

# Prognostic Value of NT-proBNP versus Killip Classification in Patients with Acute Coronary Syndromes

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## Abstract

**Background:** Plasma levels of brain natriuretic peptides have better diagnostic accuracy compared to clinical-radiologic judgment for acute heart failure. In acute coronary syndromes (ACS), the prognostic value of acute heart failure is incorporated into predictive models through Killip classification. It is not established whether NT-proBNP could increment prognostic prediction.

**Objective:** To evaluate whether NT-proBNP, as a measure of left ventricular dysfunction, improves the in-hospital prognostic value of the GRACE score in ACS.

**Methods:** Patients admitted due to acute chest pain, with electrocardiogram and/or troponin criteria for ACS were included in the study. The plasma level of NT-proBNP was measured at hospital admission and the primary endpoint was defined as cardiovascular death during hospitalization. P-value < 0.05 was considered as significant.

**Results:** Among 352 patients studied, cardiovascular mortality was 4.8%. The predictive value of NT-proBNP for cardiovascular death was shown by a C-statistic of 0.78 (95% CI = 0.65–0.90). After adjustment for the GRACE model subtracted by Killip variable, NT-proBNP remained independently associated with cardiovascular death (p = 0.015). However, discrimination by the GRACE-BNP logistic model (C-statistics = 0.83; 95%CI = 0.69–0.97) was not superior to the traditional GRACE Score with Killip (C-statistic = 0.82; 95%CI = 0.68–0.97). The GRACE-BNP model did not provide improvement in the classification of patients to high risk by the GRACE Score (net reclassification index = –0.15; p = 0.14).

**Conclusion:** Despite the statistical association with cardiovascular death, there was no evidence that NT-proBNP increments the prognostic value of GRACE score in ACS. (Arq Bras Cardiol. 2020; 114(4):666-672)

**Keywords:** Acute Coronary Syndrome; Heart Failure; Natriuretic Peptide, Brain; Mortality; Ventricular Dysfunction, Left; Biomarkers.

## Introduction

Brain natriuretic peptide is a prohormone, biologically measured by its active fragment or its inactive terminal portion (NT-proBNP). These molecules are biomarkers of left ventricular dysfunction, released to the bloodstream by myocytes undergoing wall tension due to volumetric or pressure overload.<sup>1</sup> In the detection of heart failure, these peptides present better accuracy than clinical-radiological evaluation, being able to identify sub-clinical levels of decompensation.<sup>2</sup>

The presence of left ventricular dysfunction is an important determinant of prognosis in patients with acute coronary

syndromes (ACS). In this context, multivariate predictive models<sup>3,4</sup> take into account the presence of clinically manifested left ventricular dysfunction, well represented by the classification of Killip and Kimball.<sup>5</sup> Two reasons support the hypothesis that the use of plasma biomarkers may increase the prognostic value of these models: the capacity to numerically quantify the degree of cardiac decompensation and the higher sensitivity for subclinical changes, without impairing specificity.<sup>2</sup>

In the context of ACS, the concentration of NT-proBNP has a well-documented prognostic accuracy.<sup>6</sup> However, from a predictive point of view, whether NT-proBNP has an incremental value in relation to probabilistic models that already contain Killip as a predictor variable is a controversial matter.<sup>7-9</sup> Among the models validated for risk prediction, GRACE score is the one with the best prognostic accuracy, containing Killip class as the marker of heart failure.<sup>10-12</sup> In this cohort, we tested the hypothesis that NT-proBNP incorporation increases the prognostic value of the GRACE score in patients with ACS. NT-proBNP was measured at admission and the primary outcome was defined as cardiovascular death during hospitalization.

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Manuscript received November 13, 2018, revised manuscript May 20, 2019, accepted June 05, 2019

**DOI:** <https://doi.org/10.36660/abc.20180345>

## Methods

### Sample selection

Patients consecutively admitted to the coronary care unit (CCU) of a tertiary-care hospital, between September 2007 and October 2013, due to suspected ACS (unstable angina and myocardial infarction) were prospectively included in the study. Inclusion criteria was chest discomfort in addition to at least one of the three objective criteria:

- 1) positive biological marker of myocardial necrosis, defined as troponin T  $\geq 0.01$   $\mu\text{g/L}$  or troponin I  $> 0.034$   $\text{g/L}$ , corresponding to values above the 99<sup>th</sup> percentile;<sup>13</sup>
- 2) ischemic electrocardiographic alteration, consisting of T wave inversion ( $\geq 0.1$  mV) or ST segment changes ( $\geq 0.05$  mV); and
- 3) previously documented coronary artery disease, defined as a history of myocardial infarction with Q wave or previous angiography demonstrating coronary obstruction  $\geq 70\%$ .

Patients without NT-proBNP dosage or who did not agree to participate in the study were excluded. The protocol was in compliance with the Declaration of Helsinki, was approved by the Research Ethics Committee of the Institution, and all participants provided written informed consent.

### NT-proBNP Measurement

NT-proBNP measurement was performed on a blood sample collected at patient's arrival at the hospital, aiming for a minimum delay between the onset of symptoms and the collection of material. Plasma was frozen at  $-70$  °C for simultaneous dosing of the samples. The immunoassay method (Biomérieux) was used, considering the following definitions of high NT-proBNP:

- 1) Values above 450 pg/ml in patients under 50 years of age;
- 2) Values above 900 pg/ml in patients over 50 years of age.<sup>2</sup>

### GRACE score

The GRACE score calculation was based on clinical data at admission, electrocardiogram performed within 6 hours of admission, troponin T or troponin I dosages in the first 12 hours, and the first plasmatic creatinine. Elevation of myocardial necrosis markers (as a component of the scores) was defined as troponin above the 99<sup>th</sup> percentile. The GRACE score includes eight variables: five semi-quantitative, meaning different weight for each age stratum (systolic blood pressure, heart rate, plasma creatinine and Killip class), and three dichotomic ones (ST segment depression, elevation of myocardial necrosis marker, and cardiac arrest at admission). The final score can range from 0 to 372.<sup>3</sup>

### Clinical end-point

The clinical end-point was cardiovascular death during hospitalization, defined by one of the following mechanisms: cardiac failure, arrhythmia or due to complications from treatments related to ACS.

## Statistical Analyses

Numerical variables were expressed as mean and standard deviation as they presented normal distribution or small deviation from normality, while median and interquartile range were preferred in case of significant deviation from normality. Categorical variables were expressed in proportions. Preliminary results were accompanied by a 95% confidence interval as a measure of uncertainty. Initially, predictive values of NT-proBNP and Killip class were evaluated by the area under the ROC curve (C-statistic), considering cardiovascular death as an outcome. These two curves were statistically compared by the Hanley-McNeil paired test.<sup>14</sup> In addition, Kappa Test was used to assess concordance between high NT-proBNP and Killip  $> 1$  in the definition of heart failure.

Logistic regression was used to assess the incremental value of NT-proBNP to the GRACE Score. The technique of modifying the GRACE Score was used, by replacing Killip for NT-proBNP, and then comparing this GRACE-BNP model to the traditional GRACE. The modification of GRACE was performed in two ways, one numerical and another categorical. In the first case, the regression coefficient of NT-proBNP represented the change in log odds promoted by each unit of NT-proBNP. In this case, the logistic regression equation determined the weight of NT-proBNP (numerical GRACE-BNP). In the second case, a high NT-proBNP added 20 points to the Killip free GRACE, which is the equivalent of Killip II value in the score (categorical GRACE-BNP).

The C-statistics of both models were compared with the traditional GRACE Score by the Hanley-McNeil test. Finally, net reclassification index analysis by Pencina<sup>15</sup> was used to evaluate the reclassification value of logistic and categorical GRACE-BNP in relation to the definition of high risk. For this reclassification, the best cutoff points for these new scores were used in the ROC curve.

Regarding sample size definition, two criteria were used. Firstly, aiming to reach a power of 80% to detect a difference of 0.05 between two ROC curves (referring to scores) and predicting a correlation of 0.80 between the scores, it would be necessary to enroll 192 patients. Secondly, in order to insert two variables in a logistic regression model, 10 to 20 events would be necessary.<sup>16</sup>

All of the tests above were considered statistically significant if  $p$ -value  $< 0.05$ . The SPSS Version 21 was the software used for the analysis.

## Results

### Sample characteristics

The sample consisted of 352 patients, mean age  $63 \pm 14$  years, 60% male, 26% presenting with ST-segment elevation myocardial infarction. The GRACE score had a median of 104 (IIQ 82 - 131), which corresponds to intermediate risk. The median of NT-proBNP was 340 pg/ml (IIQ 86-1212), elevated in 29% of patients. The median time between symptom onset and NT-pro-BNP dosage was 15.5 hours (IIQ 8.2 - 32.5). The incidence of cardiovascular death in the hospital phase was 4.8%. Sample characteristics are described in Table 1.

**Table 1 – Clinical and laboratorial characteristics of the selected sample**

Variable	N
Sample Size	352
Age (years)	63 ± 14
Male Gender	210 (60%)
<b>ACS</b>	
Unstable angina	102 (29%)
NSTEMI	90 (26%)
STEMI	160 (45%)
Triarterial disease or LMCA	170 (48%)
Ischemic ECG	223 (63%)
Positive Troponin	250 (71%)
Creatinine	1.0 ± 0.62
<b>Killip Classification</b>	
Killip I	308 (88%)
Killip II	18 (5%)
Killip III	25 (7%)
Killip IV	1 (0.3%)
NT-proBNP (pg/ml)	340 (86 – 1212)
GRACE score	104 (82 – 131)
Mortality	17 (4.8%)

\*ACS: acute coronary syndrome; NSTEMI: Non ST elevation myocardial infarction; STEMI: ST elevation myocardial infarction; LMCA: left main coronary artery; ECG: electrocardiogram; BNP: brain natriuretic peptide.

### NT-proBNP and Killip: Univariate Predictor Value

NT-proBNP demonstrated a moderate predictive capacity for cardiovascular death, according to C-statistic of 0.78 (95% CI = 0.65-0.90,  $p < 0.001$ ), while the Killip score presented C-statistic of 0.69 (95% CI = 0.54-0.84,  $p = 0.008$ ), with no statistical difference between the two curves ( $p = 0.29$ ) (Figure 1). The two markers agreed in the definition of heart failure in 75% of the cases (8% with heart failure and 67% without heart failure), meaning low level of agreement according to the Kappa test ( $\kappa = 0.26$ ; 95% CI = 0.54-0.84;  $p < 0.001$ ).

### Independent and Incremental NT-pro-BNP Value

In the logistic regression analysis, numerical NT-proBNP did not maintain statistical significance after adjustment for the traditional GRACE score ( $p = 0.11$ ). On the other hand, numerical NT-proBNP remained an independent predictor when adjusted for GRACE score without Killip ( $p = 0.015$ ; for each 500 pg/ml increase in NT-proBNP, a Beta of 0.029 was observed, OR = 1.03; 95% CI = 1.006 - 1.05) (Table 2). Categorical NT-proBNP was not an independent predictor after adjustment for the GRACE score ( $p = 0.91$ ) or for the GRACE score without Killip ( $p = 0.36$ ).

For analysis of the incremental value of NT-proBNP to GRACE, we compared the C-statistics of the logistic GRACE-BNP, categorical GRACE-BNP and traditional GRACE

score. The results of the analysis were, respectively, 0.83 (95% CI = 0.69-0.97), 0.82 (95% CI = 0.68-0.96), and 0.82 (95% CI = 0.68-0.97). Therefore, no incremental value of the new approaches was identified (Figure 2).

### Reclassification of GRACE Score by NT-pro-BNP

Regarding the net reclassification analysis, of the 17 patients who died, 3 were correctly reclassified by logistic GRACE-BNP from low to high risk, with no incorrect reclassification, resulting in a positive net reclassification index (+ 0.18%). Among the 335 patients who survived, 9 were erroneously reclassified from low to high risk, while there was no correct reclassification. This resulted in a negative net reclassification ratio (-0.02%). In the final analysis, considering all patients, the total net reclassification index (NRI) was - 0.15% ( $p = 0.14$ ) (Table 03). Reclassification based on categorical GRACE-BNP showed similar results (NRI = 0.08;  $p = 0.44$ ). (Table 3)

### Discussion

The present study demonstrates the independent prognostic value of numeric NT-proBNP after adjustment to the GRACE score. However, the NT-proBNP did not improve discrimination of the GRACE Score, nor its reclassification ability. Its findings are in line with the notions that not every independent predictor offers incremental value to traditional models.<sup>17</sup>

In an explanatory point of view, our findings reinforce that the status of cardiac decompensation increases the risk of patients with ACS. On the other hand, from the predictive point of view, refining the prognostic evaluation with a biomarker of heart failure that is more accurate than clinical evaluation was not enough to increase the accuracy of multivariate models. This discussion is intended to debate the potential explanations for the absence of an incremental value, to confront this paper's results with external evidence, to recognize methodological limitations and to address the relevance of the present results.

Different hypotheses may explain the absence of NT-proBNP incremental value. Three possibilities will be pointed out, which comprises the generic properties of predictors and the specificities of the clinical context in question. First, probabilistic models are created with variables that simultaneously contribute to risk prediction, each with a predictive weight that is proportional to its independent strength of association. The improvement of a single predictor (detection of ventricular dysfunction) among many may not represent a relevant change. In the present case, the incorporation of a marker related to a new phenomenon was not proposed, but rather only the replacement of the evaluation of the phenomenon of heart failure with a theoretically better marker. Second, the predictive capacity of NT-proBNP theoretically lies in its continuous characteristic (numerical variable) and in its ability to identify subclinical ventricular dysfunction. It is possible that the prognostic value of heart failure is not at initial levels, limiting to more advanced and clinically manifested degrees. Finally, the prognostic accuracy of the traditional GRACE Score is already satisfactory, represented by C-statistic above 0.8, making more difficult to improve a marker that functions with good predictive capacity.

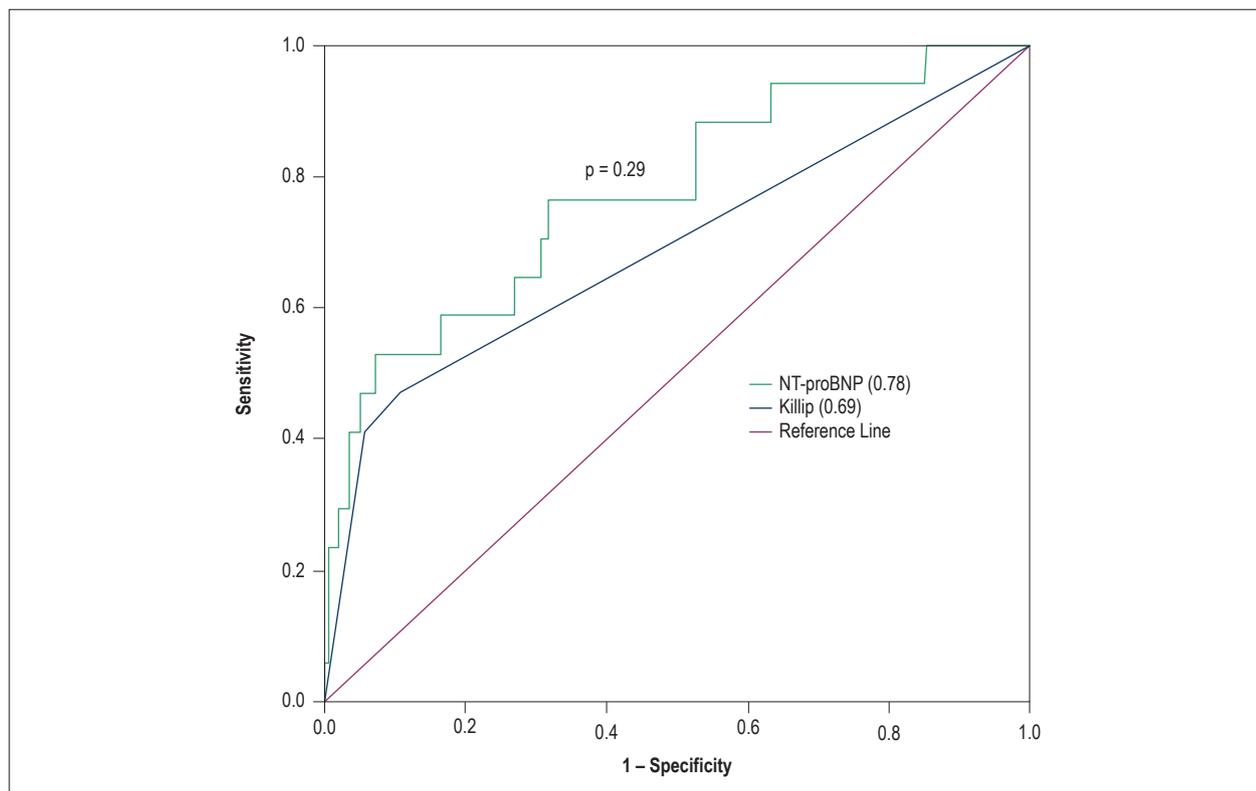


Figure 1 – The accuracy of NT-pro-BNP in the prediction of death has a value of 0.78 (95% CI 0.65 – 0.9) in the C statistic and in the Killip classification 0.69 (95% CI 0.54 – 0.84).

Table 2 – Logistic regression model containing GRACE without Killip and numerical NT-proBNP in predicting deaths

Variables	Beta Coefficient	Odds ratio (95% IC)	p Value
GRACE without Killip	0.043	1.04 (1.02 – 1.06)	<0.001
NT-proBNP / 500 pg/ml	0.000	1.03 (1.006 – 1.05)	0.015

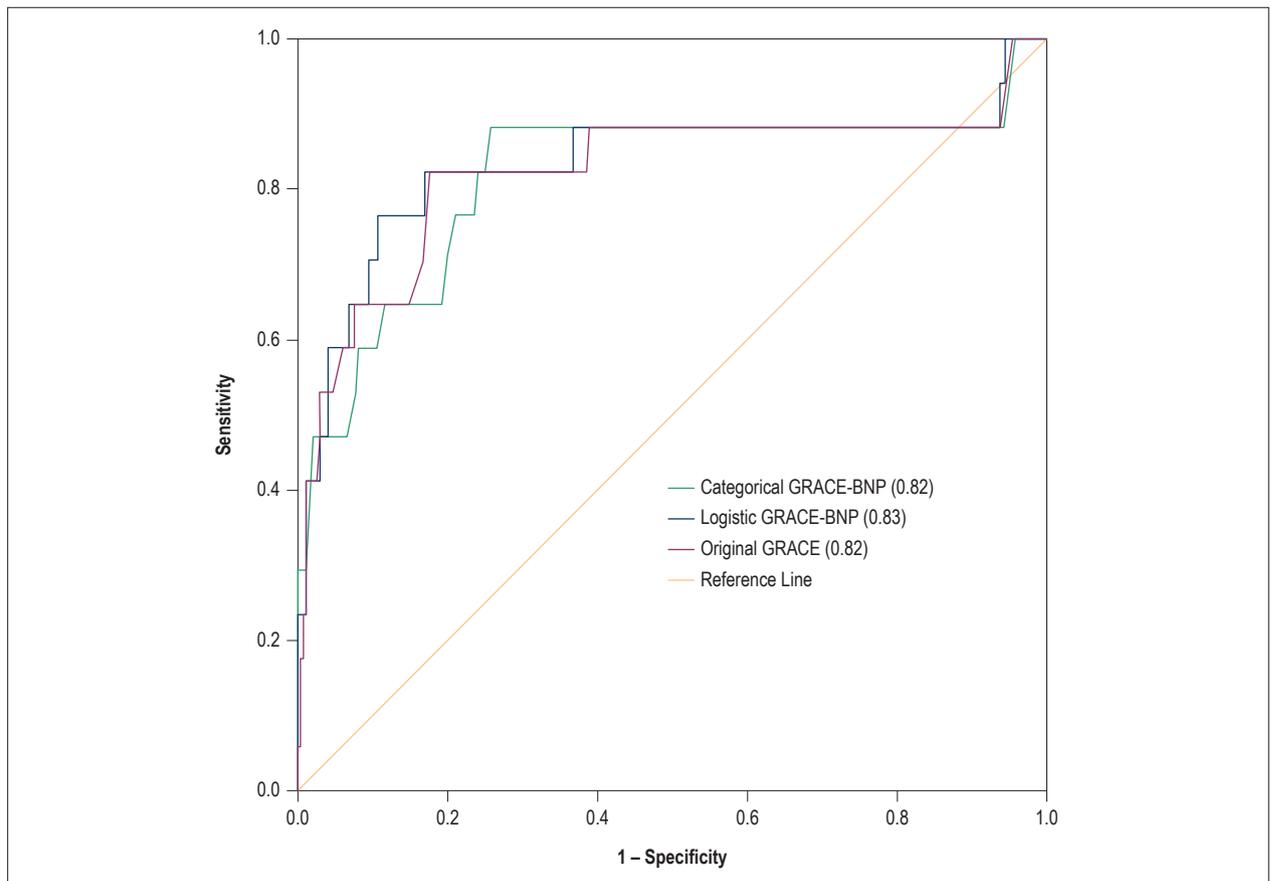
Some previous studies have tested the prognostic value of brain natriuretic peptides in ACS. Although there is a disagreement between studies, a careful analysis of the results shows that they all point in the same direction. Three studies conclude positively regarding the prognostic value of this type of marker; however, these studies evaluated the independent predictive value but did not test incremental value (discrimination or reclassification).<sup>18-22</sup> In this context, our results are not discordant. However, our negative conclusion resides in a more comprehensive analysis that was not previously done. In concordance, the two studies that evaluated the incremental value of multivariate models presented the same conclusion as ours.<sup>7,8</sup>

Two aspects are original in the present work: it was the first study to aggregate the analysis of reclassification proposed by Pencina and the only one to adjust for the GRACE score after removal of Killip, avoiding eventual collinearity between Killip and NT-proBNP which could induce type II error. These approaches bring more veracity to our negative outcome.

Methodological limitations must be recognized here, which may have promoted a false negative result. Firstly, it is known that, ideally, a risk marker should be tested in an environment where the care team is not aware of its outcome. As this marker

is already available in our clinical practice, the team became aware of the NT-proBNP result, predisposing to performance bias, which could improve the prognosis of patients with high NT-proBNP. Secondly, although this study had the planned sample size, it lacked additional power for exploratory analyses. For example, it was not possible to test the incremental value of the best NT-proBNP cutoff point. To do so, it would require a sample to identify the best cutoff point and another one to test for its incremental value. However, given our sample size, we chose not to split the sample.

The value of a negative result should be contextualized. Frequently, improper evaluation of markers modifies clinical reasoning with no probabilistic basis. That is, after estimating the risk based on the GRACE score, our evaluation would become less accurate if we mentally increased the risk after observing a high NT-proBNP value. It would be an improper reclassification. Therefore, it must be considered that the GRACE score has better accuracy than NT-proBNP, which should not modify the message of the first. On the other hand, the absence of prognostic value should not discredit the value of BNP in diagnosing symptoms of dyspnea during hospitalization or in monitoring the volemic status of patients who developed acute heart failure.



**Figure 2** – Comparison of the ROC curves between original GRACE (0.82; 95% CI = 0.68-0.97) and GRACE-BNP logistic (0.83; 95% CI = 0.69-0.97) and categorical (0.82; 95% CI = 0.68-0.96) shows similar C-statistics among the three scores.

**Table 3** – Analysis of net reclassification by the GRACE-BNP numerical score in relation to the GRACE score in the definition of high risk

	N	Reclassification to high risk	Reclassification to low risk	NRI	p Value
Outcome	17	3	0	+0.18%	
Without outcome	335	9	0	-0.02%	
Total	352	12	0	-0.15%	0.14

## Conclusion

Despite its association with risk in a univariate approach, it has not been proven that the use of NT-proBNP as a measure of left ventricular dysfunction increases the in-hospital prognostic value of GRACE score in ACS.

## Author contributions

Conception and design of the research: Souza TMB, Cerqueira Jr. AMS, Correia L; Acquisition of data: Souza TMB, Cerqueira Jr. AMS, Suerdieck JG, Sá NC, Sodr  GS, Correia VCA, Lacerda YF, Fonseca LL, Noya-Rabelo MM; Analysis and interpretation of the data: Souza TMB, Cerqueira Jr. AMS, Suerdieck JG, S  NC, Sodr  GS; Statistical analysis: Souza TMB, Cerqueira Jr. AMS, Suerdieck JG, S  NC, Correia VCA, Lacerda YF, Fonseca LL, Noya-Rabelo MM,

Correia L; Critical revision of the manuscript for intellectual content: Souza TMB, Cerqueira Jr. AMS, Suerdieck JG, S  NC, Sodr  GS, Correia VCA, Lacerda YF, Fonseca LL, Noya-Rabelo MM, Correia L.

## Potential Conflict of Interest

No potential conflict of interest relevant to this article was reported.

## Sources of Funding

There were no external funding sources for this study.

## Study Association

This study is not associated with any thesis or dissertation work.

### Ethics approval and consent to participate

This study was approved by the Ethics Committee of the Monte Tabor under the protocol number 36/11. All the

procedures in this study were in accordance with the 1975 Helsinki Declaration, updated in 2013. Informed consent was obtained from all participants included in the study.

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