8	0.9
	SR_X	2.2295	2.2899	2.4286	2.6697	3.0628	3.7103	4.8474	7.1243	13.1615
S_p	SR_P	2.2301	2.2916	2.4317	2.6754	3.0746	3.7386	4.9257	7.3876	14.3773
\sim_p	SR_S	2.1470	2.2032	2.3390	2.5796	2.9761	3.6249	4.7660	6.9956	11.8843
	SR_K	2.1459	2.2022	2.3369	2.5730	2.9579	3.5827	4.6581	6.6673	11.0515
	SR_X	2.2082	2.2665	2.3955	2.6144	2.9626	3.5175	4.4560	6.2641	10.8761
Q.	SR_P	2.2387	2.3011	2.4424	2.6880	3.0899	3.7581	4.9525	7.4295	14.4629
S_h	SR_S	1.5498	1.5433	1.5415	1.5476	1.5699	1.6228	1.7580	2.1406	3.4626
	SR_K	1.5508	1.5493	1.5556	1.5722	1.6087	1.6789	1.8356	2.2493	3.6324
	SR_X	1.9265	1.9686	2.0616	2.2160	2.4528	2.8147	3.3906	4.4035	6.6993
C	SR_P	2.2821	2.3454	2.4892	2.7393	3.1485	3.8290	5.0456	7.5686	14.7325
S_t	SR_S	1.6928	1.7098	1.7427	1.7932	1.8643	1.9614	2.0913	2.2770	2.7136
	SR_K	1.6920	1.7096	1.7442	1.7977	1.8741	1.9821	2.1356	2.3810	2.9659
	SR_X	2.2326	2.2949	2.4359	2.6805	3.0802	3.7436	4.9265	7.3708	14.2385
C	SR_P	2.2313	2.2934	2.4341	2.6787	3.0790	3.7444	4.9337	7.3994	14.3982
S_a	SR_S	2.1590	2.2187	2.3507	2.5813	2.9461	3.5381	4.5339	6.4251	11.0119
	SR_K	2.1571	2.2177	2.3520	2.5872	2.9648	3.5873	4.6611	6.7816	12.2714
	SR_X	2.2965	2.3595	2.5026	2.7510	3.1563	3.8265	5.0136	5.0136	14.0770
Q	SR_P	2.3022	2.3663	2.5118	2.7645	3.1782	3.8662	5.0966	5.0966	14.9067
S_{e}	SR_S	2.2352	2.2953	2.4333	2.6654	3.0398	3.6439	4.6804	4.6804	11.8162
	SR_K	2.2359	2.2965	2.4371	2.6732	3.0580	3.6863	4.7813	4.7813	12.6448
	SR_X	2.2445	2.3069	2.4486	2.6947	3.0974	3.7668	4.9636	7.4481	14.5038
C	SR_P	2.2457	2.3082	2.4501	2.6967	3.1004	3.7720	4.9735	7.4678	14.5614
S_l	SR_S	2.1617	2.2138	2.3385	2.5521	2.8966	3.4566	4.4232	6.3403	11.3488
	SR_K	2.1615	2.2166	2.3461	2.5694	2.9336	3.5311	4.5755	6.6912	12.4021

TABLE II SRs Comparison of the Resultants of LM2

with ρ_S and ρ_K ; 2) in cases with respect to other two deterministic signals, ρ_X has the smallest biasedness; 3) for the three stochastic signals, ρ_K and ρ_S surpass ρ_X ; and 4) ρ_P has the largest biasedness in all cases. Moreover, r_P never approaches ± 1 as $\rho \to \pm 1$, that is, r_P underestimates the strength of association when nonlinearity is involved. On the other hand, $r_X = \pm 1$ as $\rho \to \pm 1$, which indicates the validity of property 2 under increasing nonlinear transforms. Table III shows the effect of nonlinearity on ρ_X and ρ_P , elucidating that with increase of β , the biasedness (RAT) caused for ρ_X is significantly smaller than that for ρ_P . Sensitivity ratios ($\beta = 2$) tabulated in Table IV show that in most cases, ρ_K performs best; in almost all cases, ρ_P has the lowest SR, indicating the limitation of r_P in nonlinear scenarios.

F. Comparison of Time Complexities

The time complexities of r_ξ are analyzed and summarized in Table V based on the definitions. The fastest method is Pearson's coefficient r_P having a linear time complexity of O(N). Our new method r_X and Spearman's rho are of the same order $O(N\log N)$, since sorting operation dominants the computational time of both methods. However, because of the extra procedure of ranking involved in the calculation of r_S , we can expect that r_X is a little faster than r_S . Kendall's tau r_K is the slowest method compared to the other three coefficients. The core operation of r_K is to calculate the number of concordant and discordant pairs, which requires $C_N^2 = N(N-1)/2$ operations. Therefore, the time complexity of r_K is of $O(N^2)$.

To confirm this result, we estimate the relationship between computational loads of r_{ξ} versus the length of signal N, where N begins at 100 and increases by steps of 100 until N=1000. All the computational speed tests were performed in MATLAB 7.0 in a Pentium PC. For each pair of time series of size N,

the algorithms of r_{ξ} were run for 1000 times. The results are presented in Fig. 6, which is consistent with our analysis.

V. DISCUSSION

A. Rationality of Selection of Comparison Objects

Apart from the three methods used in our comparative study, some authors have used other techniques, such as the AAMI [7], [8], the nonlinear regression coefficient h^2 [9], and the contingency table-based method V [10]. However, we did not include these methods into our comparative studies due to 1) AAMI is unbound [9] and hence is incomparable to our new method and 2) the admissible values of h^2 and V are confined from 0 to 1, that is, they cannot distinguish positive associations from negative associations. Moreover, the analytical relationship between $V(h^2)$ and ρ under bivariate normal model is unknown, which prevents us from doing calibration as in (23) and (27). On the other hand, Pearson's coefficient, Spearman's ${\it rho}$ and Kendall's ${\it tau}$ have similar meaning as our new measure, thus making it possible to compare their behaviors.

B. Estimation of the Hurst Parameter H by r_X

As remarked before, the power spectral density of a long-range-correlated signal is proportional to $f^{-(2H-1)}$, where 0 < H < 1 is the Hurst parameter. Therefore, we can estimate H from the power density which is defined by the Fourier transform of the corresponding autocorrelation function R(k), $k \ge 0$. Noticing that $r_P(k)$ is the normalized version of R(k) and $r_X(k)$ behaves similarly with $r_P(k)$ under linear models, we can estimate the power density S(f) based on the Fourier transform of $r_P(k)$ as well as $r_X(k)$. The Hurst parameter H can then be measured with the slope of $\log(S(f))$ against $\log(f)$. In Fig. 7, we present the results on the full version of the long-range-correlated signal $s_{lf}(i)$. For clarity, the waveforms of r_P

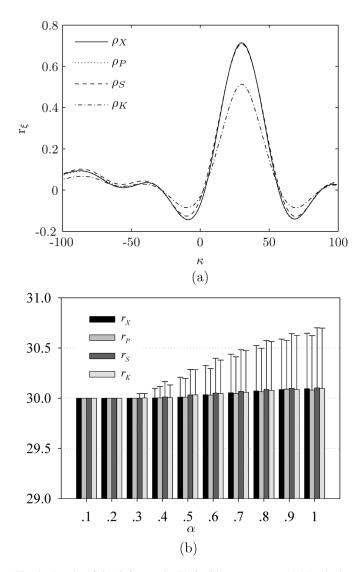


Fig. 4. Results of simulations under LM3 with respect to s_e . (a) Magnitudes of r_ξ associated with time-shift κ varying from -100 to 100 ms in the presence of a 50% SNR ($\alpha=1$). (b) Statistical results of κ_Δ versus the underlying α from 0 to 1 with a step 0.1.

in Fig. 7 are vertically shifted down, since otherwise the waveforms of r_X and r_P will be almost coincident and, therefore, unclear to observe. The corresponding power densities are plotted in a double-log scale in Fig. 7(b), where $\delta=2\hat{H}-1$. It can be easily obtained that $\hat{H}_X=0.8962$ and $\hat{H}_P=0.8960$, very close to the real value H=0.9.

C. Sufficiency of Signal Length Used in this Study

It can be shown that the variances of r_X , r_P , r_S , and r_K are all of the order of $O(N^{-1})$ [2], [5], [30], where N is the length of the signals. In other words, the larger the sample size, the smaller the variances, and the more accurate the four coefficients. For signals consisting of 1000 sample points, the variances of the four coefficients are already very small (of the order of 0.001), which means that a sample size of 1000 is sufficient for studying the behaviors of these coefficients. For the long-range-correlated time series whose length is far greater

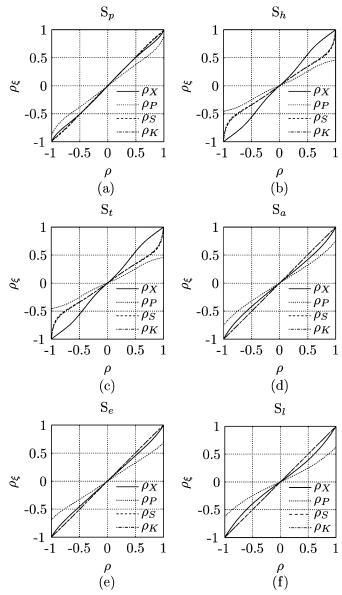


Fig. 5. Relations between $\overline{\rho}_{\xi}$ and ρ under NM. All subplots illustrate the inferiority of r_P in this nonlinear model. The wandering from positive (negative) unity of $\rho_P(r_P)$ when $\rho=\pm 1$ reveals the misleading results of r_P when nonlinearity involved.

than 1000, the variances of the four cross-correlation coefficients are very close to zero, thus assuring the accuracy of the four measurements of association.

D. Solution to Asymmetry of r_X

In general, our new measure r_X is not symmetric, namely, $r_X(x,y) \neq r_X(y,x)$ although $E\{r_X(x,y)\} = E\{r_X(y,x)\}$. This problem can be easily solved using a revised version $r_X'(x,y) = [r_X(x,y) + r_X(y,x)]/2$ when symmetry is a critical feature in practice.

E. Clinical Application of r_X

It is of great clinical importance for speedy and reliable detection of atrial fibrillation (AF) and atrial flutter (AFL) in automatic implantable atrial defibrillators [23]. AF is a type of arrhythmia (abnormal heart rhythm) exhibiting rapid and

TABLE III
RAT COMPARISON UNDER NONLINEAR MODEL NM BETWEEN TV AND TR

	S	S_p	S	S_h	S	S_t	S	S_a	S	S_e	S	$\overline{S_l}$
	$\overline{RAT_X}$	$\overline{RAT_P}$										
	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)
2	0.93	5.68	0.84	12.79	4.37	7.97	3.04	8.26	1.74	8.57	3.57	11.60
4	2.96	8.25	2.61	16.03	5.03	11.03	5.51	12.34	4.15	12.64	5.45	17.10
6	4.50	9.78	3.69	17.63	5.47	12.39	6.87	14.51	5.49	14.68	5.98	18.97
8	5.53	10.83	4.32	18.48	5.79	13.27	7.77	15.93	6.37	15.94	6.19	19.87
10	6.29	11.60	4.72	19.00	6.04	13.92	8.41	16.93	7.02	16.83	6.28	20.38

TABLE IV SRS COMPARISON OF THE RESULTANTS OF NM. THE MAXIMA AND MINIMA ARE HIGHLIGHTED WITH GRAY AREAS AND WHITE BOXES, RESPECTIVELY

	$ ho_1$	0.0	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8
	ρ_2	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8	0.9
	SR_X	1.9808	2.0212	2.1149	2.2729	2.5144	2.8715	3.4170	4.4060	7.1379
S_p	SR_P	1.7324	1.7513	1.7989	1.8794	2.0017	2.1885	2.5027	3.1359	4.8094
\mathcal{O}_p	SR_S	2.1470	2.2032	2.3390	2.5796	2.9761	3.6249	4.7660	6.9956	11.8843
	SR_K	2.1459	2.2022	2.3369	2.5730	2.9579	3.5827	4.6581	6.6673	11.0515
	SR_X	1.5219	1.5283	1.5456	1.5755	1.5975	1.5978	1.6121	1.7553	2.3737
a	SR_P	0.9321	0.9152	0.9054	0.9011	0.8900	0.8546	0.7897	0.7104	0.6535
S_h	SR_S	1.5498	1.5433	1.5415	1.5476	1.5699	1.6228	1.7580	2.1406	3.4626
	SR_K	1.5508	1.5493	1.5556	1.5722	1.6087	1.6789	1.8356	2.2493	3.6324
	SR_X	0.7710	0.7942	0.8415	0.9037	0.9843	1.1082	1.3222	1.7187	2.6497
a	SR_P	0.9064	0.8863	0.8891	0.9200	0.9696	1.0323	1.1083	1.2123	1.4081
S_t	SR_S	1.6928	1.7098	1.7427	1.7932	1.8643	1.9614	2.0913	2.2770	2.7136
	SR_K	1.6920	1.7096	1.7442	1.7977	1.8741	1.9821	2.1356	2.3810	2.9659
	SR_X	1.7298	1.7375	1.7771	1.8586	2.0021	2.2485	2.6880	3.5533	5.6944
~	SR_P	1.3216	1.3213	1.3296	1.3529	1.4024	1.4977	1.6745	2.0101	2.7336
S_a	SR_S	2.1590	2.2187	2.3507	2.5813	2.9461	3.5381	4.5339	6.4251	11.0119
	SR_K	2.1571	2.2177	2.3520	2.5872	2.9648	3.5873	4.6611	6.7816	12.2714
	SR_X	1.9998	2.0313	2.1101	2.2460	2.4591	2.7911	3.3468	4.4269	7.2043
a	SR_P	1.5529	1.5642	1.5881	1.6248	1.6769	1.7529	1.8789	2.1231	2.6991
S_{e}	SR_S	2.2352	2.2953	2.4333	2.6654	3.0398	3.6439	4.6804	6.6980	11.8162
	SR_K	2.2359	2.2965	2.4371	2.6732	3.0580	3.6863	4.7813	6.9522	12.6448
	SR_X	1.4193	1.3968	1.3618	1.3481	1.3825	1.4963	1.7389	2.2406	3.4734
G	SR_P	1.0233	0.9999	0.9652	0.9401	0.9388	0.9726	1.0573	1.2269	1.5849
S_l	SR_S	2.1617	2.2138	2.3385	2.5521	2.8966	3.4566	4.4232	6.3403	11.3488
	SR_K	2.1615	2.2166	2.3461	2.5694	2.9336	3.5311	4.5755	6.6912	12.4021

TABLE V
TIME COMPLEXITY ANALYSIS FOR FOUR METHODS

Operation	r_X	$r_{\!P}$	r_{S}	r_{K}
±	O(N)	O(N)	O(N)	$O(N^2)$
×,÷	O(N)	O(N)	O(1)	$O(N^2)$
$(\)^{2}$	_	O(N)	O(N)	
$\sqrt{}$	_	O(1)		
Sort	$O(N \log N)$		$O(N \log N)$	$O(N \log N)$
Ranking	_		O(N)	O(N)
TC	$O(N \log N)$	O(N)	$O(N \log N)$	$O(N^2)$

Fig. 6. Results of comparative CPU time test for four coefficients studied. A logarithmic scale is used for better visual effect.

random patterns. AFL is another kind of arrhythmia caused by electrical activity propagating through the atria in a fast and regular manner. Some researchers have employed r_P as

an index to detect these two intra-atrial electrograms [31]. Now, we show that r_X can serve as a useful alternative to

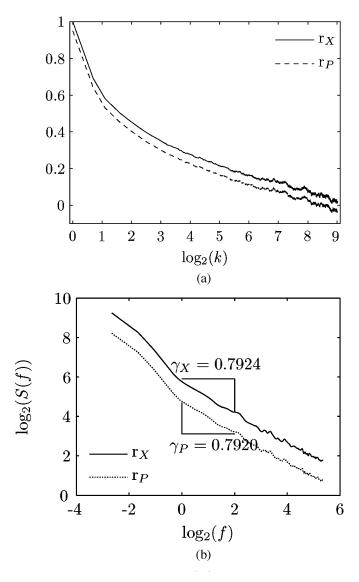


Fig. 7. Estimation of the Hurst parameter H with respect to the long-range-correlated signal $s_{lf}(i)$ by r_X and r_P . The waveforms of r_P are vertically shifted down for a better visual effect.

 r_P in discriminating AF from AFL. Bipolar intra-atrial electrograms (available on physioBank [29]) from one AF patient and one AFL patient are included in this example. The continuous recordings are parsed into nonoverlapping segments of 1-s duration (1000 samples). After a series of preprocessing steps as diagrammed in Fig. 8(a) and detailed in [31], three cross-correlation functions $r_X(k)$, $r_P(k)$, and $r_S(k)$ of time-shift $-100 \le k \le 100$ are calculated with respect to each pair of preprocessed signal segments from two channels. Three maximal values with respect to $r_X(k)$, $r_P(k)$, and $r_S(k)$ are then extracted as discriminatory indices. This operation is repeated sequentially over the entire dataset so that statistical analysis can be performed. It can be observed in Fig. 8(b) that the variance of r_X is the lowest and the average of r_X with respect to AFL is the highest, indicating the superiority of r_X over r_P and r_S . Such advantage of r_X can be quantitatively confirmed by the sensitivity ratios of r_X , r_P , and r_S , being 11, 10, and 6, respectively.

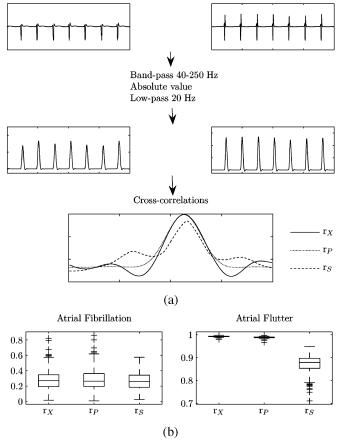


Fig. 8. Correlation analysis technique and comparison results among r_X, r_P , and r_S . (a) Demonstration of the filtering and correlation determination between a pair of simultaneous signal segments. In a parallel manner, the two data segments are first preprocessed by 1) third-order Butterworth band-pass filtering; 2) absolute valuing; and 3) low-pass filtering. Three maximal values are then extracted as discriminatory indices with respect to three cross-correlation functions $r_X(k), r_P(k)$, and $r_S(k)$ of time-shift $-100 \le k \le 100$. (b) Statistical results of $r_X(k), r_P(k)$, and $r_S(k)$. For AF data, the indices congregate near zero; whereas for AFL data, the indices congregate near unity. The larger variance of r_S for AFL data indicates the low discriminatory power of r_S in this case.

F. Summary of Main Advantages of Order Statistics Correlation Coefficient

The numerical results presented allow us to claim the following advantages of r_X as a method of quantification of association between biosignals.

- 1) Noise Robustness: The index r_X decreases slowly as the noise strength increases.
- 2) Small Biasedness: Although r_X is not an unbiased estimator of linear association, the biasedness is very small compared to rank correlations r_S and r_K . In this aspect, r_P is optimal, whereas r_S and r_K have limited power in measuring linear associations of spiky biosignals (see Fig. 3).
- 3) High Sensitivity to Changes in Association: Under linear models, r_X has sensitivity to changes in ρ similar to that of r_P and much higher than those of r_S and r_K .
- 4) High Accuracy for Time Delay Detection: The index r_X has almost perfect performance to detect time delays between two biosignals.

- 5) Fast Computational Speed: The computational load of r_X is relatively light, being much faster than r_S and r_K and a little slower than r_P .
- 6) Good Performance for Nonlinear Association Estimation: Unlike r_S and r_K , r_X is not invariant under increasing nonlinear transforms, but it always performs substantially better than r_P and for spiky signals even better than r_S and r_K .

VI. CONCLUSION

In this paper, we propose a new order statistics correlation coefficient and investigate its properties and applicability to biosignals. The proposed measure was evaluated using simulated and real biosignals and four models emulating linear and nonlinear situations. We also compared the behavior of our measure with three other correlation coefficients commonly used in the literature. The comparative studies demonstrate that our new measure r_X plays the role of a "missing link" between Pearson's coefficient and Spearman's rho and Kendall's tau. It enjoys the advantages of all the other three coefficients. In most cases, r_X is not optimal, but it usually is the second best compared to r_P , r_S , and r_K . This suboptimal feature at least avoids the worst results in practice when one has no prior knowledge as to whether nonlinearity exists in the system. The new method can be applied to a wide spectrum of biosignal processing, such as organizational indexing of atrial fibrillation [22], [23], atrial fibrillation detection [19], EEG association analysis [7]–[9], etc. In fact, the proposed measure can be used in all the fields where the other three classical methods are applicable, although our comparative studies are conducted in the context of biosignal processing.

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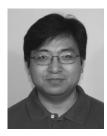
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