

Pro-B-Type Natriuretic Peptide₁₋₁₀₈ Circulates in the General Community

Plasma Determinants and Detection of Left Ventricular Dysfunction

Fima Macheret, BS, MS,*† Guido Boerrigter, MD,* Paul McKie, MD,*
Lisa Costello-Boerrigter, MD, PhD,* Brian Lahr, MS,§ Denise Heublein, CLT,*
Sharon Sandberg, CLT,* Yasuhiro Ikeda, DVM, PhD,‡ Alessandro Cataliotti, MD, PhD,*
Kent Bailey, PhD,§ Richard Rodeheffer, MD, PhD,* John C. Burnett, JR, MD*

Rochester, Minnesota

Objectives

The purpose of this study was to investigate circulating pro-B-type natriuretic peptide (proBNP₁₋₁₀₈) in the general community and evaluate its ability to detect left ventricular (LV) dysfunction.

Background

The current concept for cardiac endocrine function is that, in response to cardiac stress, the heart secretes B-type natriuretic peptide (BNP₁₋₃₂) and amino-terminal pro-B-type natriuretic peptide (NT-proBNP₁₋₇₆) after intracardiac cleavage of their molecular precursor, proBNP₁₋₁₀₈. We hypothesized that proBNP₁₋₁₀₈ circulates in normal human subjects and that it is a useful biomarker for LV dysfunction.

Methods

Our population-based study included a cohort of 1,939 adults (age ≥45 years) from Olmsted County, Minnesota, with 672 participants defined as healthy. Subjects underwent in-depth clinical characterization, detailed echocardiography, and measurement of proBNP₁₋₁₀₈. Independent factors associated with proBNP₁₋₁₀₈ and test characteristics for the detection of LV dysfunction were determined.

Results

ProBNP₁₋₁₀₈ in normal humans was strongly influenced by sex, age, heart rate, and body mass index. The median concentration was 20 ng/l with a mean proBNP₁₋₁₀₈ to NT-proBNP₁₋₇₆ ratio of 0.366, which decreased with heart failure stage. ProBNP₁₋₁₀₈ was a sensitive (78.8%) and specific (86.1%) biomarker for detecting LV systolic dysfunction, which was comparable to BNP₁₋₃₂, but less than NT-proBNP₁₋₇₆, in several subsets of the population.

Conclusions

ProBNP₁₋₁₀₈ circulates in the majority of healthy humans in the general population and is a sensitive and specific biomarker for the detection of systolic dysfunction. The proBNP₁₋₁₀₈ to NT-proBNP₁₋₇₆ ratio may provide insights into altered proBNP₁₋₁₀₈ processing during heart failure progression. Thus, this highly specific assay for proBNP₁₋₁₀₈ provides important new insights into the biology of the BNP system. (J Am Coll Cardiol 2011;57:1386-95) © 2011 by the American College of Cardiology Foundation

Pro-B-type natriuretic peptide₁₋₁₀₈ (proBNP₁₋₁₀₈) is the 108-amino acid prohormone that is cleaved, by either corin or furin, to the 32-amino acid, biologically active brain

natriuretic peptide (BNP₁₋₃₂), also known as B-type natriuretic peptide, and to the 76-amino acid, biologically inactive N-terminal pro-B-type natriuretic peptide (NT-proBNP₁₋₇₆) (1). Brain natriuretic peptide augments sodium excretion, lowers blood pressure, suppresses the renin-angiotensin-aldosterone system, inhibits cardiomyocyte hypertrophy, induces angiogenesis, and retards activation of

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cardiac fibroblasts (2), whereas its prohormone has significantly reduced function (3). The elevation of BNP₁₋₃₂ and NT-proBNP₁₋₇₆ immunoreactivity in heart failure (HF) secondary to myocardial stretch, despite the lack of BNP functionality (the “BNP paradox”), has resulted in their widespread use as diagnostic and prognostic biomarkers (4–6).

From the *Cardiorenal Research Laboratory, Division of Cardiovascular Diseases, Departments of Medicine and Physiology, Mayo Clinic, Rochester, Minnesota; †College of Medicine, Mayo Clinic, Rochester, Minnesota; ‡Department of Molecular Medicine, College of Medicine, Mayo Clinic, Rochester, Minnesota; and the §Division of Biomedical Statistics and Informatics, Department of Health Sciences Research, College of Medicine, Mayo Clinic, Rochester, Minnesota. This study was funded by National Institutes of Health grant RO1 HL36634 and PO1 HL76611, Mayo Clinic Center for Clinical and Translational Research grant TL1RR024152, and a grant from Bio-Rad. Dr. Cataliotti was also supported by the Doris Duke Charitable Foundation Scientific Development grant CSDA 2006064 and by the M.I.U.R. Progetto Rientro dei Cervelli. All other authors have reported that they have no relationships to disclose.

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Most recently, studies with mass spectroscopy and Western blot analysis have identified that the major immunoreactive form of BNP in plasma of patients with HF is proBNP_{1–108}, rather than biologically active BNP_{1–32}, which can in part explain the BNP paradox (3,7–10). Although several studies have documented elevated proBNP_{1–108} levels in patients with HF (11–13), to date there are no studies that have investigated circulating proBNP_{1–108} levels in a large sample of normal, disease-free humans to define key independent determinants of plasma levels of this important prohormone. Furthermore, no studies have analyzed the potential of unprocessed BNP forms, such as proBNP_{1–108}, to improve detection of left ventricular (LV) dysfunction.

Our central hypothesis was that proBNP_{1–108} circulates in normal human subjects without cardiac or renal disease, is increased in the presence of LV dysfunction, is modulated by age and sex, and serves as a biomarker for the detection of LV dysfunction. This hypothesis is based on previous studies by Costello-Boerrigter et al. (14), Redfield et al. (15), and Wang et al. (16) that demonstrated that these factors are independently correlated with BNP_{1–32} and NT-proBNP_{1–76}. To test this hypothesis, we utilized a novel assay for plasma proBNP_{1–108} developed by Giuliani et al. (12) that is sensitive and specific for detecting proBNP_{1–108} and does not cross-react with mature BNP_{1–32} or NT-proBNP_{1–76}. Our study cohort was a National Institutes of Health-supported study, the PAVD (Prevalence of Asymptomatic Ventricular Dysfunction) study, which consisted of a community-based cohort of 1,939 persons, of whom 672 were considered clinically normal, in Olmsted County, Minnesota (17).

Methods

The Mayo institutional review board approved this study. **Study population.** Medical history and detailed 2-dimensional and color Doppler echocardiography were done on participating residents (n = 1,939; age ≥45 years) in Olmsted County, Minnesota, and were previously reported as the PAVD study (15). Subjects were divided by cardiovascular risk factors into a reference group of stage 0 HF subjects (no risk factors) who were normal healthy humans, and a group of subjects with stages A to C HF as defined by the 2009 HF guidelines (18,19). Persons with stage D HF were previously excluded from the study (5,15,17,20).

Assays and sample processing. Blood samples were drawn into ethylenediaminetetraacetic acid-treated tubes and chilled to 4°C, then centrifuged at 2,500 rpm for 10 min and stored at –80°C. Plasma proBNP_{1–108} was measured with the Bio-Rad assay (Bio-Rad, Hercules, California) on an automated analyzer not yet commercially available. This immunoassay was developed by Giuliani et al. (12) to be specific for proBNP_{1–108} as it uses antibodies directed against the hinge region that is present only in the intact proBNP_{1–108} molecule. The lower limit of detection for proBNP_{1–108} is 2 ng/l and the interassay and intra-assay

variabilities for proBNP_{1–108} are 10.3% and 11.6%, respectively. Plasma NT-proBNP_{1–76} was previously measured using the Elecsys electrochemiluminescence immunoassay on the Elecsys 2010 platform (Roche Diagnostics, Indianapolis, Indiana), which is a sandwich assay that at that time utilized 2 different polyclonal antibodies, 1 to amino acids 1–21, and 1 to amino acids 39–50. Samples were previously frozen and sent to Biosite Diagnostics (San Diego, California) for measurement of plasma BNP_{1–32} (14,15). The nonextracted 1-ml aliquots of frozen plasma were batch analyzed with the Biosite fluorescence immunoassay. The total within-day coefficient of variation (9.4% to 15.2%) and total coefficient of variation (10.1% to 16.2%) increased from low to high BNP.

Influence of clinical characteristics. Influences on proBNP_{1–108} were assessed with univariate and multivariable analysis for age, sex, body mass index (BMI), heart rate, left atrial (LA) volume index, LV mass index, LV end-diastolic volume dimension index, systolic and diastolic blood pressure, serum creatinine, and calculated glomerular filtration rate (by Cockcroft-Gault formula). All index values were calculated by dividing by each subject's body surface area.

Statistical methods. Descriptive statistics were used to describe the cohort of subjects: median and interquartile range for continuous data, and count and percentage for categorical data. Patient and clinical characteristic were compared between subjects in the various stages of HF using rank-order tests such as Spearman correlation and the Kruskal-Wallis method. To derive normal values for proBNP_{1–108}, the 5th, 50th, and 95th percentile values from the normal subgroup of subjects were used. Multivariable linear least-squares regression was then used to detect independent associations between proBNP_{1–108} levels and clinical cardiovascular risk factors with 2-way interactions considered. Because the variability of proBNP_{1–108} increased with its mean level, a natural log transformation of the proBNP_{1–108} value +1 (since 0 is in its domain, but natural log of 0 is undefined) was applied to satisfy regression modeling assumptions. Similar transformations were applied to the BNP_{1–32} and NT-proBNP_{1–76} biomarkers.

To optimize the number of subjects retained in the multivariable regression, all continuous parameters <90% observed were each categorized, including a category for unknown values. Using stepwise selection with entry and retention criteria of p < 0.10 and p < 0.05, respectively, a final model was derived for the total population and separately for the stage 0 HF (normal subjects) and stages A to C HF subgroups. Bootstrap resampling was used to evaluate the robustness of the clinical and echocardiographic parameters that were statis-

Abbreviations and Acronyms

AUC = area under the curve
BMI = body mass index
BNP = B-type natriuretic peptide
EF = ejection fraction
HF = heart failure
LA = left atrial
LV = left ventricular
NT-proBNP = N-terminal pro-B-type natriuretic peptide

tically significantly associated with natriuretic peptide levels. The frequency of each variable retained in the model using the pre-specified model selection criteria was computed as a percentage of all 1,000 bootstrap samples. Our modeling strategy required that the factors included in the final model each be retained in at least 60% of the bootstrap samples, thus reducing the chance of type 1 error (21,22).

The diagnostic utility for detecting LV dysfunction as either ejection fraction (EF) $\leq 40\%$, or EF $\leq 50\%$, and/or moderate to severe diastolic dysfunction was assessed for BNP_{1–32}, NT-proBNP_{1–76}, and proBNP_{1–108}. Diagnostic test characteristics of proBNP_{1–108} were evaluated using receiver-operating characteristic curves to determine areas under the curve (AUCs), which were then compared by the deLong method (23) to those for BNP_{1–32} and NT-proBNP_{1–76} in the total and age- and sex-stratified populations (14). The optimal discriminatory value for each assay was estimated by the point along the receiver-operating characteristic curve that provided the minimum Euclidean distance between the curve and a perfect assay with 100% sensitivity and specificity. The positive likelihood ratio (+LR = sensitivity/[1 – specificity]) and negative likelihood ratio (–LR = [1 – sensitivity]/specificity) were calculated for the optimal discriminatory values. Additional test characteristics estimated include the percentage of patients screened who would need echocardiography secondary to an abnormal proBNP_{1–108} test result (% need echo), the percent of echocardiograms that then would be negative (% –echo), and the percent of those with a reduced EF who would be missed by the proBNP_{1–108} cutoff (% missed), and the odds ratio of having EF $\leq 40\%$ or 50% and/or diastolic dysfunction for a high proBNP_{1–108} result. Statistical significance was accepted at $p < 0.05$ for all analyses.

Results

ProBNP_{1–108} immunoreactivity in the general community.

Plasma proBNP_{1–108} values were determined in all 1,939 subjects of the community-based cohort, of whom 672 were defined as the normal reference group (i.e., stage 0 HF/healthy normals) because they had no cardiovascular or renal disease by clinical examination and normal LV structure and function by echocardiography. The rest of the cohort was considered clinically abnormal, and further classified as having stage A, B, or C HF. Table 1 provides characteristics for the total study population and for subgroups according to stage of HF. Figure 1 displays proBNP_{1–108}, NT-proBNP_{1–76}, and BNP_{1–32} distributions by HF stage. As illustrated, proBNP_{1–108} was detected at the lowest levels in the reference group. By univariate assessment, proBNP_{1–108} levels increased significantly with advancing stage of HF, as did BNP_{1–32} and NT-proBNP_{1–76} levels. As shown in Figure 2, although higher in female subjects at younger ages, plasma proBNP_{1–108} levels increased with age more markedly in male subjects in the total cohort and in subjects

with HF. In normal subjects, however, higher levels of plasma proBNP_{1–108} were associated with increasing age as well as with female sex across the entire age range.

We sought to investigate whether these age and sex trends for proBNP_{1–108} held after adjustment for potential confounders. Table 2 reports the clinical characteristics that significantly influence proBNP_{1–108} levels within the total population, the reference group (stage 0 HF/healthy normals), and the abnormal group (stages A to C HF). In the multivariable analysis of the reference group, sex, age, heart rate, and BMI were independently associated with proBNP_{1–108} ($p < 0.001$ for age and sex, $p = 0.006$ for heart rate, $p = 0.004$ for BMI 25 to 30 kg/m², and $p = 0.021$ for BMI > 30 kg/m²). Echocardiographic measurements were not associated with proBNP_{1–108} in normal subjects, but LV mass index and LV dimension index were significantly associated with proBNP_{1–108} in the total population and in subjects with stages A, B, or C HF. Table 3 displays age- and sex-specific ranges for normal values of proBNP_{1–108}, noting that there are few subjects older than 75 years in the normal subgroup. Table 4 illustrates the proBNP_{1–108}/NT-proBNP_{1–76} ratio by stage of HF. The proBNP_{1–108}/NT-proBNP_{1–76} ratio decreased significantly from stage 0 to stages A to C.

Detection of LV dysfunction. In this community-based sample, 35 (2%) subjects had an EF $\leq 40\%$, 114 (6%) had an EF $\leq 50\%$, and 135 (8%) had moderate or severe diastolic dysfunction. For detection of LV systolic dysfunction, the AUCs for BNP_{1–32}, NT-proBNP_{1–76}, and proBNP_{1–108} are each summarized in Online Table 1 for the total population and for subsets stratified by age and sex. In general, the performance for detecting LV systolic dysfunction was comparable between proBNP_{1–108} and BNP_{1–32}, although proBNP_{1–108} was more diagnostic of an EF $\leq 50\%$ in the younger stratum of the total and male populations. Despite this, NT-proBNP_{1–76} was superior to proBNP_{1–108} for detecting EF $\leq 40\%$ and $\leq 50\%$ except in the younger subjects. Across the various population groups, each of these natriuretic peptides was consistently more diagnostic of LV dysfunction based on an EF cutoff of 40% as opposed to 50%.

Using a single group-specific cutpoint to optimize sensitivity and specificity in detecting reduced EF, the test characteristics of the proBNP_{1–108} assay were summarized for the total population and for age- and sex-stratified subgroups of the population (Online Table 2). For optimal discrimination of EF $> 40\%$ or $< 40\%$ in the total population, the cutpoint proBNP_{1–108} value was 65 ng/l, reflecting a sensitivity of 78.8% and a specificity of 86.1%. The best performance of this assay was in female and younger subgroups.

Similar cutpoint analyses were performed for detecting diastolic dysfunction and overall LV dysfunction, a composite of either systolic (EF $\leq 40\%$) or diastolic (moderate or severe based on echocardiography) dysfunction. For discriminating subjects with and without diastolic dysfunction, the AUCs were on average comparable between NT-

Table 1 Characteristics of the Study Population by Stage of HF

Variable*	Total Population (n = 1,939)	Stage 0 HF (n = 672)	Stage A HF (n = 788)	Stage B HF (n = 415)	Stage C HF (n = 64)
Females	1,003 (52%)	347 (52%)	391 (50%)	245 (59%)	20 (31%)
Age, yrs	61.8 (53.6–70.5)	54.5 (50.6–61.3)	64.9 (57.2–72.2)	67.9 (59.3–76.4)	73.8 (64.4–82.0)
Age categories, yrs					
45–54	560 (29%)	346 (51%)	146 (19%)	62 (15%)	6 (9%)
55–64	595 (31%)	222 (33%)	252 (32%)	110 (27%)	11 (17%)
65–74	500 (26%)	89 (13%)	264 (34%)	128 (31%)	19 (30%)
75+	284 (15%)	15 (2%)	126 (16%)	115 (28%)	28 (44%)
BMI, kg/m ²	27.7 (25.0–31.2)	27.1 (24.5–29.8)	28.1 (25.0–31.7)	28.1 (25.3–31.9)	28.0 (25.7–31.9)
Obesity, BMI >30 kg/m ²	623 (32%)	162 (24%)	287 (36%)	152 (37%)	22 (34%)
ProBNP _{1–108} , ng/l	20.0 (9.0–42.0)	14.0 (7.0–26.0)	20.0 (9.0–41.0)	37.0 (16.0–79.0)	150.0 (61.5–391.0)
NT-proBNP _{1–76} , ng/l	69.9 (28.3–147.5)	39.4 (18.0–82.0)	72.1 (31.3–140.7)	138.5 (61.0–332.2)	777.8 (234.2–1271.0)
BNP _{1–32} , ng/l	24.0 (9.5–56.3)	15.3 (6.4–29.8)	24.4 (9.7–53.9)	50.5 (20.3–106.2)	138.5 (80.4–313.6)
Calculated GFR	75.0 (60.4–92.6)	81.2 (66.8–97.0)	72.9 (58.9–91.3)	69.3 (54.6–90.5)	62.4 (44.8–78.5)
Estrogen†	460/961 (48%)	160/325 (49%)	194/381 (51%)	101/235 (43%)	5/20 (25%)
Diabetes mellitus	145 (7%)	0 (0%)	85 (11%)	47 (11%)	13 (20%)
Past or current atrial fibrillation	94 (5%)	0 (0%)	34 (4%)	31 (7%)	29 (45%)
Coronary artery disease	233 (12%)	0 (0%)	113 (14%)	78 (19%)	42 (66%)
Past heart failure diagnosis	49 (3%)	0 (0%)	0 (0%)	0 (0%)	49 (77%)
Hypertension	543 (28%)	0 (0%)	340 (43%)	168 (40%)	35 (55%)
Systolic blood pressure, mm Hg	131.0 (116.0–146.0)	122.0 (111.0–134.0)	135.0 (121.0–149.0)	138.0 (123.0–155.0)	134.0 (119.0–154.0)
Diastolic blood pressure, mm Hg	73.0 (67.0–80.0)	72.0 (66.0–78.0)	75.0 (68.0–82.0)	74.0 (67.0–80.0)	70.0 (66.0–82.5)
LV dimension index	2.61 (2.40–2.82)	2.59 (2.39–2.77)	2.55 (2.36–2.74)	2.77 (2.54–2.97)	3.04 (2.73–3.29)
LV mass index	93.7 (81.9–108.8)	88.2 (77.2–97.0)	92.2 (81.8–104.2)	113.1 (97.7–128.0)	137.2 (116.5–163.3)
LA volume index	23.3 (19.4–28.1)	20.9 (18.0–24.6)	22.7 (19.4–26.2)	33.0 (25.2–36.9)	35.8 (29.0–51.0)
Beta-blocker	284 (16%)	0 (0%)	162 (22%)	99 (25%)	23 (36%)
ACEI/ARB	204 (11%)	0 (0%)	115 (15%)	57 (14%)	32 (50%)
Diuretic	330 (18%)	0 (0%)	193 (26%)	98 (25%)	39 (61%)
EF <40%	35 (2%)	0 (0%)	0 (0%)	0 (0%)	35 (55%)
EF <50%	114 (6%)	0 (0%)	0 (0%)	71 (17%)	43 (67%)
Moderate or severe DD	135/1,709 (8%)	0/672 (0%)	52/668 (8%)	63/330 (19%)	20/39 (51%)

Values are n (%) or median (interquartile range). *The top block of variables were each tested for an association with advancing stage of heart failure (HF) using a Spearman correlation rank-order test or a Cochran-Armitage trend test as appropriate for continuous and categorical variables (each had a highly significant association [$p < 0.001$], except for sex [$p = 0.796$] and female estrogen use [$p = 0.047$]; the lower block of variables was used in defining the subgroups of HF, so no statistical tests were conducted. †Percent of estrogen use based only on female subgroup.

ACEI = angiotensin-converting enzyme inhibitor; ARB = angiotensin-receptor blocker; BMI = body mass index; BNP = B-type natriuretic peptide; DD = diastolic dysfunction; EF = ejection fraction; GFR = glomerular filtration rate; LA = left atrial; LV = left ventricular; NT-proBNP = N terminal pro-B-type natriuretic peptide; proBNP = pro-B-type natriuretic peptide.

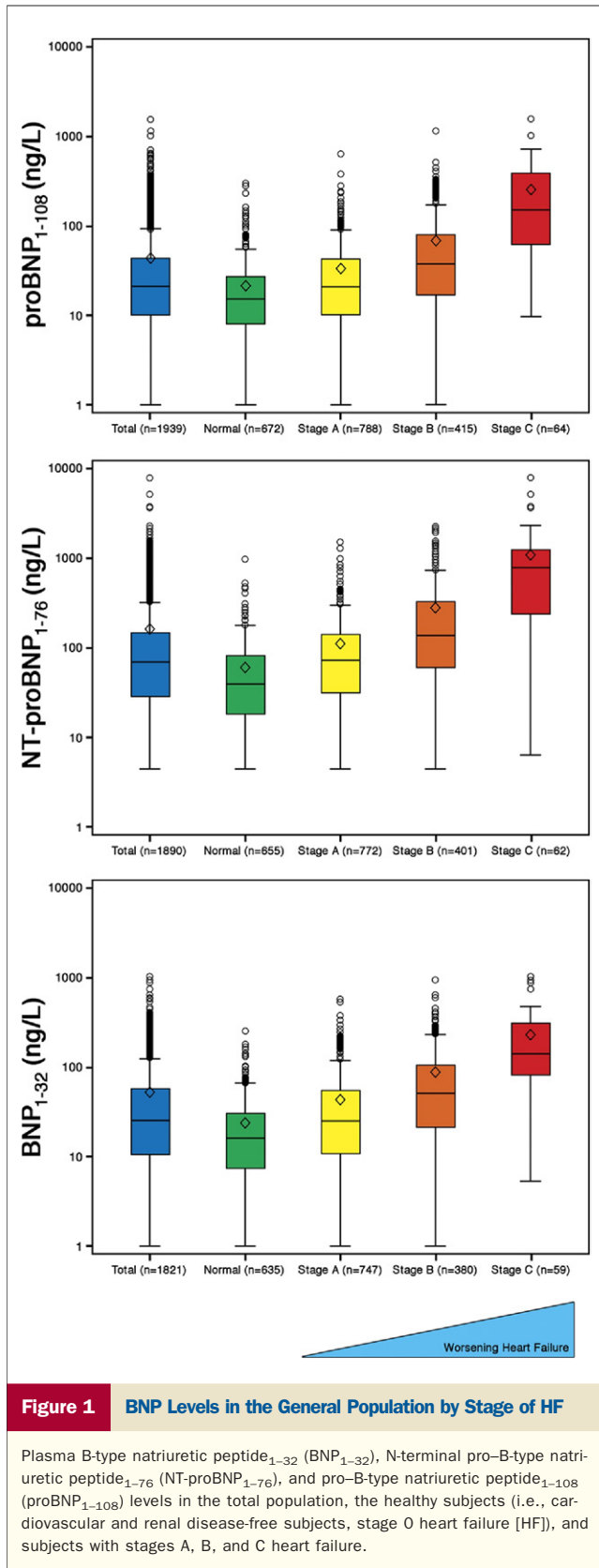
proBNP_{1–76} and BNP_{1–32}, and significantly greater than those of proBNP_{1–108} (Table 5). The optimal cutpoint for detection of diastolic dysfunction by proBNP_{1–108} was 25.1 ng/l, with a resulting sensitivity of 66.1% and a specificity of 63.1% in the total population. Finally, the optimal proBNP_{1–108} cutpoint for detection of overall LV dysfunction as a composite of either systolic or diastolic dysfunction was 39 ng/l, with a sensitivity of 56.8% and a specificity of 74.5% in the total population (AUCs not displayed for composite LV dysfunction).

Given their strong influence on proBNP_{1–108} levels, age- and sex-specific cutpoints were obtained for detecting a reduced ejection, with their corresponding test characteristics compared to those based on 1 unadjusted proBNP_{1–108} cutpoint (Online Tables 3 and 4). In the total population, the use of age- and sex-adjusted (vs. unadjusted) proBNP_{1–108} cutpoints increased the relative risk estimate of having an EF ≤40% by a factor of 2 (95% confidence interval of odds ratio: 23 to 46).

Discussion

ProBNP_{1–108} is a circulating prohormone. The major finding in our study is that proBNP_{1–108} circulates in the majority of normal, disease-free humans, which changes our understanding of the secretion of proBNP_{1–108} and its processing to mature, biologically active BNP_{1–32}. Although prior studies have shown that proBNP_{1–108} exists in circulation, our translational study is the first to our knowledge to confirm at the population level the relative predominance of this prohormone compared to multiple immunoreactive BNP forms that circulate and which are nonspecifically detected by conventional BNP_{1–32} and NT-proBNP_{1–76} assays (7,9,10,24).

To date, studies have focused on circulating proBNP_{1–108} in humans with advanced HF (8,10–13). With BNP_{1–32} reported to be at low levels or undetectable in human HF despite high immunoreactivity using conventional assays, the concept has been proposed that the presence of proBNP_{1–108} in plasma reflects a defect in proBNP_{1–108}



processing in the heart with spillover of unprocessed and nonbiologically active proBNP₁₋₁₀₈ into the plasma (3,6,9). Our study changes that paradigm with the use of this newly developed and specific assay for proBNP₁₋₁₀₈ that does not cross-react with NT-proBNP₁₋₇₆ or BNP₁₋₃₂. Specifically, our data are consistent with the concept of the physiological release of proBNP₁₋₁₀₈ from the normal heart, implying peripheral processing to mature BNP₁₋₃₂ either in plasma and/or at target organs. Indeed, we have recently documented the presence of the proBNP₁₋₁₀₈ processing convertase corin in normal human plasma and the in vitro processing of proBNP₁₋₁₀₈ to BNP₁₋₃₂ in normal human plasma (25) together with a step-up in the proBNP₁₋₁₀₈ gradient across the normal human heart (26). With

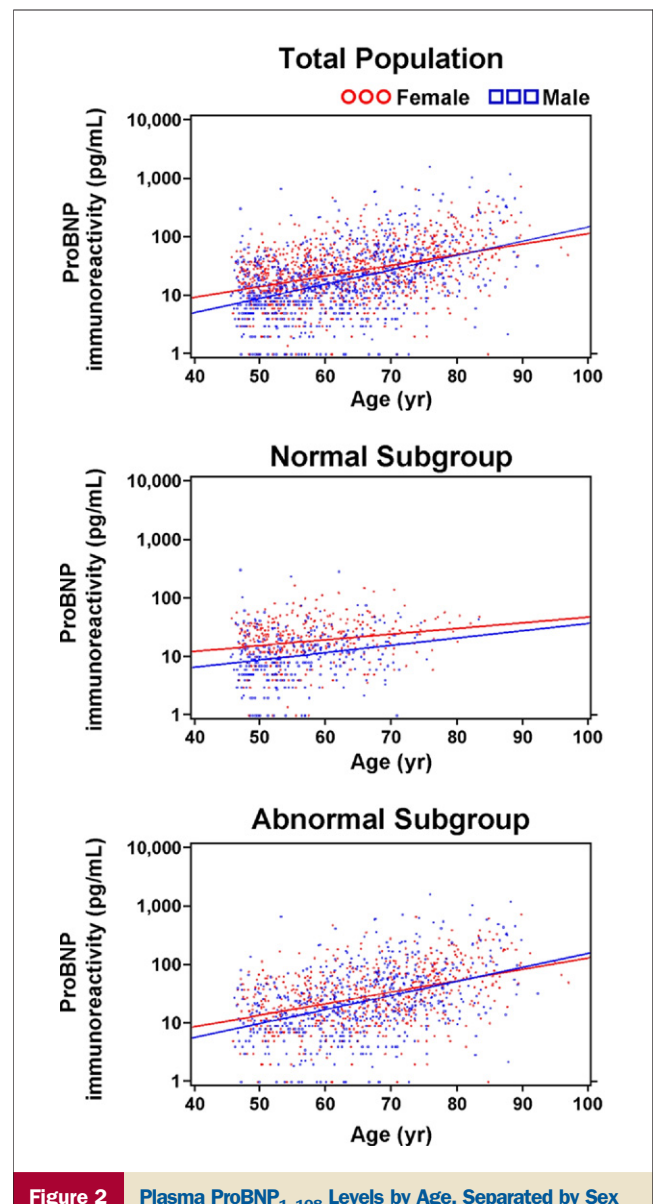


Table 2 Parameters That Significantly Contribute to ProBNP_{1–108}* in Multivariable Analysis

Population	Parameters Included in Model	Regression Coefficient	SE	p Value
Total population†	Age‡			
	Males	0.431	0.035	<0.001
	Females	0.308	0.032	<0.001
	BMI ≤25 kg/m ² (reference)	—	—	—
	25–30 kg/m ²	–0.193	0.060	0.001
	>30 kg/m ²	–0.184	0.067	0.006
	LVMi high (>134 males, >110 females)	0.203	0.085	0.017
	LVDI high (>2.6)	0.151	0.059	0.010
Normal subgroup (stage 0 HF)	Creatinine	0.297	0.129	0.021
	Age‡	0.250	0.046	<0.001
	Sex, female	0.485	0.073	<0.001
	Heart rate	–0.010	0.004	0.006
	BMI ≤25 kg/m ² (reference)	—	—	—
25–30 kg/m ²	–0.244	0.085	0.004	
>30 kg/m ²	–0.227	0.099	0.021	
Abnormal subgroup (stages A–C HF)	Age‡	0.406	0.031	<0.001
	Sex, female	0.162	0.063	0.010
	BMI ≤25 kg/m ² (reference)	—	—	—
	25–30 kg/m ²	–0.153	0.079	0.054
	>30 kg/m ²	–0.179	0.085	0.036
	LVMi high (>134 males, >110 females)	0.225	0.092	0.014
LVDI high (>2.6)	0.146	0.077	0.059	

*Dependent variable in all linear regression models above was log-transformed proBNP value+1. †For left ventricular mass index (LVMi) and left ventricular dimension index (LVDI) covariates, each was fit as categorical with categories abnormal/high (effect shown), normal (reference level), and unknown (effect not shown). ‡Age, expressed per 10-year change.

Abbreviations as in Table 1.

proBNP_{1–108} now a circulating hormone, it is logical that there now is a need for such an assay to better assess direct secretion of this cardiac hormone from the heart and its peripheral processing from a biological perspective as well as to assess its potential diagnostic significance across a wide spectrum of human cardiovascular disease.

Importantly, we have documented in a large population both free of HF and in stages A to C what Dries et al. (11) has labeled the “natriuretic peptide processing efficiency,” and which has previously been investigated in the PENN HF study, which was mostly in patients with advanced systolic HF. Our results show that progressing from stage 0 to stages A to C is associated with increasing processing efficiency due to either decreased secretion of proBNP_{1–108} or an increase in peripheral processing to NT-proBNP_{1–76} as demonstrated by the decrease in the proBNP_{1–108}/NT-

proBNP_{1–76} ratio. It appears that NT-proBNP_{1–76} is increasing faster than proBNP_{1–108} in stages A to C. We speculate as one moves into stage D, secretion outpaces processing and efficiency decreases. Because of our population, we are unable to prove whether HF worsens processing efficiency or if decreasing processing efficiency pre-dates advanced HF, as we had no stage D subjects. To show this, an important future use of the proBNP_{1–108} assay, using step-up studies in subjects with varying degrees of HF and at different time points, will be needed to determine exactly what proportion of BNP forms the heart secretes in physiological and pathophysiological states, and where and by what mechanism precisely extracardiac processing occurs. We speculate that local processing of proBNP_{1–108} may be an efficacious way of promoting renal function without causing systemic actions such as hypotension.

Table 3 Age- and Sex-Specific Median Values and Ranges for Plasma ProBNP_{1–108} in Normal Subjects

	Age, yrs			
	45–54	55–64	65–74	≥75
Women				
Median (5th–95th percentiles), ng/l	17 (3–48)	21 (3–81)	23 (4–128)	37 (5–56)
n	168	117	49	13
Men				
Median (5th–95th percentiles), ng/l	7 (1–40)	12 (2–44)	16 (2–53)	25 (8–42)
n	178	105	40	2

ProBNP = pro-B-type natriuretic peptide.

Table 4 ProBNP_{1–108} to NT-proBNP_{1–76} Ratio by Stage of Heart Failure

Ratio	Stage 0	Stage A	Stage B	Stage C	p Value
ProBNP _{1–108} /NT-proBNP _{1–76}	0.366	0.300	0.271	0.301	<0.001
	(0.209, 0.612)	(0.178, 0.504)	(0.147, 0.444)	(0.140, 0.416)	

Median (quartile 1, quartile 3); both tested for an association with advancing stage of heart failure using a Spearman correlation rank-order test. Abbreviations as in Table 1.

Association of clinical and echocardiographic parameters on proBNP_{1–108} levels. We present the first analysis of the factors independently associated with circulating proBNP_{1–108} in subjects without cardiovascular and renal disease and then compare those factors to those of BNP_{1–32} and NT-proBNP_{1–76} in the same population. To our knowledge, this is the first study of natriuretic peptides that utilized the internal validation technique of bootstrap resampling to ensure the robustness of the multivariable modeling. We found that age and sex were the principal determinants of proBNP_{1–108} levels, with higher levels associated with increasing age and female sex, but we also observed that this association increased at a higher rate with age in males in the total population. Despite significant associations, the effects of age and sex on proBNP_{1–108} in our study were not as strong as those on NT-proBNP_{1–76}, suggesting a difference between the effects of age and sex on cardiac proBNP_{1–108} production and downstream processing and renal clearance (13,14,27,28). These important findings underscore the need for considering age and sex when developing ranges for normals; they also emphasize that production and processing are distinct processes with unique determining characteristics. Of relevance is the recent report by Ichiki et al. (25) that corin levels in the plasma are also determined by sex, with males possessing higher levels than females.

Although the mechanisms by which age contributes to an increase in proBNP_{1–108} levels are not clearly understood, we demonstrate that the aging heart produces more proBNP_{1–108}, even with normal myocardial function and structure in the general population. Expression of the BNP gene has been shown to be up-regulated with aging (29), but some studies have speculated that circulating BNP_{1–32} immunoreactivity rises because of impaired renal function

with aging, including reduced natriuretic peptide receptor function (30,31). Thus, the increase in proBNP_{1–108} with aging may be due to 1 or more mechanisms: increased cardiac production and release or impaired downstream processing or clearance.

Previous studies have shown a relationship between sex and BNP_{1–32} and NT-proBNP_{1–76}, and some have noted an association between estrogen use (hormone replacement therapy) and increased BNP_{1–32} levels (14,15). We did not find an independent association between estrogen use and proBNP_{1–108} in our study, whereas a significant association was found for NT-proBNP_{1–76} in the Costello-Boerrigter et al. (14) study, suggesting a differential effect by androgens and estrogens on production and processing. Studies of sex differences in natriuretic peptide production and processing may lead to better explanation of the predominance of men with HF at younger ages than women.

Another important finding in our study was that cardiac structure and function parameters, LV mass index and LV dimension index, were independently associated with increased proBNP_{1–108} in stage A, B, and C HF subgroups, suggesting that decreasing myocardial function and worsening fibrosis, dilation, and hypertrophy may contribute to increasing proBNP_{1–108} by an alteration in production or processing capability in persons with cardiovascular and renal disease.

Previously, NT-proBNP_{1–76} and BNP_{1–32} were found to be independently associated with LA volume index, but in our study, proBNP_{1–108} was not. It was believed that LA volume index and LV dimension index had effects on the variability of BNP_{1–32} because both atrial and ventricular myocardium are responsible for BNP_{1–32} production (5,32–34). Importantly, our results suggest a stronger role for ventric-

Table 5 Detection of Moderate or Severe Diastolic Dysfunction by ProBNP_{1–108}, NT-proBNP_{1–76}, and BNP_{1–32}

Population/Subgroup	n	AUC (95% CI)			p Value of Overall Difference
		ProBNP _{1–108}	NT-proBNP _{1–76}	BNP _{1–32} (Bio-site)	
Total (n = 1,558)	118	0.71 (0.66, 0.76)	0.77 (0.72, 0.82)*	0.78 (0.73, 0.82)*	0.028
Age ≥65 yrs (n = 594)	77	0.65 (0.58, 0.72)	0.73 (0.67, 0.79)*	0.76 (0.70, 0.82)*	0.006
Age <65 yrs (n = 964)	41	0.71 (0.63, 0.79)	0.73 (0.63, 0.83)	0.72 (0.63, 0.81)	0.871
All men (n = 742)	54	0.77 (0.71, 0.83)	0.78 (0.71, 0.86)	0.78 (0.72, 0.85)	0.929
All women (n = 816)	64	0.66 (0.58, 0.74)	0.76 (0.70, 0.83)*	0.79 (0.72, 0.85)*	0.008
Men, age ≥65 yrs (n = 265)	33	0.71 (0.61, 0.80)	0.76 (0.67, 0.86)	0.78 (0.70, 0.87)	0.228
Women, age ≥65 yrs (n = 329)	44	0.61 (0.50, 0.72)	0.70 (0.61, 0.80)	0.75 (0.67, 0.83)*	0.022
Men, age <65 yrs (n = 477)	21	0.77 (0.67, 0.87)	0.72 (0.57, 0.86)	0.71 (0.59, 0.84)	0.644
Women, age <65 yrs (n = 487)	20	0.66 (0.53, 0.80)	0.76 (0.64, 0.89)	0.76 (0.62, 0.89)	0.345

*Area under the curve (AUC) of biomarker is significantly different from that of proBNP_{1–108}. CI = confidence interval; other abbreviations as in Table 1.

ular myocardium than atrial myocardium in secretion of unprocessed proBNP_{1–108} in diseased subjects compared with normal subjects, on the basis of a lack of significant association between LV echocardiographic characteristics and proBNP_{1–108} in subjects with stage 0 HF (healthy normals) and the presence of significant associations between LV dimension index and LV mass index and proBNP_{1–108} in subjects with cardiovascular disease. Atrial myocardium may play a part in secretion of mature (BNP_{1–32}) or may be involved in processing.

We did not find a statistically significant correlation between glomerular filtration rate and proBNP_{1–108} levels in our multivariable analysis of the normal population. This finding is similar to previous studies with NT-proBNP_{1–76} in our study population (14). However, several groups have reported that BNP_{1–32} and NT-proBNP_{1–76} levels are inversely correlated with renal function in abnormal patients, such as in those with varying degrees of HF or renal disease (35–37). It was further shown that cardiovascular disease was required for BNP_{1–32} elevation in patients on dialysis for end-stage renal disease (38). Given that our study had so few subjects with the severity of disease that was examined in those studies, it was unlikely that we would have found an association. Another consideration is that proBNP_{1–108}, NT-proBNP_{1–76}, and BNP_{1–32} may all have different mechanisms for renal clearance (39–41), which could explain the variability in the associations among these 3 forms of BNP and renal function, especially in the stages A to C HF group.

Finally, we note that the decrease in proBNP_{1–108} with obesity may support the overall concept that decreased circulating BNP_{1–32} levels in obesity may indeed be secondary to reduced BNP_{1–32} production by the heart in this clinical setting (42). This observation supports the need for better understanding the impact of obesity on the heart, which may influence BNP production, and, here, use of proBNP_{1–108} measurements may have an important role beyond HF alone.

In our analysis of clinical factors, because of lack of data, we were unable to correlate proBNP_{1–108} secretion with functional status as measured by New York Heart Association functional class of HF.

Detection of LV dysfunction. We are the first to report that the Bio-Rad proBNP_{1–108} assay detected LV systolic dysfunction in the general community with high sensitivity and specificity and was comparable to BNP_{1–32} in discriminative performance (14). We also show the efficacy of an unprocessed natriuretic peptide for detection of diastolic dysfunction in a large population. Importantly, because of a lack of cross-reactivity, we are able to emphasize that this novel assay is a true reflection of physiological and pathological secretion of proBNP_{1–108}, and, therefore, of the endocrine function of the heart. A specific proBNP_{1–108} assay will be of tremendous value for future studies evaluating the peripheral processing of proBNP_{1–108}. Regarding its use as a novel biomarker for HF, it is also important to

state that proBNP_{1–108} is a more robust biomarker for the detection of systolic versus diastolic dysfunction, and that our findings will have to be replicated in other populations and in other scenarios.

Based on statistically significant differences in their AUCs, the NT-proBNP_{1–76} assay performed better than the proBNP_{1–108} and BNP_{1–32} assays in several subgroups for the detection of reduced EF, whereas the proBNP_{1–108} assay was comparable with BNP_{1–32} in most others. These results illustrate that all of these assays hold some diagnostic value for the detection of LV systolic dysfunction in the general population, but individual differences in their biologic meaning must be taken into account in the clinical setting. We caution against over-interpretation of their differences because of the small number of subjects in some of the subgroups, especially females. In fact, the lack of a large number of diseased persons in the general population is an important limitation of the cohort study design.

Our findings illustrate the heterogeneity of factors in cardiac disease that may influence proBNP_{1–108} secretion and processing. We noted that age- and sex-adjusted cutpoints generally improved test characteristics for detecting EF ≤ 40%. Furthermore, we noted that processed forms of BNP, including BNP_{1–32} and NT-proBNP_{1–76} were for the most part better at detection of systolic HF, implying that processes that adversely affect ventricular filling may have a differential effect on processing compared to secretion. The heterogeneity is also evident from the fact that when proBNP_{1–108} is used to detect composite, namely, systolic and/or diastolic dysfunction, the sensitivity and specificity both decrease relative to systolic dysfunction alone.

A limitation of our study was that we had no information regarding right ventricular function. Therefore, we had no way to assess the diagnostic accuracy of proBNP_{1–108} for right ventricular dysfunction.

Physiologic, pathophysiologic, and diagnostic significance of proBNP_{1–108}. In addition to its value as a novel diagnostic tool for HF, the Bio-Rad proBNP_{1–108} assay reveals both the physiologic and pathophysiologic importance of proBNP_{1–108} in human plasma. It confirms and extends previous studies by Lam et al. (24) that utilized an assay for proBNP_{3–108}, which reported its presence in the general population. With the proBNP_{1–108} assay, which has been documented not to cross-react to BNP_{1–32} or NT-proBNP_{1–76} we can now be confident that the heart releases this prohormone of BNP in normal humans (12). Indeed, the presence of proBNP_{1–108} in most normal humans in this large population study and the recent report that proBNP_{1–108} can be processed to mature BNP_{1–32} in human plasma clearly change our understanding of the heart as an endocrine organ and the BNP system (25). These observations strongly suggest that in part proBNP_{1–108} is an important carrier protein, which delivers the mature peptide to the plasma or tissues for local processing and conversion. From a biological perspective, measurement of circulating proBNP_{1–108} may be important

from the perspective of fully understanding the intactness of proBNP_{1–108} secretion from the heart and its processing in the circulation.

In HF, the BNP paradox is the disparity between diagnostically elevated BNP_{1–32} immunoreactivity in patients with HF and a lack of BNP-derived vasodilatory and cardiorenal protective effects. We can now state that the elevation of proBNP_{1–108}, which has reduced biologic activity and cross-reacts with conventional BNP assays, in patients with HF can account for at least some of this phenomenon (3,8). Yet, in stages A, B, and C HF, there may be increased processing efficiency based upon the decrease in the proBNP_{1–108}/NT-proBNP_{1–76} ratio. Perhaps the increased secretion of proBNP_{1–108} in stages A to C, and its efficient conversion, represents successful compensation to maintain a state of compensated HF. We would predict that this efficiency is impaired in advanced HF, as supported by studies by Dries et al. (11). The exact reason for this possible proBNP_{1–108} processing deficiency is unknown, but there may be an enzymatic deficiency, whether due to saturation or constitutive dysfunction, that underlies the progression of HF. Such a deficiency may also represent a therapeutic target and opportunity. Use of either corin or a corin-like drug could have potential as a therapy to delay disease progression and warrants investigation, especially in experimental models of progressive HF. In addition to aberrant upstream processing, there may be downstream receptor insensitivity or decreased renal clearance, as other authors have speculated. These may be similar pathophysiological mechanisms to those that cause decreasing BNP_{1–32} activation and processing in early stages of hypertension as well (43). It should also be noted, however, as indicated in the preceding text, that decreased overall proBNP_{1–108} processing may serve to achieve high local levels of BNP_{1–32} activity in areas with convertase activity while at the same time avoiding systemic BNP_{1–32} activity, which could result, for example, in hypotension.

Conclusions

ProBNP_{1–108} circulates in the majority of normal persons in the general population, supporting the conclusion that proBNP_{1–108} is normally released from the heart and may serve a role as a carrier protein to deliver the biologically active peptide into the circulation or to target tissues for processing. Importantly, age and sex were the major factors influencing proBNP_{1–108} in the general population. In addition, proBNP_{1–108} was a sensitive and specific biomarker for the detection of systolic dysfunction and less so for diastolic dysfunction equal to that of BNP_{1–32}, but in general, less so than NT-proBNP_{1–76}. Importantly, the ratio of proBNP_{1–108}/NT-proBNP_{1–76} provides insight into possible changes in proBNP_{1–108} processing during the progression of HF. Thus, this highly specific assay for proBNP_{1–108} provides important new insights into the

biology of the BNP system as well as potential diagnostic applications for this important new technology.

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Reprint requests and correspondence: Mr. Fima Macheret, Cardiorenal Research Laboratory, Guggenheim 915, Mayo Clinic and Foundation, 200 First Street S.W., Rochester, Minnesota 55905. E-mail: macheret.fima@mayo.edu.

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Key Words: biomarker ■ BNP ■ heart failure ■ natriuretic peptide ■ NT-proBNP ■ proBNP.

 **APPENDIX**

For supplementary Tables 1 through 4, please see the online version of this article.