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Please cite this article as: Monneret D. High-sensitive cardiac troponin after CPAP in obstructive sleep apnoea: the adjusted analytical change limit (adjACL) for small variations at low concentrations. *Eur Respir J* 2022; in press (https://doi.org/10.1183/13993003.03022-2021).

This manuscript has recently been accepted for publication in the *European Respiratory Journal*. It is published here in its accepted form prior to copyediting and typesetting by our production team. After these production processes are complete and the authors have approved the resulting proofs, the article will move to the latest issue of the ERJ online.

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High-sensitive cardiac troponin after CPAP in obstructive sleep apnoea: the adjusted analytical change limit (adjACL) for small variations at low concentrations

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Keywords: cardiac troponin, CPAP, obstructive sleep apnoea, analytical imprecision

Even below the 99<sup>th</sup> percentile upper reference limit (99PURL), elevated high-sensitive cardiac troponin concentration (hs-cTn) is associated with increased cardiovascular risk [1]. However, the lower the troponin concentration, the higher the analytical imprecision [2]. It is unclear to what extent a change in hs-cTn (Δhs-cTn) is significant between the limit of detection (LOD) and the 99PURL. Recently, Lui et al. compared, through a recent randomized control trial, the effect of 8 weeks of continuous positive airway pressure (CPAP) on \( \Delta hs-cTnI \) in subjects with moderate-to-severe obstructive sleep apnoea (OSA) and hypertension [3]. They showed a statistically significant variation compared to untreated subjects (adjusted mean difference -1.74 ng·L<sup>-1</sup>; p=0.006). The adjusted mean of hs-cTnI decreased from 5.6 to 4.8 ng·L<sup>-1</sup> (-14.3%) after CPAP, and increased from 7.9 to 8.5 ng·L<sup>-</sup> <sup>1</sup> (+7.6%) in the control group. These values are above the LOD of 1.2 ng·L<sup>-1</sup>, but are close to the limit of quantification of 5 ng·L<sup>-1</sup> (hs-cTnI reagent kit 3P25, Abbott, Chicago, US). From a diagnostic standpoint, at the individual scale, the reference change value (RCV) concept assumes that the change between two serial results is significant if greater than the sum of the variations for each result. These variations are mostly analytical (analytical coefficient of variation, CVa) and biological (intraindividual, CVi) [2]. Considering two serial results, the **RCV** equal  $Z^*[(CVa_1^2+CVi_1^2)+(CVa_2^2+CVi_2^2)]^{1/2}$ , with Z=1.96 for a probability of change of 95%. Considering only analytical imprecision -which mainly depends on assay methods and analyzers- as the minimal unavoidable variation, and applying it to serial results, the adjusted analytical change limit (adjACL) is equal to  $\pm 1.96*[(CVa_1^2)+(CVa_2^2)]^{1/2}$  [4]. To test whether the  $\Delta$ hs-cTnI observed after 8 weeks of CPAP was greater than the adjACL of hs-cTnI, we used CVa for hs-cTnI concentrations ranging from 50 to 1 ng·L<sup>-1</sup> (mean imprecision profile), provided in the supplemental data from [2], and measured on an Architect Abbott analyser, as did Lui et al. [3]. The regression equation of the best-fit curve was CVa = 3.997+28.17/hs-cTnI (Figure 1). Applying this formula on serial hs-cTnI means of 5.6 and 4.8 ng·L<sup>-1</sup> (before and after CPAP), the adjACL was  $\pm 26.2\%$ . Hence, at such low concentrations, the  $\Delta$ hscTnI of -14.3% is within the adjACL, meaning that the effect of CPAP is not great enough to decrease troponin more than analytical variability. According to this model, a decrease from 5.6 to at least 4.0 ng·L<sup>-1</sup> ( $\Delta$ hs-cTnI  $\leq$  -28.6%, i.e., out of adjACL  $\pm 28.0\%$ ) would have been considered as significantly due to CPAP, at least in part. As expected,  $\Delta$ hs-cTnI of the control group (7.9 to 8.5 ng·L<sup>-1</sup>,  $\pm 7.6\%$ ) remains within the adjACL, calculated at  $\pm 20.6\%$ . As demonstrated, the closer the concentrations are to the LOD, the greater the imprecision. This regression model reflects real analytical imprecisions for low ranges of hs-cTnI in the laboratory, with concentrations of 10%CV and 20%CV at 4.7 and 1.8 ng·L<sup>-1</sup> [5]. Another analytical consideration is lot-to-lot bias between hs-cTnI reagents and/or calibrators, which can occur over weeks and could partly bias the mean difference of  $\pm 1.74$  ng·L<sup>-1</sup> observed by Lui *et al.* Illustrating this, Wu *et al.* recently showed for Abbott hs-cTnI a lot-to-lot bias at low ranges (below 5.4 ng·L<sup>-1</sup>) between  $\pm 1.7$  to  $\pm 1.2$  ng·L<sup>-1</sup> [6]. A large difference of 2.5 ng·L<sup>-1</sup> between two successive lots was even observed (from 4.7 to 7.2 ng·L<sup>-1</sup>), leading the authors to conclude that "a total analytic error  $\pm 1.5$  ng·L<sup>-1</sup> for long-term studies with hs-cTnI concentrations  $\pm 1.5$  ng·L<sup>-1</sup> could prevent erroneous reporting of results" [7]. This bias is advantageously considered in the adjACL, since it is also determined over the course of weeks.

Analytical imprecision of troponin is rarely considered in clinical studies, and even less when outcomes rely on serial results. Few studies have evaluated the effect of CPAP on hs-cTnT in OSA, but they overall concluded to an absence of significant effect [8, 9]. Lui *et al.* were the first to focus on Δhs-cTnI under CPAP, furthermore in a well-framed RCT. They argued that hs-cTnT is "*less sensitive than hs-cTnI* in *detecting subclinical myocardial injury*", but this is not consensually admitted to date. Hs-cTnI is more sensitive in the sense that a greater proportion of the general population has detectable concentrations (roughly 75% *versus* 53% for hs-cTnT), but it also has the drawback of being more strongly associated with age, male sex, BMI, and SBP (p <0.0001 for all *versus* hs-cTnT) [10], making these factors potentially more confounding. Cardiovascular co-morbidities are generally numerous in OSA, most of them impacting troponin levels. This is probably why most studies failed to show beneficial effects of CPAP on cardiovascular outcomes [11, 12]. Further studies interpreting small changes in cardiac troponin at low concentrations under CPAP in OSA should consider the adjACL as an objective criterion, requiring a close collaboration between clinicians and laboratorians.

**Acknowledgements**: The author is grateful to Vincent Fitzpatrick for his English proofreading.

**Conflict of interest:** None declared.

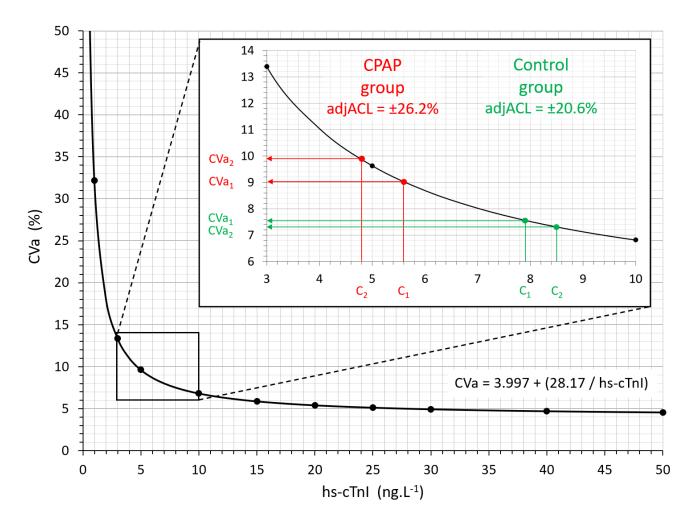
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## Legend

Figure 1.



Mean imprecision profile of high-sensitive cardiac troponin I (hs-cTnI) determined on Architect analyser (Abbott, Chicago, US), using CVa and mean hs-cTnI concentrations provided in supplemental data from [2]. CVa<sub>1</sub> and CVa<sub>2</sub> are the analytical coefficients of variation corresponding to the hs-cTnI concentrations before (C<sub>1</sub>) and after (C<sub>2</sub>) an 8-week period with (red) or without (green) CPAP treatment, and calculated according to the equation of the nonlinear regression curve. The adjusted analytical change limits (adjACL) were calculated according to the formula  $\pm 1.96*[(CVa_1^2) + (CVa_2^2)]^{1/2}$  [4].