

Baroreflex Sensitivity Evolution Before Weaning From Mechanical Ventilation

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Abstract

Weaning is the process of withdrawing mechanical ventilation at the Intensive Care Units. The problem is that around 20% of weaned patients were not actually ready for discontinuation. Studies suggest that vagal dysfunction is lower in patients successfully weaned. Therefore, the Baroreflex Sensitivity (BRS) and Heart Rate Variability (HRV) are estimated to see if they can provide additional information to improve the prediction of weaning outcomes.

9 successfully weaned patients (S-group) and 6 unsuccessfully weaned (F-group) were monitored in the last hour prior to the Spontaneous Breathing Trial. The BRS is estimated through spectral analysis, to obtain the α parameter in the low and high frequency bands, and through the capacity, C , estimated by the Bivariate Phase Rectified Signal Average (BPRSA) method. The current clinic parameters of weaning readiness do not show statistical differences. However, the capacity to changes of the BRS, C , estimated via BPRSA, exhibits significant differences between the two groups. Negative values of C , and with higher absolute values, were obtained for the S-group. Temporal indices of HRV also show differences, but not significant. These results suggest that BRS should be further explored for predicting weaning outcomes.

1. Introduction

Patients admitted to Intensive Care Units (ICU) generally need the support of Mechanical Ventilation (MV) to maintain breathing. Weaning is the process of withdrawing this MV. It is a challenging and very delicate procedure and it must be considered as soon as possible [1]. However, weaning failure and reintubation after a premature weaning is really critical and can lead to an increased risk of severe respiratory complications such as pneumonias, lung injuries, or diaphragmatic dysfunctions. Indeed,

weaning failure increases the risk of mortality in the range from 25% to 50%. The Spontaneous Breathing Trial (SBT) is the diagnostic test to determine weaning readiness [1], and some parameters based on respiration, cardiovascular status and hemodynamics help clinicians to determine if a weaning attempt will be successful or not.

The cardiovascular control involves several mechanisms to dynamically couple arterial pressure, heart rate and respiration. The cross-talk between heart, lungs and nervous system is mainly coordinated by the Autonomic Nervous System (ANS). Indices derived from Baroreflex Sensitivity (BRS) and Heart Rate Variability (HRV) have already been proven to assess the regulatory mechanisms of the ANS [2]. In fact, recent studies revealed high incidence of autonomic dysfunction in patients admitted to the ICU, resulting in chronic adrenergic activations and acute respiratory distress syndromes [3]. However, no previous work investigated the role of the BRS. Moreover, none looked for differences just before the SBT, in order to improve the predictive value of the weaning indices.

This study aims to evaluate the BRS in the last hour prior to the SBT and explore if this can provide additional information to improve the prediction of weaning outcomes. Different approaches exist to assess BRS. In this work, the BRS will be calculated via the spectral analysis of HRV and SAP, to derive the well-known α index [2]. Furthermore, the method referred to as Bivariate Phase Rectified Signal Averaging (BPRSA) will also be computed [4].

2. Materials and Methods

2.1. Data set

Patients were monitored in the ICU's attached to the Hospital Universitari Parc Taulí of Sabadell (Barcelona), during the hour prior to the SBT. Patients with neurological disorder, dementia or focal brain injury at ICU admission

were excluded for the analysis. For this study, only those patients ventilated with assist/support ventilation modes during the last hour were considered.

When the clinical status and health condition of a patient improved in the ICU, they are considered presumably ready for weaning [1]. These patients had to perform the SBT. The SBT was carried out by a T-tube test or by low-level inspiratory pressure support. The criteria in [1], shows the indicators used to evaluate the success of the SBT, in order to decide if MV could be withdrawn. Weaning was considered successful for patients who passed SBT [1], and did not require reintubation or reconnection to MV in the 48 hours after SBT. These patients are in the successful weaning group, namely *S-group*. Patients presenting at least one item of the intolerance criteria [1], were not ready for discontinuation and weaning failure was considered. These patients were classified in the *F-group*. With all this premises, 9 patients from the *S-group* and 6 from the *F-group* remained.

Physiological signals were recorded using the proprietary system *BetterCare®*. This system for data collection standardizes and synchronizes all the biosignals from all the different bedside monitors and ventilators. The signals are sampled at 200 samples per second. The present study uses the recordings of the lead II of the electrocardiogram (ECG), the blood pressure (BP) signal, acquired using an invasive arterial catheter, and the respiratory frequency signal, $f_r(t)$.

2.2. HRV estimation

The ECG signal is upsampled at 1000 Hz with cubic spline interpolation. Then, the R-waves are detected by means of a wavelet-based method [6]. Detections are manually corrected using the R-DECO application [7]. Once ECG detections are correct, the RR interval is defined by the time between two consecutive R-waves, and the normal-to-normal (NN) interval is defined after correcting ectopic beats [8]. From the NN series, temporal indices of HRV are calculated: standard deviation of the NN intervals (SDNN) and the root mean-square of successive differences (RMSSD).

In order to explain the regulation of the heart rate by the ANS, the HRV signal is estimated using the Time-Varying Integral Pulse Frequency Modulation model [8]. Given a particular beat time occurrence series, the instantaneous HR, $d_{HR}(t)$, can be derived and expressed as $d_{HR}(t) = (1 + m(t))/T(t)$. The term $m(t)$ represents the modulating signal, which is assumed to contain the ANS modulation. The instantaneous mean HR, $d_{HRM}(t) = 1/T(t)$, is obtained by low-pass filtering $d_{HR}(t)$ at 0.03 Hz. The HRV signal is represented by the term $d_{HRV}(t) = d_{HR}(t) - d_{HRM}(t)$. Finally, the modulating signal is estimated as $m(t) = d_{HRV}(t)/d_{HRM}(t)$. The evenly-sampled version,

$m(n)$, is obtained by resampling $m(t)$ at $f_s = 4$ Hz.

Since respiration affects HRV and SAP through changes in the respiratory frequency and the respiratory pattern, the analysis is performed guided by respiration. Therefore, the High Frequency (HF) band is redefined to be centred at the respiratory frequency, f_r , and time-varying: $\Omega_{HF}(t) = [f_r(t) - 0.15, f_r(t) + 0.15]$ Hz. This is mainly motivated since f_r in MV patients, is around 24 respirations per minute (rpm), and this is in the limit of the classic HF band. The HF power, P_{HF} , is defined as the power within the HF band. Low Frequency (LF) power, P_{LF} , is defined as the power in the classic LF band: $\Omega_{LF} = [0.04, 0.15]$ Hz. The sympathovagal balance is estimated with the normalized power: $P_{LF}^n = P_{LF}/(P_{LF} + P_{HF})$.

2.3. SAP estimation and BRS analysis

An ad hoc algorithm is implemented to delineate the BP signal. The SAP values are essentially detected using a pulse detection algorithm based on a low pass differentiator (LPD) filter [9], and a time-varying threshold defined by an enveloping procedure [7].

First, the BP signal, $x_{BP}(n)$, is filtered using the LPD filter, to accentuate the maximum up-slope of the pulses. This signal is referred to as $x'_{BP}(n)$. After that, the enveloping procedure computes the upper and lower envelope of the filtered BP signal, $x'_{BP}(n)$. A flattened-positive version, $x_F(n)$, is obtained, which enhances the up-slopes. The location of the pulses, n_{D_i} , are found by detecting the peaks in this flattened version, applying the adaptive thresholding procedure of the Pan-Tompkins algorithm [7]. Then, once the locations of the pulses, n_{D_i} , are identified, the BP signal is used to obtain, at the maximum of each pulse, n_{SAP_i} , the corresponding SAP value, $x_{BP}(n_{SAP_i})$. Erroneous SAP detections are manually corrected using the R-DECO as well [7]. An illustration of the procedure is in Fig. 1.

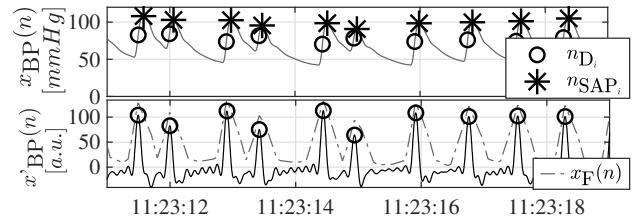


Figure 1. Blood Pressure Delineation. The LPD-filtered BP signal, $x'_{BP}(n)$, is in bottom. The flattened-positive envelope, $x_F(n)$, of $x'_{BP}(n)$ is represented with the dashed-dotted line. From $x_F(n)$, the location of the pulses, n_{D_i} , are identified. The BP signal, $x_{BP}(n)$, plotted in the top, is used to obtain at the maximum of each pulse, n_{SAP_i} , the corresponding SAP value, $x_{BP}(n_{SAP_i})$, for each heartbeat.

Now, the SAP signal, $d_{SAP}^u(n)$, can be estimated as:

$$d_{SAP}^u(n) = \sum_i [x_{BP}(n_{SAP_i})] \delta(n - n_{SAP_i}), \quad (1)$$

where $\delta(\cdot)$ denotes the Kronecker delta function, and the superscript “u” denotes that the signal is unevenly sampled, since heartbeats occur unevenly in time. The evenly-sampled version, $d_{\text{SAP}}(n)$, is obtained by cubic spline interpolation at $f_s = 4$ Hz.

At this point, the BRS indices based on the α index can be calculated from the spectral analysis of HRV and SAP:

$$\alpha_B(n) = \sqrt{\frac{\int_{\Omega_B} \hat{S}_{\text{HRV}}(n, f) df}{\int_{\Omega_B} \hat{S}_{\text{SAP}}(n, f) df}}, \quad B \in \{\text{LF}, \text{HF}\}, \quad (2)$$

where $\hat{S}_{\text{HRV}}(n, f)$ and $\hat{S}_{\text{SAP}}(n, f)$ are the time-varying power spectral densities of $m(n)$ and $d_{\text{SAP}}(n)$, respectively, calculated by means of the Cohen’s Class Wigner Ville Distribution [10], with the parameters adjusted for a time and frequency resolution of 11.25 s and 0.039 Hz, respectively. This time-frequency distribution is used to deal with non-stationarities, since recordings are 1-hour length.

2.4. BPRSA analysis

An alternative method to analyse BRS function has been proposed based on BPRSA [4]. Essentially, an averaged HRV profile is obtained of the overall heart response to SAP increases. For this, anchor points (AP) must be identified firstly in the $d_{\text{SAP}}(n)$ series. Usually, AP’s are defined in samples where the average of their T prior samples is greater than the average of their T subsequent samples [4]:

$$\frac{1}{T} \sum_{i=0}^{T-1} d_{\text{SAP}}(n+i) > \frac{1}{T} \sum_{i=1}^T d_{\text{SAP}}(n-i) \quad (3)$$

However, in this work, AP’s are considered only if they are also local maxima in the corresponding series of Eq. (3). With this approach, only one AP is defined for each up-slope. Now, windows of length $2L$ are segmented around each AP over the HRV series. Finally, the BPRSA curve is obtained by averaging all segmented windows in the HRV. In this study, L is set to $10 * f_s$. T sets an upper frequency limit for the periodicities that can be detected by BPRSA [4]. For each patient, T is adjusted with their mean respiratory frequency, $T \approx f_s / (2.5 \cdot f_r)$. An illustrative example of the methodology is shown in Fig. 2.

The BRS, estimated from the BPRSA curve, is quantified with the capacity term, C [11]:

$$C = \frac{1}{2s} \sum_{i=1}^s \text{BPRSA}(L+i) - \frac{1}{2s} \sum_{i=0}^{s-1} \text{BPRSA}(L-i). \quad (4)$$

The index C is based on the Haar wavelet, but evaluated at a scale s and location $(L+1)$, in the center of the curve. Then, C can be either positive or negative. The scale, s , selects the oscillations in the BPRSA curve that most affect C . In this study, s is taken to be equal to T , since using $s = T$ avoids the need of optimization [11].

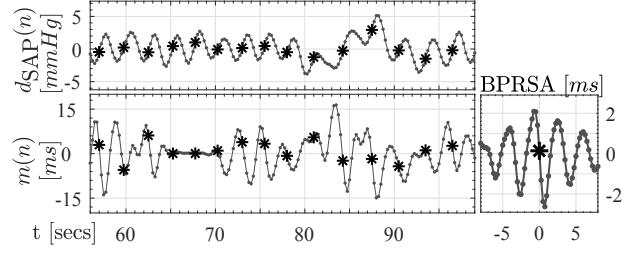


Figure 2. Estimation of the BPRSA curve for a patient. In top-left, the SAP signal, $d_{\text{SAP}}(n)$, as driver mechanism, in which AP’s (asterisks) are defined. In bottom-left, HRV signal, $m(n)$. Around each AP, segments of $2L$ length are extracted from the HRV signal. In bottom-right, the BPRSA curve, obtained averaging all segments of HRV.

2.5. Statistical analysis

The parameters $\overline{\text{HR}}$, $\overline{\text{SAP}}$, $\overline{f_r}$, SDNN, RMSSD, $\overline{\alpha_{\text{LF}}}$, $\overline{\alpha_{\text{HF}}}$, $\overline{P_{\text{LF}}^n}$ and \overline{C} are averaged for the whole recordings of 1-hour (Tab. 1). In addition, α_{LF} , α_{HF} , P_{LF}^n and C , are also averaged in periods of 5 minutes (Fig. 3). The Mann-Whitney U-test is used to compare the values of the *S-group* vs. *F-group*. Statistical significance is considered for $p \leq 0.05$.

3. Results and Discussion

The results for the scalar variables are in Tab. 1. The capacity, \overline{C} , obtained from the BPRSA curve, is the only parameter significantly different comparing *S-group* vs *F-group*. No significant differences can be found for SDNN and RMSSD of the HRV, but they are notably higher for the *S-group*. In fact, other works did find significant differences for these between the two groups when comparing values half an hour before the SBT, and half an hour after SBT [3].

Table 1. Scalar indices. Inter-subject median (Q_1 , Q_3), computed throughout the whole hour before SBT.

	<i>S-group</i>	<i>F-group</i>	p
$\overline{\text{HR}}$ [bpm]	80 (75,89)	84 (81,123)	0.33
$\overline{\text{SAP}}$ [mmHg]	132 (110,139)	126 (101,141)	0.78
$\overline{f_r}$ [rpm]	19 (18,23)	22 (18,24)	0.61
SDNN [ms]	32 (27,53)	22 (15,26)	0.11
RMSSD [ms]	11 (6,14)	5 (3,8)	0.18
$\overline{\alpha_{\text{LF}}}$ [ms/mmHg]	6.0 (3.5,11.8)	5.1 (2.3,7.1)	0.46
$\overline{\alpha_{\text{HF}}}$ [ms/mmHg]	7.1 (3.1,8.7)	6.7 (1.6,8.6)	0.78
$\overline{P_{\text{LF}}^n}$	0.56 (0.30,0.65)	0.66 (0.41,0.80)	0.39
\overline{C} [ms]	-1.7 (-2.7,-0.8)	0.9 (0.5,1.4)	0.02

Since no strong conclusions can be extracted from the averages of the whole hour (see Tab. 1), the evolution of α_{LF} , α_{HF} , P_{LF}^n and C is calculated in 5-minutes averaged-periods, throughout the hour before SBT (Fig. 3). Clear statistical significant differences in C are visible comparing the *S-group* vs *F-group* in many 5-minutes periods.

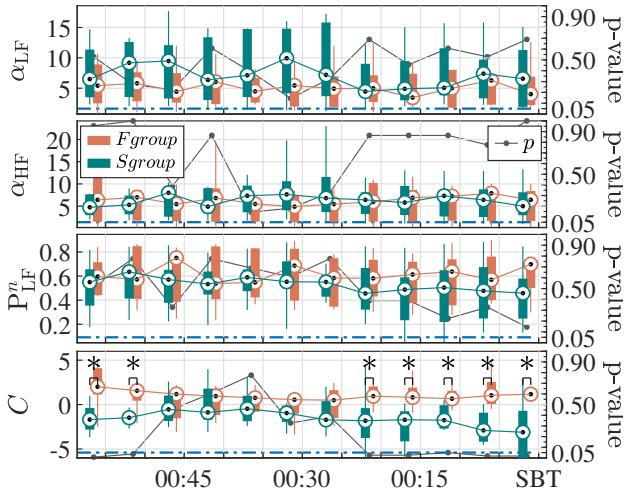


Figure 3. Evolution, throughout the hour before SBT, of the BRS with α_{LF} and α_{HF} , in $[ms/mmHg]$, P^n_{LF} and C in $[ms]$. Green and Red boxplots represent the average of each 5-minutes period for the *S-group* and *F-group*, respectively. The *p-value* is represented in the right axis, from 0 to 1, comparing each 5-minutes period and asterisk represents statistical significance.

Again, no significant differences are found in any period for α_{LF} , α_{HF} or P^n_{LF} either. In spite of that, the α_{LF} median values are generally higher for the *S-group* during the whole hour.

Maybe, no strong differences can be found since these patients present strong non-linear dynamics because of the disease severity [5]. This fact reflects complex influences of respiration on HR and SAP variability and might limit the ability of α_{LF} and α_{HF} for BRS assessment, even though analysis was guided by respiration.

It is important to remark that C in absolute values, is higher for the *S-group*. This could be because of the effect of the strength of SAP changes in HRV, reflected in the amplitude of the BPRSA curve. Besides, notice that C is negative for the *S-group* but positive for the *F-group*, since sympathetic activity is associated with higher latency shifts in the response of the HRV for the spontaneous increases in BP. In fact, related to this, higher adaptability to changes can be deduced from SDNN and higher parasympathetic activity from RMSSD, for the *S-group*. Note that the scale $s = T$, used for the computation of C , is not distorting the results, since \bar{f}_r is similar for both groups (see Tab. 1).

The selection of the AP's for the BPRSA analysis is of relative importance. Using all the increasing samples as AP's [4,5], produces a reduction in the amplitude of the average curve, because of the distortion introduced by the average of few-samples-delayed ensembles. This fact compromises the BPRSA curve, and some investigation should be done in order to standardize the selection of AP's.

4. Conclusions

BRS was evaluated in a sample of critical patients in MV. Results revealed that the capacity of the BRS, measured via BPRSA analysis, is stronger and with negative values for patients who were actually ready for weaning.

Results suggest that BRS quantification is potentially useful for predicting weaning outcomes in the ICU. Future studies will provide useful insights of the role of the ANS in the weaning process that will help reduce weaning failure rates and the very adverse effects of reintubation.

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