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# New Biomarkers in the Prognostic Assessment of Acute Heart Failure with Reduced Ejection Fraction: Beyond Natriuretic Peptides

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Keywords: Systolic heart failure; biomarkers; prognosis; GDF 15; sST2



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Article

# New Biomarkers in the Prognostic Assessment of Acute Heart Failure with Reduced Ejection Fraction: Beyond Natriuretic Peptides

Marcelino Cortés 1,2,\*, Jairo Lumpuy-Castillo 3,4, Camila Sofía García-Talavera 5, María Belén Arroyo Rivera 6, Lara de Miguel 1, Antonio José Bollas 1, JoseMaria Romero-Otero 1, Jose Antonio Esteban Chapel 1, Mikel Taibo-Urquía 1,2, Ana María Pello 1,2, María Luisa González-Casaus 7, Ignacio Mahillo 8, Oscar Lorenzo 3,4 and José Tuñón 1,9,10

- <sup>1</sup> Cardiology Department, Fundación Jiménez Díaz University Hospital, 28040, Madrid, Spain
- <sup>2</sup> Faculty of Medicine and Biomedicine, Universidad Alfonso X el Sabio (UAX), 28691, Madrid, Spain
- <sup>3</sup> Laboratoryof Diabetes and Vascular Pathology, IIS-Fundación Jiménez Díaz, Universidad Autónoma, 28040 Madrid, Spain
- <sup>4</sup> Biomedical Research Network on Diabetes and Associated Metabolic Disorders (CIBERDEM), Carlos III National Health Institute, 28029 Madrid, Spain
- <sup>5</sup> Cardiology Department, ComplejoHospitalario San Millán San Pedro, 26004, Logroño (La Rioja), Spain
- 6 Cardiology Department, HM Montepríncipe University Hospital, 28660. Madrid, Spain
- Department of Laboratory Medicine, La Paz University Hospital, 28046 Madrid, Spain
- 8 Laboratory of Epidemiology, IIS-Fundación Jiménez Díaz, Madrid, Spain
- 9 Department of Medicine, Faculty of Medicine, Universidad Autónoma de Madrid, 28049 Madrid, Spain
- 10 CIBERCV, 28029 Madrid, Spain
- \* Correspondence: mcortesg@quironsalud.es

Abstract: Natriuretic peptides are established biomarkers related to the prognosis of heart failure. New biomarkers have emerged in the area of cardiovascular disease. The prognostic value of these biomarkers in heart failure with reduced LVEF (HFrEF) is not well established. We conducted a prospective, single-centre study, including consecutively (July 2019 to March 2023) 104 patients admitted with a diagnosis of acute HFrEF decompensation. Median follow-up was 23.5 months, during which 20 deaths (19.4%) and 21 readmissions for heart failure (20.2%) were recorded. Plasma biomarkers such as NT-proBNP, GDF-15, sST2, uPAR, and FGF-23 were associated with an increased risk of all-cause mortality. However, a Cox regressionanalysis showed that the strongest predictors of mortality were estimated glomerular filtration rate (HR 0.96 [0.93-0.98]), GDF-15 (HR 1.3 [1.16-1.45]), and sST2 (HR 1.2 [1.11-1.35]). The strongest predictive model was formed by the combination of glomerular filtration rate and sST2 (C-index 0.758). In conclusion, in patients with acute decompensated HFrEF, GDF-15 and sST2 showed the highest predictive power for all-cause mortality, superior to other established biomarkers such as natriuretic peptides. GDF-15 and sST2 may provide additional prognostic information to improve the prognostic assessment.

**Keywords:** Systolic heart failure; biomarkers; prognosis; GDF 15; sST2

### 1. Introduction

Heart failure (HF) remains a prevalent and relevant health problem today. It is estimated that approximately 1-2% of the adult population suffers from it, reaching a prevalence of more than 10% in elderly patients [1,2]. Despite major advances in the treatment and management of these patients in recent years, the mortality and morbidity associated with HF remains high [3]. Several markers and prognostic models have been studied over the last decades in order to predict which patients are

at increased risk of events [4]. Among these risk markers, biomarkers, elements detectable in analytical samples, stand out. Their prognostic and diagnostic role has been analysed in the cardiovascular field, and also specifically in the field of HF [5]. Markers have been described at the neurohormonal level, inflammatory mediators, cell damage, etc., with natriuretic peptides standing out in particular. They have been fully implemented in clinical practice, playing a prognostic role, guiding treatment, or even in the very definition of HF [6]. In recent years, new biomarkers(related to inflammation, oxidative stress, tissue damage, renal function, etc.) have been sought to provide new advances in the management of patients with HF. To date, these new biomarkers have not been successfully used in routine clinical practice [7]. However, some of them such as soluble Suppression of Tumorigenicity 2 (sST2), Growth Differentiation Factor-15 (GDF-15), soluble urokinase Plasminogen Activator Receptor (suPAR), Fatty Acid Binding Protein 4 (FABP4), or mineral metabolism (MM) biomarkers (Fibroblast Growth Factor 23 (FGF23), klotho, phosphorus (P), parathyroid hormone (PTH), or 1-25-dihydroxyvitamin D (calcidiol) have shown promising results in relation to the diagnosis and prognosis of HF.

The aim of our study was to analyse the prognostic role of these new biomarkers in HF with reduced ejection fraction (HFrEF), in the setting of discharge after admission for acute heart failure, assessing and comparing the prognostic power of these biomarkers and their associations, as well as their added value to natriuretic peptides.

#### 2. Results

#### 2.1. Baseline characteristics of patients

We included 104 patients in our study (figure 1). The median age of our population was 66.7 years, with a majority of male patients (78.8%). The percentage of patients with comorbidities was relatively high. Thus, 29.8% had chronic lung disease (COPD, asthma, OSA), 31.7% had chronic kidney disease, 10.6% had a history of stroke, and30.8% of patients were in atrial fibrillation at inclusion. The percentage of diabetics in our population reached almost 50%, with more than 66% hypertensive. In 31% of the study population, the main underlying cause of LV systolic dysfunction was ischaemic heart disease, with 27.9% of patients having a history of previous STEMI. After hospital discharge, patients were followed up in the HFU according to the study protocol, achieving treatment rates with BB greater than 90%, ARBS-ACEIS-ARNI 87%, MRAs 74%, and SGLT2i 72.1%.

Following the described methodology, we analysed plasma samples obtained fromour study population at admission. Table 1 shows the results of the main biochemical blood parameters (renal function, iron profile, haemogram...) in our population. It also shows the results of the wide range of biomarkers determined in our study: the most classical ones (CK-MB, NT-proBNP, TnI), as well as a wide representation of new biomarkers (mineral metabolism biomarkers, GDF-15, sST2, suPAR, etc.).

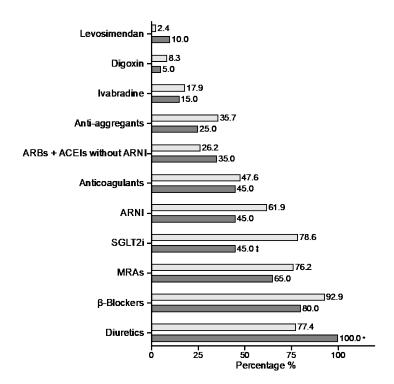


Figure 1. Baseline characteristics: clinical and treatment. Comparison according to all-cause mortality.

		All-cause death		
	Total	No	Yes	p-Value
	(n=104)	(n=84)	(n=20)	
Anthropometric parameters				
Age (years)	66.7 (18.3)	65.5 (16.1)	76.6 (13.5)	0.009
Male [n (%)]	82 (78.8)	66 (78.6)	16 (80)	0.888
Obesity [n (%)]	39 (37.5)	34 (40.5)	5 (25)	0.199
Risk factors and comorbidities				
Stroke [n (%)]	11 (10.6)	9 (10.7)	2 (10)	0.926
Peripheral vascular disease [n (%)]	9 (8.7)	7 (8.3)	2 (10)	0.683
CPD [n (%)]	31 (29.8)	23 (27.4)	8 (40)	0.268
CKD [n (%)]	33 (31.7)	22 (26.3)	11 (55)	0.013
Cancer [n (%)]	15 (14.4)	9 (10.7)	6 (30)	0.038
STEMI [n (%)]	29 (27.9)	24 (28.6)	5 (25)	0.749
LVEF (%)	20 (15)	20 (15)	20 (10)	0.936
Atrial fibrillation [n (%)]	32 (30.8)	26 (31.1)	6 (30)	0.934
NYHA III-IV [n (%)]	13 (12.5)	3 (3.6)	10 (50)	< 0.001
HF [n (%)]	46 (44.2)	31 (36.9)	15 (75)	0.002
Prior coronary revasc. [n (%)]	21 (20.2)	18 (21.4)	3 (15)	0.520
Smoking [n (%)]	37 (35.6)	30 (35.7)	7 (35)	0.952
Diabetes [n (%)]	49 (47.1)	41 (48.8)	8 (40)	0.448
Hypertension [n (%)]	69 (66.3)	56 (66.7)	13 (65)	0.887
Dyslipidemia [n (%)]	58 (55.8)	47 (56)	11 (55)	0.939

<sup>\*</sup> ACEI: angiotensin converting enzyme inhibitor; ARB: angiotensin receptor blocker; ARNI: angiotensin receptor/neprilysin inhibitor; CPD: Chronic pulmonary disease; CKD: chronic kidney disease; HF: admission for heart failure prior to inclusion; LVEF: left ventricular ejection fraction; MRAs: mineralocorticoid receptor antagonists; SLGT2i: sodium-glucose co-transporter-2 inhibitors; STEMI: ST-elevation myocardial infarction. Bold p-values and asterisk indicate statistical significance.

**Table 1.** This is a table. Tables should be placed in the main text near to the first time they are cited.

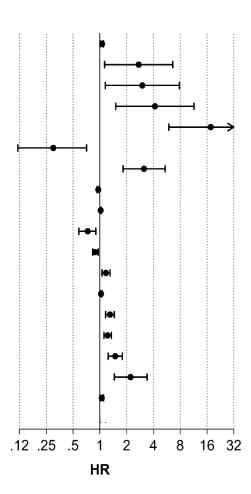
		All-cause death			
	Total	No	Yes	p-Value	
	(n=104)	(n=84)	(n=20)		
Biochemestry					
Glucose (mg/dL)	113 (45)	111.5 (45)	117.5 (49)	0.954	
Creatinine (mg/dL)	1.1 (0.6)	1 (0.4)	1.5 (1)	< 0.001	
eGFR (mL/min/1.73 m²)	66.9 (38)	70.3 (36.7)	46 (27.9)	< 0.001	
BUN (mg/dL)	25 (16)	23.5 (14)	38.5 (26)	0.03	
Serum iron level (µg/dL)	54 (37.8)	54 (39)	48 (42.5)	0.615	
Ferritin (ng/mL)	147.4 (220)	137.5 (231)	163 (183)	0.961	
HB (g/dL)	13.6 (3.6)	13.9 (3.2)	11.8 (3)	0.006	
Hct (%)	41.9 (9.4)	43 (7.9)	36.9 (9.9)	0.008	
ProteinBiomarkers					
CRP (mg/L)	0.96 (2.4)	0.9 (2)	2.6 (4.6)	0.027	
TnI (ng/mL)	0.04(0.1)	0.04 (0.07)	0.04 (0.08)	0.834	
CK-MB (ng/mL)	1.1 (0.7)	0.99 (1.4)	1.12 (1.4)	0.091	
NT-proBNP (pg/mL)	6.4 (10.7)	6.1 (8.7)	10.1 (14.5)	0.029	
NT-proANP (ng/mL)	29.7 (10)	28.9 (11.4)	31.8 (6.8)	0.175	
GDF-15 (ng/mL)	3.1 (2.4)	2.9 (2.1)	5 (6.4)	< 0.001	
sST2 (x10 ng/mL)	3.53 (3.5)	3.09 (2.9)	5 (5.82)	< 0.001	
suPAR (ng/mL)	2.9 (1.5)	2.8 (1.4)	3.5 (2.1)	0.004	
FABP4 (ng/mL)	44.21 (32.6)	43.2 (32.2)	50 (54.2)	0.152	
MM Biomarkers					
PTH (pg/mL)	71 (49.5)	67.5 (46)	85 (80)	0.416	
Calcium (mg/dL)	9.4 (0.8)	9.4 (0.9)	9.6 (0.6)	0.048	
Phosphorus (mg/dL)	3.7 (1)	3.7 (1)	3.6 (1.3)	0.948	
25(OH)D (ng/mL)	24.5 (27.2)	25.5 (26.5)	19.3 (22.2)	0.345	
FGF-23 (x10 <sup>3</sup> RU/mL)	0.36 (0.5)	0.33 (0.4)	0.90 (1.8)	0.034	
Klotho (pg/mL)	458.5 (242)	458.5 (235)	461 (264)	0.603	

<sup>\* 25(</sup>OH)D: 1-25-dihydroxyvitamin D;eGFR: estimated glomerular filtration rate; FABP4: Fatty Acid Binding Protein 4;FGF23: Fibroblast Growth Factor 23; GDF-15: Growth Differentiation Factor-15;HB: haemoglobin; Hct: haematocrit; CK-MB: creatine kinase-MB; CRP: C-reactive protein; NT-ProANP: N-terminal Proatrial Natriuretic Peptide; NT-ProBNP: N-terminal Probrain Natriuretic Peptide; PTH: parathormone; sST2: soluble Suppression of Tumorigenicity 2; TnI: troponin I; suPAR: soluble urokinase Plasminogen Activator Receptor. Bold p-values indicate statistical significance.

# 2.2. Association of biomarkers and all-cause death

After a median follow-up of 23.5 months, 20 deaths were recorded in our population. Seven of these deaths were of cardiac origin (including 3 sudden deaths). In up to 8 cases death was due to a non-cardiac cause. In the remaining 5 patients the origin of death could not be determined. Table 1 and figure 1 show comparatively different variables (clinical, treatment and biochemical parameters) with respect to all-cause mortality. Variables such as age, CKD, previous cancer, previous admissions for HF or advanced functional class were associated with higher mortality in the univariate study. Treatment with SGLT2i was shown to be a protective factor, with a significantly lower rate of use in patients who died. In terms of biochemical parameters, glomerular filtration rate and haemoglobin were associated with total mortality, as expected. As for biomarkers, several of them were associated with worse prognosis in our study population. Higher levels of C-reactive protein, NT-proBNP, GDF-15, sST2 and suPAR were associated with an increased risk of mortality. Regarding biomarkers of mineral metabolism, FGF-23 was also associated with an increased risk of all-cause mortality, with a borderline significant relationship with calcium.

As described in the methodology, wedesigned multivariable predictive models for all-cause mortality considering for the selection of variables those that showed a C index 0.7 in the univariable Cox regression analysis (figure2). Following this methodology, we found 3 variables with adequate predictive power: glomerular filtration rate, GDF-15, and sST2. These three variables showed greater predictive power than the rest of the clinical and biochemical variables. We used these 3 variables to generate different predictive models of mortality by combining them. In this way, three predictive models could be generated. The model combining GDF-12 and sST2 showed adequate predictive power (C-index 0.744), although the most powerful model resulted from the combination of sST2 and estimated glomerular filtration rate. Figure3 shows comparatively the different predictive models for mortality obtained in our analysis.



**Figure 2.** All-cause mortality: Univariate Cox regression analysis (statistically significant variables).

		All-cause death		
	HR	(95% CI)	p-Value	C-index
Age (years)	1.07	1.01-1.10	0.012	0.63
CKD [n (%)]	2.73	1.13-6.60	0.025	0.41
Cancer [n (%)]	3	1.15-7.83	0.025	0.29
HF [n (%)]	4.2	1.51-11.45	0.006	0.52
NYHA III-IV [n (%)]	9.62	3.97-23.408	< 0.001	0.048
SGLT2i [n (%)]	0.3	0.12-0.71	0.007	0.45
Creatinine (mg/dL)	3.13	1.82-5.4	< 0.001	0.66

eGFR (mL/min/1.73 m <sup>2</sup> )	0.96	0.93-0.98	< 0.001	0.7
BUN (mg/dL)	1.02	1.01-1.04	0.007	0.65
HB(g/dL)	0.73	0.58-0.90	0.004	0.66
Hct (%)	0.89	0.83-0.96	0.002	0.66
CRP (mg/L)	1.17	1.06-1.30	0.003	0.6
NT-ProBNP (pg/mL)	1.03	1.01-1.06	0.010	0.54
GDF-15 (ng/mL)	1.3	1.16-1.45	< 0.001	0.7
sST2 (x10 ng/mL)	1.2	1.11-1.35	< 0.001	0.7
suPAR (ng/mL)	1.49	1.24-1.79	< 0.001	0.65
FGF-23 (x10 <sup>3</sup> RU/mL)	2.2	1.45-3.39	< 0.001	0.56
NT-ProANP (ng/mL)	1.05	1.01-1.10	0.024	0.57

<sup>\*</sup> eGFR: estimated glomerular filtration rate; FGF23: Fibroblast Growth Factor 23; GDF-15: Growth Differentiation Factor-15; HB: haemoglobin; HF: admission for heart failure prior to inclusion; Hct: haematocrit; CRP: C-reactive protein; NT-ProANP: N-terminal Proatrial Natriuretic Peptide; NT-ProBNP: N-terminal Probrain Natriuretic Peptide; sST2: soluble Suppression of Tumorigenicity 2; suPAR: soluble urokinase Plasminogen Activator Receptor.

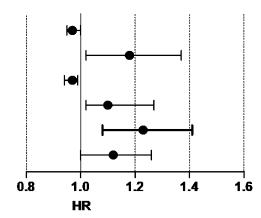


Figure 3. All-cause mortality: Multivariate Cox regression analysis and predictive models.

	HR	(95% CI)	p-Value	C-index
eGFR (mL/min/1.73 m <sup>2</sup> )	0.97	0.95-1.00	0.069	0.727
GDF-15(ng/mL)	1.18	1.02-1.37	0.031	0.727
eGFR (mL/min/1.73 m <sup>2</sup> )	0.97	0.94-0.99	0.009	0.750
sST2 (x10 ng/mL)	1.1	1.02-1.27	0.020	0.758
GDF-15(ng/mL)	1.23	1.08-1.41	0.002	0.744
sST2 (x10 ng/mL)	1.12	1-1.26	0.051	0.744

<sup>\*</sup> eGFR: estimated glomerular filtration rate; GDF-15: Growth Differentiation Factor-15; sST2: soluble Suppression of Tumorigenicity 2. Bold p-values indicate statistical significance.

# 2.3. Hospital Readmissionsfor Heart Failure

At the end of the follow-up period of our study population, there were 21 patients with HF readmissions. After univariate survival analysis using Cox regression, biomarkers such GDF-15, suPAR, calcidiol and FGF23 were associated with readmissions. Other variables such as advanced NYHA functional class (NYHA III or IV), HF admissions prior to study inclusion, or previous history of coronary revascularisation were significantly associated also with HF readmissions. However, none of these variables achieved sufficient predictive ability according to the statistical methodology described, with a C-index in all cases of less than 0.7. Tables 2 to 4 show the results of the statistical analyses of our study population with respect to readmissions for heart failure.

**Table 2.** Baseline characteristics: clinical and treatment. Comparison according to heart failure readmission.

	Heart Failure readmission			
	Total	No	Yes	p-Value
	(n= 104)	(n=83)	(n=21)	
Anthropometric parameters				
Age (years)	66.7 (18.3)	66.7(20.1)	64.8 (12.14)	0.310
Male [n (%)]	82 (78.8)	66 (79.5)	16 (76.2)	0.739
Obesit [n (%)]	39 (37.5)	30 (36.1)	9 (42.9)	0.570
Risk factors and comorbidities				
Stroke [n (%)]	11 (10.6)	8 (9.6)	3 (14.3)	0.691
Peripheral vasc dis.[n (%)]	9 (8.7)	6 (7.2)	3 (14.3)	0.381
COPD [n (%)]	31 (29.8)	22 (26.5)	9 (42.9)	0.183
CKD [n (%)]	33 (31.7)	23 (27.7)	10 (47.6)	0.080
Cancer [n (%)]	15 (14.4)	14 (16.9)	1 (4.8)	0.295
STEMI [n (%)]	29 (27.9)	20 (24.1)	9 (42.9)	0.087
LVEF (%)	20 (15)	20 (15)	20 (10)	0.953
Atrial fibrillation [n (%)]	32 (30.8)	23 (27.7)	9 (42.9)	0.179
NYHA III-IV [n (%)]	13 (12.5)	4 (4.8)	9 (42.9)	<0.001
HF [n (%)]	46 (44.2)	29 (34.9)	17 (81)	<0.001
Prior coronary revasc. [n (%)]	21 (20.2)	12 (14.5)	9 (42.9)	0.012
Smoking [n (%)]	37 (35.6)	28 (33.7)	9 (42.9)	0.435
Diabetes [n (%)]	49 (47.1)	39 (47)	10 (47.6)	0.959
Hypertension [n (%)]	69 (66.3)	55 (66.3)	14 (66.7)	0.972
Dyslipidemia [n (%)]	58 (55.8)	49 (59)	9 (42.9)	0.182
Pharmacology				
Anticoagulants [n (%)]	49 (47.1)	36 (43.4)	13 (61.9)	0.129
Anti-aggregants [n (%)]	35 (33.7)	28 (33.7)	7 (33.3)	0.972
MRAs [n (%)]	77 (74)	61 (73.5)	16 (76.2)	0.801
SGLT2i [n (%)]	75 (72.1)	62 (74.7)	13 (61.9)	0.243
ARBs + ACEIs without ARNI	29 (27.9)	25 (30.1)	4 (19)	0.312
3-Blockers [n (%)]	94 (90.4)	75 (90.4)	19 (90.5)	0.987
Diuretics [n (%)]	85 (81.7)	66 (79.5)	19 (90.5)	0.350
Digoxin [n (%)]	8 (7.7)	7 (8.4)	1 (4.8)	0.573
Ivabradine [n (%)]	18 (17.3)	16 (19.3)	2 (9.5)	0.518
Levosimendan [n (%)]	4 (3.8)	2 (2.4)	2 (9.5)	0.181
ARNI [n (%)]	61 (58.7)	51 (61.4)	10 (47.6)	0.250

<sup>\*</sup> ACEI: angiotensin converting enzyme inhibitor; ARB: angiotensin receptor blocker; ARNI: angiotensin receptor/neprilysin inhibitor; CPD:Chronic pulmonary disease; CKD: chronic kidney disease; HF: admission for heart failure prior to inclusion; LVEF: left ventricular ejection fraction; MRAs: mineralocorticoid receptor antagonists; SLGT2i:sodium-glucose co-transporter-2 inhibitors; STEMI: ST-elevation myocardial infarction. Bold p-values indicate statistical significance.

**Table 3.** Baseline characteristics: biochemical analysis. Comparison according to heart failure readmission.

-		HF read	mission	
	Total	No	Yes	p-Value
	(n= 104)	(n=83)	(n=21)	
Biochemestry				
Glucose (mg/dL)	113 (45)	113 (35)	99 (73)	0.489
Creatinine (mg/dL)	1.1 (0.6)	1.1 (0.49)	1.2 (0.64)	0.047
eGFR (mL/min/1.73 m <sup>2</sup> )	66.9 (38)	68 (35.9)	54 (37.83)	0.111
BUN (mg/dL)	25 (16)	25 (15)	29 (19)	0.395
Serum iron level ( $\mu g/dL$ )	54 (37.8)	54 (41.5)	47 (28)	0.672
Ferritin (ng/mL)	147.4 (220)	137.6 (265)	127 (143)	0.101
HB (g/dL)	13.6 (3.6)	13.7 (3.3)	13 (4.05)	0.709
Hct (%)	41.9 (9.4)	42.5 (8.9)	40 (12.8)	0.755
ProteinBiomarkers				
CRP (mg/L)	0.96 (2.4)	0.92 (2.64)	0.99 (2.08)	0.288
TnI (ng/mL)	0.04 (0.1)	0.04 (0.07)	0.05 (0.1)	0.893
CK-MB (ng/mL)	1.1 (0.7)	1.01 (0.75)	1.05 (0.87)	0.929
NT-proBNP (pg/mL)	6.4 (10.7)	7.61 (10.96)	5.08 (5.35)	0.195
NT-proANP (ng/mL)	29.7 (10)	29.69 (9.84)	28.57 (13.71)	0.442
GDF-15 (ng/mL)	3.1 (2.4)	3 (2.25)	4.04 (3.23)	0.072
sST2 (x10 ng/mL)	3.53 (3.5)	3.37 (3.05)	3.98 (3.86)	0.229
uPAR (ng/mL)	2.9 (1.5)	2.8 (1.41)	3.18 (1.4)	0.093
FABP4 (ng/mL)	44.21 (32.6)	44.36 (33.99)	52.95 (29.17)	0.574
MM Biomarkers				
PTH (pg/mL)	71 (49.5)	71 (54)	71 (55)	0.156
Calcium (mg/dL)	9.4 (0.8)	9.4 (0.95)	9.5 (0.95)	0.810
Phosphorus (mg/dL)	3.7 (1)	3.6 (1)	3.9 (1.05)	0.305
25(OH)D (ng/mL)	24.5 (27.2)	23 (21.3)	34 (36)	0.211
FGF-23 (x10 <sup>3</sup> RU/mL)	0.36 (0.5)	0.32 (0.36)	0.71(1.58)	0.104
Klotho (pg/mL)	458.5 (242)	452 (230)	529 (278)	0.135

<sup>\* 25(</sup>OH)D: 1-25-dihydroxyvitamin D; eGFR: estimated glomerular filtration rate; FABP4: Fatty Acid Binding Protein 4; FGF23: Fibroblast Growth Factor 23; GDF-15: Growth Differentiation Factor-15; HB: haemoglobin; Hct: haematocrit; CK-MB: creatine kinase-MB; CRP: C-reactive protein; NT-ProANP: N-terminal Proatrial Natriuretic Peptide; NT-ProBNP: N-terminal Probrain Natriuretic Peptide; PTH: parathormone; sST2: soluble Suppression of Tumorigenicity 2; TnI: troponin I; suPAR: soluble urokinase Plasminogen Activator Receptor. Bold p-values indicate statistical significance.

**Table 4.** Heart failure readmission: Univariate Cox regression analysis (statistically significant variables).

All-cause death					
	HR	(95% CI)	p-Value	C-index	
Creatinine (mg/dL)	2.20	1.14-4.22	0.018	0.58	
GDF-15 (ng/mL)	1.22	1.07-1.38	0.003	0.59	
suPAR (ng/mL)	1.41	1.12-1.77	0.003	0.60	
Calcidiol (ng/mL)	1.02	1.01-1.04	0.006	0.53	
FGF-23 (x10 <sup>3</sup> RU/mL)	2.12	1.36-3.33	0.001	0.53	
CKD [n (%)]	2.40	1.02-5.67	0.046	0.37	
HF [n (%)]	7.38	2.47-22.0	< 0.001	0.56	
NYHA III-IV [n (%)]	12.0	4.58-31.3	< 0.001	0.51	
Prior coronary revasc. [n (%)]	3.43	1.44-8.15	0.005	0.40	

<sup>\*</sup> CKD: chronic kidney disease; FGF-23: Fibroblast Growth Factor 23; GDF-15: Growth Differentiation Factor-15; HF: admission for heart failure prior to inclusion; suPAR: soluble urokinase Plasminogen Activator Receptor.

#### 3. Discussion

HF is a clinical syndrome of marked relevance today, with a high prevalence and incidence [6,8,9]. Mortality and morbidity associated with HF remain significant, with a mortality rate of around 8% per year and a one-year hospitalization rate of over 28% according to some registries [10]. Properly identifying those patients with the worst prognosis allow us to select patients with the greatest care needs, allowing us to carry out a more rational management of health system resources. In this regard, several prognostic markers and models have been evaluated in recent decades within HFrEF [4]. Among these risk markers are biomarkers. The prognostic and diagnostic role of these biomarkers has been analysed in different cardiovascular diseases, also specifically in the field of HF [5,11]. The most widely used in routine clinical practice are natriuretic peptides, having shown utility in the diagnosis, risk stratification, and clinical follow-up of patients with HF[12,13]. However, natriuretic peptides have some limitations. Their blood levels are influenced by several factors, like age, renal failure, hypertrophy or obesity [14,15]. Moreover, natriuretic peptides are produced almost exclusively in the heart, in response to increased end-diastolic wall stress in the left ventricle [16], so their blood levels are determined solely by this condition. Nevertheless, HF is a much more complex clinical syndrome, with diverse aetiologies and pathophysiological mechanisms involved, including inflammatory and immunomodulatory processes not measurable by natriuretic peptides [17,18]. For these and other reasons, in recent years several studies have evaluated the role of new biomarkers that may add diagnostic and prognostic value to natriuretic peptides [7]. In our work, we have collected some of these promising new biomarkers and analysed their prognostic role in the setting of discharge after admission for HFrEF. Our analysis shows a significant relationship of NTproBNPwith mortality, but also other biomarkers such as CRP, GDF-15, sST2, suPAR, or FGF-23. Moreover, the predictive power of sST2 and GDF-15 was superior to other biomarkers (including natriuretic peptides), leading to more powerful predictive models (in association with estimated glomerular filtration rate).

GDF-15 and sST2 are biomarkers belonging to the TGF-β, and interleukin-1 receptor families respectively [19,20]. In situations of myocardial stress or cellular overload, they are highly expressed in cardiomyocytes, but also in other cell types. In addition, they are also associated with different pathophysiological conditions such as oxidative stress, hypoxia, tissue injuryand inflammatory and immune processes [21–23]. Several publications have shown a prognostic relationship of these biomarkers with cardiovascular disease [24–29], and specifically with HF. In this setting, increased levels of GDG-15 have been found in patients with HF [30], as well as an increased risk of developing HF [31]. Several studies have shown a worse prognosis in patients with chronic stable HFrEF and elevated levels of GDF-15or sST2[32–40], even witha stronger prognostic power than other more traditional variables, including natriuretic peptides [41]. However, In the setting of acute HF in

patients with HFrEF, the evidence is scarce. Although several studies have been published showing a prognostic value of these biomarkers in acute HF, most of them are based on a very heterogeneous population, analyzing HFpEF and HFrEF together, or not differentiating both entities [23,42,48], or with HFrEF criteria different from current recommendations [49]. In contrast to these publications, wefocused on a specific and homogeneus population of patients with decompensated HFrEF, providing a greater robustness to our results in relation to this subgroup of patients. This subgroup has a particularly poor prognosis, as demonstrated by the high mortality in our study group. Our results show an important prognostic role of GDF-15 and sST2, allowing the identification of those patients with a higher risk and facilitating a better allocation of resources.

We also analysed other biomarkers that in recent years have been related to cardiovascular disease, such as suPAR, FABP4 and MM biomarkers (P, PTH, vitamin D, FGF-23, klotho). In this setting, changes in the different components of the MMcascade have been associated with cardiac alterations (functional and structural) and heart diseases, playing a prognostic role even the general population and incertain CVD [50–55]. Specifically, alterations of several MM biomarkershave been associated with an increased incidence of HF[56–64], as with suPAR [65], and FABP4 [66]. Some ofthese biomarkers have demonstrated a prognostic role in HF,including in HFrEF [67–73]. However, there is little or no data on the prognostic role of these biomarkers in acute HF, and generally without differentiating HFrEF and HFpEF [74,75]. In our study population of patients with acute HFrEF, only FGF-23 and suPAR showed a statistically significant relationship with prognosis, losing its significance in multivariate analysis. It is possible that a larger study population could change our results regarding these biomarkers.

In summary, results such as those obtained in our population of patients with decompensated HFrEF, together with those published by other authors in other populations of patients with HF, support the prognostic utility of these new biomarkers (specifically sST2 and GDF-15). HF is a complex clinical syndrome, with various pathophysiological mechanisms involved that are reflected in these new biomarkers (immune processes, inflammation, tissue injury etc.). Their use could provide additional prognostic information improving the prognostic assessment of our patients with HF.

#### 4. Materials and Methods

# 4.1. Patients and Study Design

We carried out a single-centre, observational prospective study. Between July 2019 and March 2023, patients admitted to our centre with a principal diagnosis of decompensated HFrEF were consecutively included. Inclusion criteria were as follows: 1) diagnosis prior to or during admission of HFrEF, according to the 2021 recommendations of the European Society of Cardiology (symptoms and signs of HF, and LVEF <40%) [6]; 2) HF as the main cause for admission; and 3) referral at discharge to the HF Unit (HFU) of our centre for follow-up. Exclusion criteria for the study, as well as for follow-up of patients in the HFU were: a) HFrEF due to heart disease potentially reversible with cardiac surgery or programmed short-term intervention (revascularization, surgical valve replacement-repair, percutaneous aortic prosthesis implantation, mitral valvuloplasty, etc.); b) non-cardiac end-stage disease with life expectancy of less than 6 months; c) decompensation of HF secondary to non-cardiac cause; and d) patients expected to be unable to follow the protocol.

During admission, several clinical and demographic variables were collected from the included patients. After patients gave their consent to be included in the study, blood samples were drawn after 12 h of fasting. Blood sampling was performed as soon as possible after the patient's admission date. These venous blood samples were collected in tubes with and without EDTA, and were centrifuged at 2500 g for 10 minutes. The obtained plasma samples were stored in 2 ml cryovials at -80°C. After hospital discharge, all patients were referred to the HFU of our hospital and included in the specific follow-up programme of this unit. This programme included follow-up visits by both physicians and specialised nurses, with early visits after discharge, as well as repeated medical checkups throughout the follow-up, according to the patient's needs. During this follow-up, patients were

clinically assessed, medical treatment was optimised and specific patient education activities, among other actions, were carried out. During patient follow-up in the HFU, several clinical and follow-up variables were collected for further analysis.

This investigation was carried out in accordance with the principles outlined in the Declaration of Helsinki. A written informed consent was obtained from all participants. Moreover, the study design and protocol have been approved by the Clinical Research Ethics Committee of our institution (Ref. PIC157-18\_FJD).

#### 4.2. Clinical Outcomes

The outcomes analysed in our study were the rate of all-cause death and admissiondue to HF. HF admission was defined as admission to a healthcare facility lasting > 24 h due to the worsening of HF symptoms and followed by specific treatment for HF (regardless of the cause of decompensation). Clinical events and death during follow-up were collected from patients' electronic health records or, if not available, from telephone interviews with patients or relatives.

# 4.3. Biochemical Analysis

Serum and plasma samples were collected and stored (at -80 °C) during hospital admission (at inclusion of patients in the study). We measured the usual blood parameters (complete blood count, lipid profile, kidney function, liver function, etc.). Additionally, we analysed the levels of several specific biomarkers. Plasma concentrations of human GDF-15, sST2, and suPAR were measured using the automated immunoassay system ELLA from Protein Simple (Bio-Techne, Minnesota, USA), following the manufacturer's instructions. The detection kits used were SPCKB-PS-000269 (GDF-15), SPCKB-PS-000221 (sST2), and SPCKB-PS-007370 (suPAR). Each plasma sample was run in triplicate, and the inter-plate coefficient of variation (CV%) was less than 4% in all cases. Also, plasma levels of human NT-ProANP and FABP4 were measured by immunoassay using Quantikine® colorimetric sandwich ELISA kits (ref: DANP00 and DFBP40, respectively) from R&D Systems. The absorbance was set at 450nm with a wavelength correction at 570 nm using a plate reader (EnSpire® Multimode Reader, Perkin Elmer, Waltham, MA, USA). For both assays, the intra-assay CV% was less than 4.5%, and the inter-plate CV was less than 7.5%. Additionally, the creatine kinase-myocardial band (CK-MB) levels were measured by immunoassay using VITROS Immunodiagnostic products (CK-MB reagent pack, ref: 1896836, VITROS Immunodiagnostic, Raritan, NJ, USA) at the Analytical Service of the Fundación Jiménez Díaz. For MM biomarkers, plasma calcidiol levels were quantified by chemiluminescent immunoassay (CLIA) on the LIAISON XL analyzer (LIAISON 250H-Vitamin D Total Assay, Dia Sorin, Saluggia, Italy). FGF-23 was measured by enzyme-linked immunosorbent assay (ELISA) recognizing epitopes within the carboxyl-terminal portion of FGF23 (Human FGF23, C-Term, Immutopics Inc, San Clemente, CA). Klotho levels were measured by ELISA (Human Soluble Alpha Klotho Assay Kit, Immuno-Biological Laboratories Co., Hokkaido, Japan). Finally, intact PTH was analyzed using a second-generation automated chemiluminescent method (Elecsys 2010 platform, Roche Diagnostics, Mannheim, Germany).

# 4.4. Statistical Analysis

Qualitative variables were presented as absolute and relative frequencies. Associations between qualitative variables were assessed using the Chi-squared test or Fisher's exact test. Subsequently, the relative risk (RR) was calculated. On the other hand, quantitative variables were described using medians and interquartile ranges (IQR), and comparisons were performed with the Mann–Whitney U test for independent samples. Subsequently, relationships between variables were explored using both univariable and multivariable Cox regression models. Initially, univariable Cox regression analysis was conducted to identify variables associated with all-cause mortality and HF admissions. For each variable, the hazard ratio with its 95% confidence interval, p-value, and C-statistic (C-Index) were reported, with the latter being derived through the Leave-One-Out Cross-Validation method. This method was employed to select variables generating univariable models with the best predictive

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capacity (C-index  $\geq$  0.7) [76]. A multivariable Cox regression analysis was then performed to identify significant predictors. All statistical analyses were conducted using the Statistical Package for the Social Sciences (SPSS v.26.0, IBM, Armonk, NY, USA), the R statistical language version 4.0.5 (R Foundation for Statistical Computing, Vienna, Austria) and the statistical package for the biomedical sciences (MedCalc v.23.0.2, Ostend, Belgium; https://www.medcalc.org).

#### 5. Conclusions

In our population of patients with acute heart failure and HFrEF, GDF-15 and sST2 showed the highest predictive power for all-cause mortality, superior to more established biomarkers (natriuretic peptides). Their use would provide additional prognostic information and could improve the prognostic assessment of our acute HF patients.

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**Data Availability Statement:** Further information and requests for resources and data base should be directed to and will be fulfilled by the corresponding author, Dr.Marcelino Cortés (mcortesg@quironsalud.es). All data reported in this paper will be shared by the corresponding author upon request.

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#### **Abbreviations**

The following abbreviations are used in this manuscript:

25(OH)D 1-25-dihydroxyvitamin D

ACEI Angiotensin Converting Enzyme Inhibitor

ARB Angiotensin Receptor Blocker

ARNI Angiotensin Receptor/Neprilysin Inhibitor COPD Chronic Obstructive Pulmonary Disease

CKD Chronic Kidney Disease

eGFR estimated Glomerular GiltrationRate

FABP4 Fatty Acid Binding Protein 4
FGF23 Fibroblast Growth Factor 23
GDF-15 Growth Differentiation Factor-15

HF Heart Failure

HFrEF Heart Failure with reduced ejection fraction

HFU Heart Failure Unit MM Mineral Metabolism

MRA Mineralocorticoid Receptor Antagonists

OSA Obstructive Sleep Apnea

P Phosphorus PTH Paratohormone 12

SLGT2i Sodium-Glucose Co-Transporter-2 Inhibitors sST2 Soluble Suppression of Tumorigenicity 2 STEMI ST elevation myocardial infarction

suPAR soluble urokinase Plasminogen Activator Receptor.

TnI Troponin I

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