

BRS Analysis from Baroreflex Sequences and Baroreflex Events Compared Using Spontaneous and Drug Induced Data

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Abstract

Spontaneous time domain BRS estimation is based on the SBP-RR slope, which can be computed from either baroreflex sequences (BS) or baroreflex events (BE). BRS analysis from BEs was recently shown to be advantageous particularly in the cases of reduced BRS or when BS are not identified. Also, it offers a superior discrimination between lying and standing positions.

In this work, the methods developed for spontaneous BRS analysis are further compared using spontaneous and drug induced data. The results corroborate that spontaneous and drug induced estimates are different although correlated. In particular, if BEs are used the differences and the correlation between the estimates is higher. No precision improvement is achieved if the BRS is estimated from drug induced data. In spontaneous, the higher number of beats in BEs in comparison with BSs (at the expense of a lower SBP-RR correlation) allows a higher BRS estimate precision using recordings of the same length.

1. Introduction

Over the past years, the arterial-cardiac baroreflex sensitivity (BRS) quantification has been useful in the study of many pathological states, including myocardial infarction, hypertension and congestive heart failure [1]. Briefly, lower levels of BRS have been associated with an increased cardiovascular disease-related mortality.

Time domain BRS is traditionally quantified as the regression slope obtained from the cross-analysis between systolic blood pressure (SBP) and RR interval values, either using drug induced or spontaneous data. The advantages and limitations of these techniques have been reviewed elsewhere [2]. Briefly, the noninvasive nature of the spontaneous methods simplifies the test procedure and allows the BRS measurement under a broad range of daily

life conditions, making these methods more appropriate in many research settings. The principle of the invasive technique is to make use of a drug which changes arterial pressure by producing vasoconstriction (or vasodilatation) while not having direct effect on the RR interval, so that the observed RR changes are mediated reflexively via the baroreceptors. In comparison with the spontaneous, drug induced techniques stimulate a larger and clearer SBP increase (or decrease) in order to force a pronounced RR response (i.e, a clearer baroreflex activation). Several comparisons between spontaneous and drug induced BRS estimates evidence that they are different although being correlated (e.g, [3]). Therefore spontaneous BRS estimates can have the same clinical and predictive power as the invasive, even beyond that provided by heart rate variability indices alone [4].

The most frequently used spontaneous BRS method is the sequences technique, which is based on the identification of baroreflex sequences (BS) followed by the BRS estimation as the mean of the SBP-RR slopes computed in each BS [5]. The events technique, recently proposed to improve spontaneous BRS assessment, consists of a global slope estimation from the SBP-RR values in baroreflex events (BE) [6]. The BRS analysis from BEs has shown to be advantageous as it provides a larger number of beats for slope estimation, allowing BRS analysis in cases where BSs cannot be identified. However, the higher number of beats is obtained at the expense of a lower SBP-RR correlation (although being close to 0.8) with possible repercussion on the precision of the BRS estimate.

In this study, a precision measure over time domain BRS estimates from global slope estimation is proposed. Also, the BRS analysis from BSs and from BEs, previously proposed for spontaneous BRS assessment, are further compared using spontaneous and drug induced data. The methods performance is evaluated in terms of the ability to discriminate different conditions and in terms of precision.

2. Time domain BRS estimation

The methods for global BRS estimation combined either with BSs or BEs have been previously detailed [6] and are exemplified in Fig. 1. As illustrated in 1(a) and (b), both BSs and BEs are identified considering $x_{\text{SBP}}(n)$ paired with $x_{\text{RR}}(n+1)$, with n being the beat number. Briefly, each valid BS_{*k*} requires a minimum beat length ($N_k \geq 3$), minimum SBP and RR beat-to-beat changes in the same direction ($\Delta_k^{\text{SBP}} \geq 1$ mmHg and $\Delta_k^{\text{RR}} \geq 5$ ms) and a minimum x_{SBP} and x_{RR} correlation ($r_k \geq 0.8$), whereas for the identification of each BE_{*k*} only the thresholds N_{min} and r_{min} are enforced. After the identification, the mean is removed from x_{SBP} and x_{RR} at each segment and the result concatenated in \mathbf{d}_{SBP} and \mathbf{d}_{RR} vectors, respectively. Finally, the global slope $\mathcal{B}_{\text{G.O}}$ is obtained from the regression analysis $\mathbf{d}_{\text{RR}} = \mathcal{B}_{\text{G.O}} \mathbf{d}_{\text{SBP}} + \epsilon$, where the parameter $\mathcal{B}_{\text{G.O}}$ is estimated by ordinary least squares (OLS) minimization and ϵ a noise vector. Additional variables can be retrieved from global BRS analysis, namely the number of points for slope estimation (N) and the \mathbf{d}_{SBP} - \mathbf{d}_{RR} correlation (r). Figure 1(c-d) presents the \mathbf{d}_{SBP} and \mathbf{d}_{RR} dispersion diagrams for the example in (a-b), illustrating that BRS analysis from BEs provides a higher N at the expense of a lower r , as a consequence of the less restrictive identification thresholds.

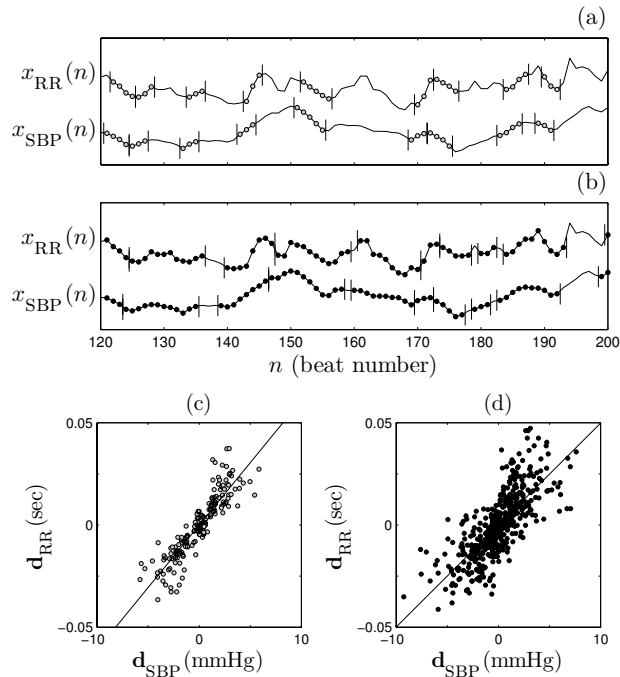


Figure 1. Example of BRS estimation from BSs (a,c) and from BEs (b,d) in a 512 beats spontaneous recording: (a,b) x_{SBP} and x_{RR} plot for the identified baroreflex segments and (c,d) corresponding \mathbf{d}_{SBP} and \mathbf{d}_{RR} dispersion diagrams with the global regression line with slope $\hat{\mathcal{B}}_{\text{G.O}}$. For BSs, $N=171$ and $r=0.89$ and for BEs $N=448$ and $r=0.76$.

2.1. BRS confidence intervals

Standard parametric confidence intervals (CI) over the regression slope can provide a measure of precision, if the regression residuals ($\hat{\epsilon}$) satisfy some requirements (namely, non correlation, zero mean, homoscedasticity and normal distribution). In this study, the precision was computed from a nonparametric bootstrap CI [7], as the $\hat{\epsilon}$ analysis did not support these assumptions, except for the zero mean normal distribution (with 95% confidence).

The linear regression was computed from 1000 Bootstrap (B) replications of \mathbf{d}_{RR} following $\mathbf{d}_{\text{RR}}^{\text{B}} = \hat{\mathcal{B}}_{\text{G.O}} \mathbf{d}_{\text{SBP}} + \hat{\epsilon}^{\text{B}}$, with $\hat{\mathcal{B}}_{\text{G.O}}$ the OLS estimate of $\mathcal{B}_{\text{G.O}}$ and $\hat{\epsilon}^{\text{B}}$ a realization of ϵ reproducing the mean, variance, autocorrelation and heteroscedastic pattern in $\hat{\epsilon}$. The limits of the 95% CI were defined by the 2.5th and 97.5th percentiles of the empirical distribution of the bootstrap slope estimates $\hat{\mathcal{B}}_{\text{G.O}}^{\text{B}}$. Finally, the $\hat{\mathcal{B}}_{\text{G.O}}$ relative maximum error (δ), a measure inversely related to precision, was calculated as the CI half amplitude divided by $\hat{\mathcal{B}}_{\text{G.O}}$.

For the $\hat{\epsilon}^{\text{B}}$ simulation, it was considered $\hat{\epsilon} = \hat{\epsilon}_c + \hat{\epsilon}_h$, with $\hat{\epsilon}_c$ being the constant variance component and $\hat{\epsilon}_h$ the remaining part (i.e. the heteroscedastic pattern, maintained constant in all realizations). The $\hat{\epsilon}_c$ component was estimated from weighted LS regression [7], considering the weights $w_i = (|p_i - 1| / s_p + 1)^{-1}$, $i=1, 2, \dots, N$ with $p_i = \hat{\mathcal{B}}_i / \hat{\mathcal{B}}$, $\hat{\mathcal{B}}_i$ the OLS regression slope excluding the i^{th} ($\mathbf{d}_{\text{SBP}}, \mathbf{d}_{\text{RR}}$) pair and s_p the standard deviation of $\mathbf{p} = [p_1 \ p_2 \ \dots \ p_N]$. The bootstrapped residuals $\hat{\epsilon}_c^{\text{B}}$ were generated as white gaussian noise with the same mean and variance as $\hat{\epsilon}_c$. Its autocorrelation was introduced by means of filtering, after $\hat{\epsilon}_c$ autoregressive modeling (minimum AIC order, from Yule-Walker equations and Levinson-Durbin algorithm [8]). The remaining part was obtained by $\hat{\epsilon}_h = \hat{\epsilon} - \hat{\epsilon}_c$ and set as zero for $|\hat{\epsilon}|$ lower than 1.96 times the $\hat{\epsilon}_c$ standard deviation (i.e. non significant heteroscedasticity). Finally, each noise realization was obtained from $\hat{\epsilon}^{\text{B}} = \hat{\epsilon}_c^{\text{B}} + \hat{\epsilon}_h$. Figure 2 illustrates the dispersion diagram of two bootstrap replicas, showing the similarity with the dispersion diagrams in Fig. 1(c-d).

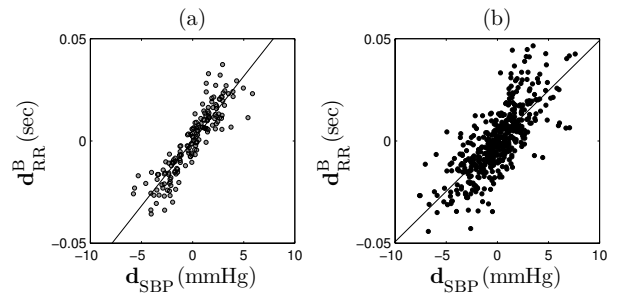


Figure 2. Dispersion diagrams of two bootstrap replicas (a-b) reproducing the real data in Figure 1(c-d), with the global regression line with slope $\hat{\mathcal{B}}_{\text{G.O}}^{\text{B}}$.

3. Experimental data

The data was collected from 15 healthy male subjects (20-36 years) in supine rest condition [9]. Each subject was monitored 5 min in spontaneous (SP) condition and during the 3 min modified Oxford protocol. The latter consists of a bolus injection of 150 μg sodium nitroprusside (NT) followed 1 min later by a bolus of 150 μg phenylephrine HCl (PH). Figure 3 presents the data from one subject, illustrating that the consecutive NT (vasodilator) and PH (vasoconstrictor) bolus acutely decrease/increase SBP and produce a baroreflex mediated shortening/lengthening of the RR interval, respectively. Also, it can be observed that in this protocol there is a short time gap between the successive injections and the PH is administrated still under the NT effect, leading to a mixture of effects. Therefore, this experimental setting increases the range of SBP changes observed in SP acquisition, but only by lowering the x_{SBP} and x_{RR} values with respect to the baseline.

In this work, the BRS in SP was assessed from the 5 min recordings, whereas in NT and PH it was computed from consecutive 45 sec segments starting 30 sec after the NT bolus (Fig. 2). The 30 sec lag assures that the NT effect is present, as its onset of action is within 30 sec [10], whereas 45 sec is the maximum time interval for all subjects between the beginning of NT and PH effects.

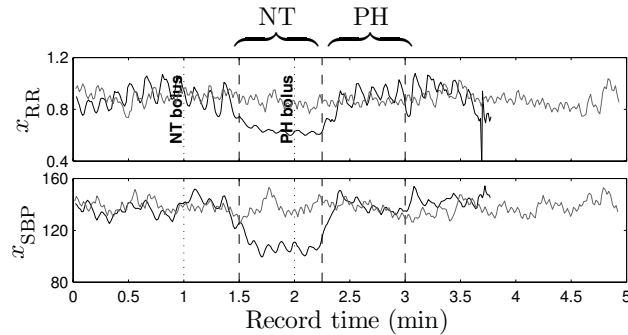


Figure 3. Plot of x_{SBP} and x_{RR} in SP (grey) and during the modified Oxford method (black). Dotted lines identify the timing of NT and PH bolus and the brackets delimitate the time intervals for BRS analysis in NT and PH conditions.

4. Results

The methods were compared with respect to $\hat{\mathcal{B}}$ and δ was studied as a function of N and r . Indices (S,E) and (SP,NT,PH) were added to each variable according to its evaluation: e.g., $\hat{\mathcal{B}}_{\text{SP}}^{\text{E}}$ refers to $\hat{\mathcal{B}}$ from BEs in SP.

It is expected that RR changes after induced SBP increase or decrease are asymmetrical, being the RR response to falling SBP lower than of that to rising SBP [11]. Figure 4 shows the comparison between the $\hat{\mathcal{B}}$. For BEs, $\hat{\mathcal{B}}_{\text{SP}}^{\text{E}} > \hat{\mathcal{B}}_{\text{NT}}^{\text{E}}$ for 15/15 subjects, $\hat{\mathcal{B}}_{\text{SP}}^{\text{E}} > \hat{\mathcal{B}}_{\text{PH}}^{\text{E}}$ for 9/15 and $\hat{\mathcal{B}}_{\text{PH}}^{\text{E}} > \hat{\mathcal{B}}_{\text{NT}}^{\text{E}}$

for 14/15. With BSs, the countings are 11/15, 4/15, and 13/15, respectively. The mean paired differences $\hat{\mathcal{B}}_{\text{SP}}^{\text{E}} - \hat{\mathcal{B}}_{\text{NT}}^{\text{E}}$ and $\hat{\mathcal{B}}_{\text{NT}}^{\text{E}} - \hat{\mathcal{B}}_{\text{PH}}^{\text{E}}$ differ from zero ($p < 0.01$), whereas $\hat{\mathcal{B}}_{\text{SP}}^{\text{E}} - \hat{\mathcal{B}}_{\text{PH}}^{\text{E}}$ do not ($p > 0.1$). The pairwise correlation between $\hat{\mathcal{B}}_{\text{SP}}^{\text{E}}$, $\hat{\mathcal{B}}_{\text{NT}}^{\text{E}}$ and $\hat{\mathcal{B}}_{\text{PH}}^{\text{E}}$ is higher for BEs ($r > 0.85$ and $p < 0.01$ for the null hypothesis of no correlation).

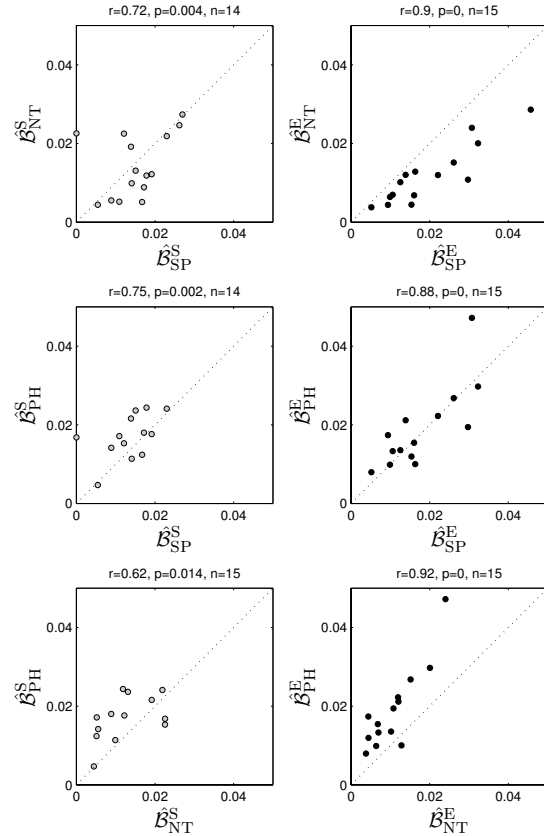


Figure 4. Dispersion diagrams comparing $\hat{\mathcal{B}}^{\text{S}}$ (grey) and $\hat{\mathcal{B}}^{\text{E}}$ (black) in SP, NT and PH conditions.

The value of δ is a trade-off between N and the $\hat{\epsilon}$ standard deviation (inversely related with r), with higher N (or r) leading to lower δ (i.e., higher precision). As illustrated in Fig. 5, the less restrictive thresholds for BE identification lead to $r_{\text{SP}}^{\text{E}} < r_{\text{SP}}^{\text{S}}$ and $N_{\text{SP}}^{\text{E}} > N_{\text{SP}}^{\text{S}}$ so that $\delta_{\text{SP}}^{\text{E}} < \delta_{\text{SP}}^{\text{S}}$ in 8/12 cases and the mean paired differences $\delta_{\text{SP}}^{\text{E}} - \delta_{\text{SP}}^{\text{S}}$ do not differ from zero ($p > 0.1$). The δ_{SP} value decays with increasing N_{SP} and r_{SP} remaining close to 0.8. Therefore, $\delta_{\text{SP}}^{\text{E}}$ is more influenced by N_{SP} than by r_{SP} . For NT and PH, N is lower and r is higher than in SP. For BSs, $\delta_{\text{NT}}^{\text{S}}$ and $\delta_{\text{PH}}^{\text{S}}$ have the same range as $\delta_{\text{SP}}^{\text{S}}$ and $\delta_{\text{SP}}^{\text{S}} < \delta_{\text{NT}}^{\text{S}}$ in 5/10 of the cases and $\delta_{\text{SP}}^{\text{S}} < \delta_{\text{PH}}^{\text{S}}$ in 5/10. For BEs, the slightly higher r^{E} and much lower N^{E} in NT and PH in comparison to SP lead to $\delta_{\text{SP}}^{\text{E}} < \delta_{\text{NT}}^{\text{E}}$ in 13/15 cases and $\delta_{\text{SP}}^{\text{E}} < \delta_{\text{PH}}^{\text{E}}$ in 12/15. That is, no precision improvement is achieved with the drug-induced protocol: for BSs, the lower N^{S} is balanced by the increased r^{S} , so that the δ^{S} is maintained; for BEs, the

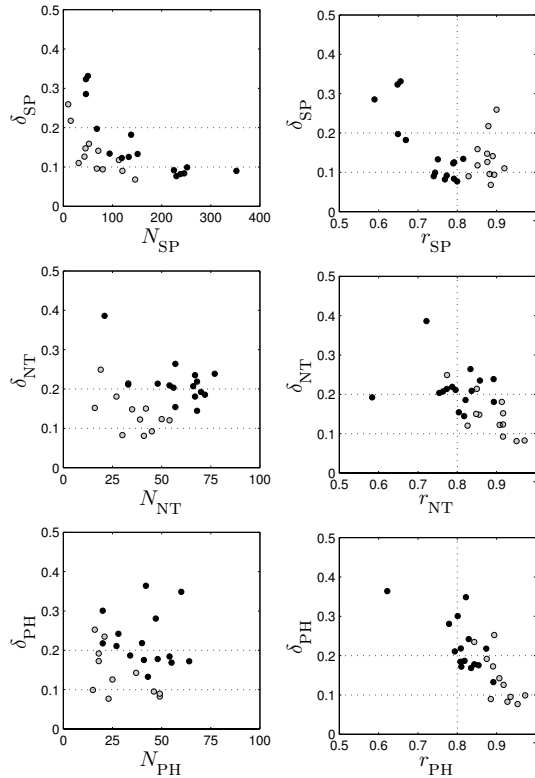


Figure 5. Plot of δ as a function of N and r for \hat{B}^S (grey) and \hat{B}^E (black) in SP, NT and PH conditions.

higher r^E does not pay off the much lower N^E , so that δ^E is higher in either NT and PH when compared to SP.

In SP there is no time span constrain as in drug-induced experiments (especially if using a bolus instead of a continuous infusion). Therefore, the possibility of increase N in SP is an advantage over increase r in a drug-induced protocol: in SP, higher N requires longer recordings whereas higher r requires higher $x_{SBP}-x_{RR}$ correlation (only obtained with stimulation). In SP, r_{SP}^S is higher than 0.8 and r_{SP}^E is constrained to be lower than 0.8 (with $r_{min}=0.8$ for BE identification, r^E will tend to be 0.8 in SP stationary recordings [6]). In contrast, the longer the recordings the higher the N^E and N^S difference is. Therefore, if the SP stationary conditions are satisfied (i.e. r_{SP}^E close to 0.8), it is expected that \hat{B}_{SP}^E outperforms \hat{B}_{SP}^S in precision.

5. Conclusions

In this work, BRS analysis from BSs and BEs is further compared using spontaneous and drug induced data. Besides its non invasive nature, the spontaneous BRS estimates are preferred to drug induced estimates, as no precision improvements are achieved with the modified Oxford method. The BRS analysis from BEs distinguishes better the different conditions, including for BRS estimates with lower precision. Also, with the use of BEs instead of BSs

in spontaneous condition, an improvement of the BRS estimate precision is achieved, for the same recording length.

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