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The influence of atrial fibrillation on the levels of NT-proBNP versus GDF-15 in patients with heart failure

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Abstract

Background In heart failure (HF), levels of NT-proBNP are influenced by the presence of concomitant atrial fibrillation (AF), making it difficult to distinguish between HF versus AF in patients with raised NT-proBNP. It is unknown whether levels of GDF-15 are also influenced by AF in patients with HF. In this study we compared the plasma levels of NT-proBNP versus GDF-15 in patients with HF in AF versus sinus rhythm (SR).

Methods In a post hoc analysis of the index cohort of BIOSTAT-CHF ($n = 2516$), we studied patients with HF categorized into three groups: (1) AF at baseline ($n = 733$), (2) SR at baseline with a history of AF ($n = 183$), and (3) SR at baseline and no history of AF ($n = 1025$). The findings were validated in the validation cohort of BIOSTAT-CHF ($n = 1738$).

Results Plasma NT-proBNP levels of patients who had AF at baseline were higher than those of patients in SR (both with and without a history of AF), even after multivariable adjustment (3417 [25th–75th percentile 1897–6486] versus 1788 [682–3870], adjusted $p < 0.001$, versus 2231 pg/mL [902–5270], adjusted $p < 0.001$). In contrast, after adjusting for clinical confounders, the levels of GDF-15 were comparable between the three groups (3179 [2062–5253] versus 2545 [1686–4337], adjusted $p = 0.36$, versus 2294 [1471–3855] pg/mL, adjusted $p = 0.08$). Similar patterns of both NT-proBNP and GDF-15 were found in the validation cohort.

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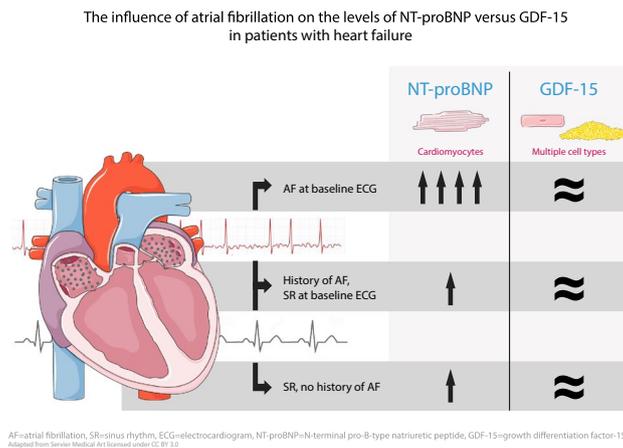
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Conclusion These data show that in patients with HF, NT-proBNP is significantly influenced by underlying AF at time of measurement and not by previous episodes of AF, whereas the levels of GDF-15 are not influenced by the presence of AF. Therefore, GDF-15 might have additive value combined with NT-proBNP in the assessment of patients with HF and concomitant AF.

Graphic abstract



Keywords Atrial fibrillation · Heart failure · Biomarkers · Natriuretic peptides · GDF-15

Abbreviations

| | |
|-------------|--|
| AF | Atrial fibrillation |
| BIOSTAT-CHF | The BIOlogy Study to Tailored Treatment in Chronic Heart Failure |
| ECG | Electrocardiogram |
| GDF-15 | Growth differentiation factor-15 |
| HF | Heart failure |
| LVEF | Left ventricular ejection fraction |
| NT-proBNP | N-terminal pro-B-type natriuretic peptide |
| SR | Sinus rhythm |

Introduction

N-terminal pro-B-type natriuretic peptide (NT-proBNP) plays an important role in the diagnosis and prognosis of heart failure (HF) [1, 2]. For the diagnosis of HF, this marker is known for its high sensitivity, but lower specificity, which makes NT-proBNP especially helpful to rule out HF [3]. Several other conditions, such as renal failure, pulmonary embolism and atrial fibrillation (AF), are also known to further elevate NT-proBNP levels in patients with concomitant HF. AF in this regards is particularly important because it is highly prevalent among patients with HF regardless of ejection fraction, and mimics the symptoms (breathlessness) and signs (left atrial enlargement) of HF. Therefore, it is often unclear how elevated NT-proBNP levels in patients with HF

and AF should be interpreted, since there are several potential explanations for these elevated levels [4–6]. First, NT-proBNP elevations may be directly related to the immediate hemodynamic alterations during the actual episode of AF [7]. Secondly, elevated levels of NT-proBNP may be related to the chronic structural or functional cardiac remodeling as a result of sustained episodes of AF, or in the third place, just reflect that patients with AF have more advanced HF. In most contemporary clinical HF trials, different NT-proBNP thresholds are being used for inclusion of patients with and without AF, often without further differentiation between patients who only have a history of AF, and those who have AF at time of enrollment.

In the past years, many markers have been shown to have strong prognostic value in HF, of which growth differentiation factor-15 (GDF-15) is amongst the best established ones [8–11]. GDF-15 is a protein belonging to the transforming growth factor-beta superfamily, and has a role in inflammatory and apoptotic cell processes, and is produced by multiple organs, including the heart [12–14]. It is, however, unknown how the plasma levels of GDF-15 are influenced by underlying AF in patients with HF. The search for a biomarker that is less influenced by underlying AF than NT-proBNP is, could be of help in the assessment of patients with both HF and AF.

To investigate whether the levels of GDF-15 are similarly elevated as NT-proBNP by concomitant AF in patients with HF, we performed a post hoc analysis of these two

biomarkers in The BIOlogy Study to Tailored Treatment in Chronic Heart Failure (BIOSTAT-CHF) [15].

Methods

Patient population and definitions

In the multinational, prospective, observational index cohort of BIOSTAT-CHF, 2516 patients with new-onset or worsening signs and/or symptoms of HF from 11 European countries were included between 2010 and 2012 [15]. Patients had to have evidence of cardiac dysfunction documented either by left ventricular ejection fraction (LVEF) of $\leq 40\%$ or plasma concentrations of NT-proBNP > 2000 pg/mL (this cutoff was the same for patients in sinus rhythm [SR] and AF). A comparable validation cohort of BIOSTAT-CHF included another 1738 patients from six centers in Scotland between 2010 and 2014, who had to have a previously documented admission for HF. No additional LVEF or NT-proBNP was used for the validation cohort, which resulted in a higher percentage of patients with heart failure with preserved ejection fraction (HFpEF, 34% in the validation cohort versus 7% in the index cohort).

In both cohorts, a standard 12-lead electrocardiogram (ECG) was performed at baseline, generally on the same day as the time of blood draw for biomarker measurements (median difference of 0 days with 25th–75th percentile [Q1–Q3] from -2 to $+2$ days). Patients were categorized into three groups based on history and baseline ECG: (1) AF at baseline, (2) history of AF but in SR at baseline, and (3) SR at baseline and no previously documented episode of AF. Patients with a rhythm other than SR or AF on the baseline ECG were excluded (pacemaker rhythm, $n = 283$, and unknown rhythm, $n = 55$) [16]. The study was conducted according to the Declaration of Helsinki, approved by the medical ethics committees of participating centers, and all patients provided informed consent.

Biomarkers

Measurement of NT-proBNP and GDF-15 was performed at baseline. The levels of NT-proBNP and GDF-15 were measured using electrochemiluminescence on a cobas e 411 analyzer, using standard methods (Roche Diagnostics GmbH, Mannheim, Germany) [17, 18].

Statistical analyses

Normally distributed variables were displayed as mean with standard deviation, non-normally distributed variables as median with 25th–75th percentile, and categorical

variables as numbers with percentages. Group differences were assessed with *t tests* and one-way analysis of variance for normally distributed variables, Kruskal–Wallis and Mann–Whitney *U tests* for non-normally distributed continuous variables, and χ^2 tests for categorical variables. Multiple linear regression models were used to investigate the associations between NT-proBNP and GDF-15 and the three rhythm groups. Natural transformed biomarkers were used in the regression analyses. Potential and known confounders of the two biomarkers were included in the regression model, including age, sex, body mass index (BMI), LVEF, heart rate, renal disease [estimated glomerular filtration rate (eGFR) using the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) formula], a previous myocardial infarction, diabetes mellitus, and the use of ACE-inhibitors/ARBs and beta-blockers at baseline. To test which variables had the strongest association with elevated levels of NT-proBNP and GDF-15, both biomarkers were analyzed as a dependent variable in uni- and multi-variable linear regression analysis. The multivariable models were built with all variables with $p < 0.10$ in the univariable analysis, after which backward elimination was performed. Variables that had the strongest associations in both the index and validation cohort were displayed in the final multivariable model. A *p* value of < 0.1 was considered significant for testing interactions. *p* values of < 0.05 were considered statistically significant in all other analyses. All analyses were conducted with R version 3.5.2 (R Foundation for Statistical Computing, Vienna, Austria).

Results

Index cohort

A total of 1941 patients with HF were studied, of whom 733 patients had AF at baseline, 183 patients had a history of AF but were in SR at baseline, and 1025 had SR at baseline and no previously documented episode of AF. The characteristics of the patients within these three rhythm groups are summarized in Table 1. Main findings were that patients with AF at baseline were significantly older, had a higher BMI and higher heart rate, and less often had a previous myocardial infarction as compared with the two other groups who were in SR at baseline.

The plasma levels of NT-proBNP were significantly higher in patients who had AF at baseline, with a median of 3417 pg/mL (1897–6486), as compared with patients who were in SR at baseline; both those who had a history of AF (1788 pg/mL [682–3870], $p < 0.001$) and those who never had AF before (1588 pg/mL [902–5270], $p < 0.001$) (Table 1), also after multivariable adjustment (Table 2). In univariable analysis, the levels of GDF-15 were also highest

Table 1 Baseline characteristics of the index cohort, stratified by heart rhythm

| Clinical characteristic | AF at baseline N= 733 (38%) | History of AF–SR at baseline N= 183 (9%) | Sinus rhythm N= 1025 (53%) | P for trend |
|---------------------------------|--------------------------------|---|-------------------------------|-------------|
| Age (years) | 76 ± 10 | 72 ± 11 | 70 ± 13 | < 0.001 |
| Women (%) | 182 (25) | 51 (28) | 301 (29) | 0.110 |
| BMI (kg/m ²) | 28.5 ± 5.6 | 28.3 ± 5.1 | 27.5 ± 5.5 | 0.001 |
| NYHA (%) | | | | 0.010 |
| I/II | 202 (28) | 68 (37) | 331 (32) | |
| III | 233 (36) | 37 (23) | 270 (31) | |
| IV | 28 (4) | 7 (4) | 33 (4) | |
| LVEF, % | 33 ± 12 | 33 ± 11 | 29 ± 10 | < 0.001 |
| Systolic blood pressure (mmHg) | 125 ± 22 | 126 ± 24 | 126 ± 22 | 0.780 |
| Diastolic blood pressure (mmHg) | 76 ± 14 | 75 ± 16 | 75 ± 13 | 0.188 |
| Heart rate (beats/min) | 93 ± 25 | 73 ± 16 | 79 ± 18 | < 0.001 |
| History of (%) | | | | |
| Myocardial infarction | 215 (29) | 67 (37) | 431 (42) | < 0.001 |
| Stroke | 83 (11) | 27 (15) | 72 (7) | < 0.001 |
| Hypertension | 470 (64) | 118 (65) | 623 (61) | 0.300 |
| Diabetes mellitus | 232 (32) | 63 (34) | 320 (31) | 0.691 |
| COPD | 133 (18) | 46 (25) | 150 (15) | 0.001 |
| Medication (%) | | | | |
| ACE-inhibitors/ARBs | 504 (69) | 135 (74) | 770 (75) | 0.012 |
| Beta-blockers | 599 (82) | 160 (87) | 850 (83) | 0.185 |
| Loop diuretics | 732 (100) | 183 (100) | 1024 (100) | 0.873 |
| Amiodarone | 92 (13) | 67 (37) | 117 (11) | < 0.001 |
| Digoxin | 282 (39) | 21 (12) | 72 (7) | < 0.001 |
| Verapamil/diltiazem | 18 (3) | 4 (2) | 7 (1) | 0.008 |
| Class 1c antiarrhythmic drugs | 2 (1) | 5 (3) | 2 (1) | < 0.001 |
| Ivabradine | 0 (0) | 2 (1) | 26 (3) | < 0.001 |
| Laboratory data | | | | |
| eGFR | 58.0 ± 21.8 | 59.2 ± 20.8 | 66.0 ± 23.4 | < 0.001 |
| NT-proBNP (pg/mL) | 3417 [1897, 6486] | 1788 [682, 3870] | 2231 [902, 5270] | < 0.001 |
| GDF-15 (pg/mL) | 3197 [2062, 5253] | 2545 [1686, 4337] | 2294 [1471, 3855] | < 0.001 |

AF atrial fibrillation, SR sinus rhythm, BMI body mass index, NYHA New York Heart Association, LVEF left ventricular ejection fraction, COPD chronic obstructive pulmonary disease, ACE angiotensin converting enzyme, ARB angiotensin receptor blockers, eGFR estimated glomerular filtration rate, NT-proBNP N-terminal pro-B-type natriuretic peptide, GDF15 growth differentiation factor 15

in patients with AF at baseline and lowest in patients who were in SR at baseline (Table 1), but after adjusting for clinical confounders, the levels of GDF-15 between patients with AF at baseline and those in SR with and without previous AF were comparable (Table 2).

Validation cohort

Baseline characteristics of the validation cohort were generally comparable to the index cohort of BIOSTAT-CHF (Supplementary Table 1). As discussed previously, a higher number of patients with HFpEF were included in the validation cohort, which resulted in a higher number of women, a higher LVEF, and lower levels of NT-proBNP in all three groups. Despite these differences in baseline characteristics,

similar patterns of plasma levels of both NT-proBNP and GDF-15 were found in the validation cohort as compared to the index cohort. Patients with AF at baseline had a median NT-proBNP of 2105 pg/mL (1015–4472), which was significantly higher than those who had a history of AF but were in SR at baseline (1063 [440–4094], $p < 0.001$) and patients in SR who never had AF before (874 [314–2758]). The levels of GDF-15 were comparable among the three groups after multivariable adjustment (Table 2). No significant interactions between the biomarkers and the rhythm groups in heart failure with reduced ejection fraction (HFrEF) versus HFpEF were found in both the index and validation cohort.

Table 2 Multivariable differences of the plasma levels of NT-proBNP and GDF-15 between the three rhythm groups in the index and validation cohort

| Index cohort | | | | | |
|------------------------------|---------------|---------------|------------------------------|--------|------|
| | NT-proBNP | | | GDF-15 | |
| AF at baseline | } $p < 0.001$ | } $p < 0.001$ | AF at baseline | } NS | } NS |
| History of AF—SR at baseline | | | History of AF—SR at baseline | | |
| Sinus rhythm | | | Sinus rhythm | | |
| Validation cohort | | | | | |
| | NT-proBNP | | | GDF-15 | |
| AF at baseline | } $p = 0.02$ | } $p < 0.001$ | AF at baseline | } NS | } NS |
| History of AF—SR at baseline | | | History of AF—SR at baseline | | |
| Sinus rhythm | | | Sinus rhythm | | |

Adjusted for age, sex, body mass index, left ventricular ejection fraction, heart rate, estimated glomerular filtration rate, a previous myocardial infarction, diabetes mellitus, and the use of ACE-inhibitors/ARBs and beta-blockers at baseline

NT-proBNP N-terminal pro-B-type natriuretic peptide, *GDF15* growth differentiation factor 15, *NYHA* New York Heart Association, *BMI* body mass index, *LVEF* left ventricular ejection fraction, *eGFR* estimated glomerular filtration rate, *NS* non-significant

Correlates of NT-proBNP versus GDF-15

The multivariable models with the correlates of elevated levels of NT-proBNP and GDF-15 are presented in Table 3. AF at baseline was strongly associated with elevated levels of NT-proBNP in both the index and validation cohort. Other variables that were strongly associated with elevated levels of NT-proBNP were age, BMI, LVEF and eGFR. AF at baseline was not associated with higher levels of GDF-15 in the multivariable model. Variables that were strongly associated with higher levels of GDF-15 in both cohorts were age, systolic blood pressure, diabetes mellitus and eGFR.

Discussion

These data suggest that in patients with HF, after adjustment for clinical confounders, the levels of NT-proBNP, but not GDF-15, are significantly influenced by the presence of AF at time of measurement. GDF-15 is mainly produced in non-cardiac and peripheral tissues, such as endothelial cells and adipocytes, and we recently showed that the levels of GDF-15 in mice are 2- to 60-fold higher in the liver, lungs and kidney than in the cardiac muscle [19]. Therefore, this marker reflects changes in many organs, not just in the cardiac ventricles and atria, and might therefore be less likely to be load-dependent as compared with NT-proBNP. Since levels of GDF-15 are independent of the presence of AF in patients with HF, it may better reflect HF patients' overall clinical condition, including non-cardiac comorbidities [20].

Previous studies have shown that the levels of GDF-15 in AF patients without HF are fairly similar to the levels in community-dwelling elderly [21]. In the AF field, GDF-15 is of increasing interest since this biomarker was the strongest predictor of major bleeding, stroke and mortality in the ARISTOTLE trial (the Apixaban for Reduction in Stroke and Other Thromboembolic Events in Atrial Fibrillation) and RE-LY (Randomized Evaluation of Long-Term

Anticoagulant Therapy), and is one of the strongest prognostic factors in the ABC (age, biomarkers, comorbidities) score, the new score for assessing risk of AF patients [22, 23].

The diagnosis of HF (especially HFpEF) in patients with AF remains a clinical challenge, since signs and symptoms, echocardiographic abnormalities and elevated NT-proBNP levels can be caused by both AF alone and by AF with concomitant HFpEF. Since GDF-15 has previously been shown to have diagnostic utility in HFpEF, with similarly or even more elevated levels as compared with patients with HFrEF, and seems to be less influenced by concomitant AF in patients with HF as shown in the present study, it could perhaps be a suitable companion marker next to NT-proBNP to diagnose the presence or absence of HFpEF in patients presenting with AF [9, 24, 25]. This novel hypothesis should be further explored by studying the potential usefulness of GDF-15 and its clinical consequences. The combination of GDF-15 and NT-proBNP to distinguish AF versus HF by additionally comparing levels in patients with AF without HF, as well as by comparing levels in patients with AF before and after cardioversion needs further investigation.

In clinical practice, NP levels are often used for therapy guidance, but these levels can fluctuate in patients with HF and paroxysmal AF, depending on whether they are in SR or AF at the time of measurement [26, 27]. Since NT-proBNP is mainly produced and secreted by the cardiomyocytes in the atria and ventricles in response to haemodynamic wall stress, this marker is known to be sensitive to heart rate and rhythm disturbances [5, 7].

As described previously, most contemporary clinical HF trials use different NP thresholds for the inclusion of patients with and without AF [28–30]. These higher thresholds in patients with AF increase the probability that these patients have actual underlying HF, instead of including patients who have merely AF—a distinction that is especially challenging in patients with HFpEF and AF [31]. For patients with a history of AF but who are in SR at time of blood collection,

Table 3 Multivariable model with the strongest associated correlates of NT-proBNP and GDF-15 in the index and validation cohort

| Variable | NT-proBNP | | | GDF-15 | | |
|---------------------------------|---------------|---------|---------|-------------------|---------|---------------|
| | Index cohort | | | Validation cohort | | |
| | Beta Coeff | T value | P value | Beta Coeff | T value | P value |
| Age | 0.014 | 3.291 | 0.001 | 0.021 | 3.613 | <0.0001 |
| BMI | -0.071 | -8.846 | <0.0001 | -0.076 | -9.015 | <0.0001 |
| Systolic blood pressure | NS | NS | NS | NS | NS | NS |
| LVEF | -0.022 | -5.232 | <0.0001 | -0.028 | -7.402 | <0.0001 |
| Atrial fibrillation at baseline | 0.438 | 4.550 | <0.0001 | 0.766 | 7.163 | <0.0001 |
| Diabetes mellitus | NS | NS | NS | NS | NS | NS |
| Heart rate | 0.017 | 8.004 | <0.0001 | 0.016 | 6.669 | <0.0001 |
| eGFR | -0.022 | -10.469 | <0.0001 | -0.023 | -9.070 | <0.0001 |
| NYHA class IV | 0.821 | 3.343 | 0.0008 | 1.597 | 3.168 | 0.002 |
| | $r^2 = 0.261$ | | | $r^2 = 0.353$ | | |
| | | | | $r^2 = 0.244$ | | |
| | | | | | | $r^2 = 0.335$ |

NT-proBNP N-terminal pro-B-type natriuretic peptide, GDF15 growth differentiation factor 15, NYHA New York Heart Association, BMI body mass index, LVEF left ventricular ejection fraction, eGFR estimated glomerular filtration rate, NS non-significant

it is often unclear which threshold to use. This study shows for the first time that patients with a history of AF but who have SR at the time of measurement, have NT-proBNP levels that are much lower and more similar to those patients who have never had AF before, as compared to patients who have AF at time of measurement. Using a higher NP threshold for patients in SR but with a history of AF could result in inappropriately high screen failure rates in clinical trials. In these clinical HF trials, GDF-15 might be considered as an additional marker to distinguish severity of HF apart from AF.

Limitations

Limitations of this study include the post hoc design. There was a lack of information about the duration between the last episode of AF and screening and number/type of AF episodes the patient had experienced before. Asymptomatic patients with paroxysmal AF could have been missed and regarded as SR patients. Furthermore, no echocardiography data apart from LVEF was available. The NT-proBNP cutoff for inclusion in the BIOSTAT-CHF index cohort could have potentially led to higher inclusion rates of SR patients with more severe HF as compared with those with AF, and as compared with patients included in the validation cohort. However, even though this inclusion criterion differed, similar biomarker patterns were observed in both the index and validation cohort, which is a strength of the present study. Unfortunately, we were not able to further stratify the index and validation cohort in HFpEF and HFrEF, since this would have importantly limited the number of patients in the three rhythm groups within these three HF subtypes.

Conclusion

The plasma levels of NT-proBNP in HF patients were significantly influenced by the presence of AF at time of measurement, whereas the plasma levels of GDF-15 were independent of underlying AF. Therefore, GDF-15 might have additive value combined with NT-proBNP in the assessment of patients with HF and concomitant AF.

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Compliance with ethical standards

Conflict of interest SD.A. reports grants from Vifor and Abbott Vascular, and fees for consultancy from Vifor, Bayer, Boehringer Ingelheim, Brahms, Janssen, Novartis, Servier, Stealth Peptides, and ZS Pharma. K.Di. has received honoraria and/or research support from Medtronic,

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