



Article **Prognostic Role of Tissue Iron Deficiency Measured by sTfR Levels in Heart Failure Patients without Systemic Iron Deficiency or Anemia**

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Abstract: Background. Iron deficiency (ID) is a significant, high-prevalence comorbidity in chronic heart failure (HF) that represents an independent predictor of a worse prognosis. However, a clear-cut diagnosis of ID in HF patients is not assured. The soluble transferrin receptor (sTfR) is a marker that reflects tissue-level iron demand and may be an early marker of ID. However, the impact of sTfR levels on clinical outcomes in non-anemic HF patients with a normal systemic iron status has never been evaluated. Methods. This is a post hoc analysis of an observational, prospective cohort study of 1236 patients with chronic HF of which only those with normal hemoglobin levels and a normal systemic iron status were studied. The final cohort consisted of 215 patients. Tissue ID was defined as levels of sTfR > 75th percentile (1.65 mg/L). Our aim was to describe the association between sTfR and clinical outcomes (all-cause death and HF hospitalization) and to explore its association with a wide array of serum biomarkers. **Results**. The sTfR level (HR 1.48, 95% CI 1.13–1.96, p = 0.005) and tissue ID (HR 2.14, 95% CI 1.22–3.75, p = 0.008) was associated with all-cause death. However, we found no association between sTfR levels and the risk of HF hospitalization. Furthermore, high sTfR levels were associated with a worse biomarker profile indicating myocardial damage (troponin and NTproBNP), systemic inflammation (CRP and albumin), and impaired erythropoiesis (erythropoietin). Conclusions. In this cohort, the presence of tissue ID defined by sTfR levels is an independent factor for all-cause death in patients with normal systemic iron parameters.

Keywords: chronic heart failure; comorbidities; iron deficiency; biomarkers; clinical outcomes

1. Introduction

Heart failure (HF) is a very prevalent disease that has an enormous clinical impact. It affects patient quality of life (QoL) and causes cardiovascular and non-cardiovascular



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Copyright: © 2024 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). deaths. Moreover, it entails a high healthcare expense due to diagnostic procedures, hospitalization, and treatments [1]. Iron deficiency (ID) is an important comorbidity in HF that has a high prevalence rate of around 40–70% [2,3]. Regardless of the presence of anemia, it represents an independent predictor of worse QoL, functional capacity, and prognosis in patients with chronic HF [4–6]. Several studies have shown that intravenous iron replacement improves functional capacity and QoL in patients with HF [7,8]. Additionally, it may reduce the risk of recurrent cardiovascular hospitalizations [9]. Therefore, the correct identification of HF patients that rightly present with ID is essential since it modifies their clinical management and can decisively affect their prognosis.

Nevertheless, a clear-cut diagnosis of ID in patients is not assured. Although the gold standard is the assessment of iron stores directly in bone marrow, its invasiveness limits its clinical application. Thus, ID definition in the context of HF is based on ferritin < 100 μ g/L or ferritin between 100 and 300 μ g/L with transferrin saturation (TSAT) < 20% [3]. However, the accuracy of these criteria is in the spotlight. Firstly, ferritin might be modified by many other conditions other than ID [10,11]. In addition, most experts suggest that systemic iron depletion in HF develops as a continuum, starting from a normal iron status and going on to overt systemic ID [12–14]. The first stage of this spectrum would be a mild functional ID at the tissue level that has no impact on iron storage and transport compartments. In this regard, it has been suggested that the soluble transferrin receptor (sTfR) better reflects iron demand at the tissue level at an exceedingly early stage. Previous works have suggested that sTfR might be the best candidate as a screening tool for ID in HF patients [15] while also showing associations with clinical outcomes, functional capacity, and QoL [15,16].

However, those studies were made in cohorts with a great proportion of patients with anemia and overt systemic ID in accordance with the current definition of ID. Therefore, the capacity of sTfR to predict outcomes in HF patients without systemic ID or anemia has never been explored.

Given the above-mentioned limitations, our study aims to determine whether tissue ID (defined by sTfR levels) can predict clinical outcomes in HF patients with a normal systemic iron status. In addition, our group also explored the relationship of sTfR with a broad panel of biomarkers that provide indicative data on myocardial damage, renin–angiotensin–aldosterone system (RAAS) activation, erythropoiesis, and inflammatory status.

2. Materials and Methods

Patient selection. This research is based on the DAMOCLES cohort, consisting of 1236 consecutive heart failure patients, enrolled between January 2004 and January 2013 in a single-center, prospective observational study [16]. Inclusion required a heart failure diagnosis based on the European Society of Cardiology criteria and a history of at least one recent acute heart failure episode necessitating intravenous diuretic treatment. Patients were excluded if they had significant primary valvular disease, clinical signs of fluid overload, pericardial disease, restrictive cardiomyopathy, hypertrophic cardiomyopathy, hemoglobin (Hb) levels below 8.5 g/dL, active cancer, or chronic liver disease.

For this specific analysis, we explored patients who had a thorough iron status assessment, including sTfR levels, normal hemoglobin (≥ 12 g/dL), and normal systemic iron parameters (serum iron > 33 µg/dL, ferritin > 100 µg/L, and transferrin saturation > 20%). This subgroup comprised 215 patients.

Ethical committee and data availability. The study was approved by the local ethics committee for clinical research and was conducted in accordance with the principles of the Declaration of Helsinki. All patients gave written informed consent before study entry. The data that support the findings of this study are available on request from the corresponding author.

Parameters collected. At study entry, all participants underwent a complete baseline evaluation. Demographic characteristics, clinical and disease-related factors like the New York Heart Association (NYHA) functional class and co-morbidities, laboratory tests, medical treatments, and the most recent left ventricle ejection fraction (LVEF) were collected. The medical history was the main source of information. Information regarding hospital admissions and survival was obtained from the HF clinical database or from the hospital system.

Outcomes. The aim of the present study was to describe the association between sTfR levels and clinical outcomes. On this matter, the primary and secondary outcomes were all-cause death and HF hospitalization, respectively. Moreover, we also investigated the association between sTfR and a wide array of serum biomarkers in non-anemic patients with HF and normal systemic iron parameters. The relationship between cardiac biomarkers (NTproBNP, troponin), neurohormonal biomarkers related to the RAAS (aldosterone, angiotensin convertor enzyme (ACE) activity, renin activity), cellular response to hypoxia (erythropoietin), and biomarkers of inflammatory status (albumin, C-reactive protein (CRP)) were also explored.

Tissue ID definition. The sTfR levels were determined using the Beckman Coulter enzyme immunoassay. While higher sTfR levels indicate a greater iron demand, there is no standardized and validated cutoff value for sTfR that defines tissue ID. In this study, tissue ID was characterized by sTfR levels exceeding the 75th percentile, equating to 1.65 milligrams per liter (mg/L).

Statistical analysis. A descriptive analysis was carried out using baseline data from the DAMOCLES cohort. Demographic and clinical characteristics, as well as laboratory test results, were summarized with basic descriptive statistics. The cohort was divided depending on the presence or absence of tissue ID as defined by the sTfR level.

Categorical variables were presented as numbers and percentages, while continuous variables were summarized using mean (standard deviation) or median (interquartile range), depending on their distribution. Comparisons across strata were conducted using χ^2 tests, Student's *t*-tests, and non-parametric tests as appropriate.

We utilized unadjusted generalized additive models (GAMs) to examine both parametric and non-parametric relationships between sTfR levels (1 mg/L increase) and the biomarkers. These findings were confirmed in a multivariate adjusted GAM model. To further investigate the associations between sTfR levels, tissue ID, and the study biomarkers, we developed univariate and multivariate linear regression models. All multivariable models were adjusted for age, sex, and LVEF. Extreme sTfR values (>3 mg/dL) were excluded for this analysis.

Finally, we constructed multivariate Cox proportional hazards models to examine the associations between tissue ID and clinical outcomes. Additionally, we used GAM to explore the parametric and non-parametric relationships between sTfR levels and the estimated β risk of primary and secondary outcomes. In this case, multivariate models were adjusted for age, sex, and prognostic factors like the LVEF, NYHA, ischemic etiology, comorbidities (diabetes mellitus, hypertension, chronic kidney disease and obesity), biomarkers (NTproBNP and albumin levels), neuro-hormonal treatment (beta-blockers, angiotensin convertor enzyme inhibitors (ACEI), angiotensin receptor blockers (ARB), and mineralocorticoids receptor antagonists (MRA)).

For the multivariable linear regression models, backward conditional stepwise methods were utilized. Afterward, a collinearity assessment was carried out to verify that all variables in the final model had a tolerance level greater than 0.3, thereby ruling out significant collinearity. Finally, all the variables included in the model had a tolerance between 0.95 and 1.00.

Statistical analyses and confidence intervals (CI) were calculated with a Type I error rate set at 5%, without adjustments for multiple comparisons. *p*-values less than 0.05 were deemed statistically significant. The analyses were conducted using SPSS software (version 22.0; IBM, Armonk, NY, USA) and R software (version 4.2.1; R Foundation for Statistical Computing, Vienna, Austria).

3. Results

The DAMOCLES study enrolled 1236 patients with HF and the entire LVEF spectrum. For the present study, only those patients without anemia and ID were studied. The cohort finally included 215 patients (Figure 1).



Figure 1. Flowchart and exclusion criteria.

3.1. Baseline Patient Characteristics

The baseline characteristics of the study sample, both overall and according to tissue ID status (tissue ID \geq 1.65 mg/L, defined as levels of sTfR > 75th percentile) are listed in Table 1. Tissue ID was present in 54 patients (25%). The mean age was 70 \pm 12 years, 62 (29%) were women and the mean LVEF was 43 \pm 15%. There were no differences between groups in terms of age, LVEF, or the etiology of HF (all *p*-value > 0.05). The mean sTfR values were 1.42 \pm 0.66 mg/L. The mean hemoglobin levels were 14.1 g/dL and were similar between groups (14.2 vs. 14.0 g/dL, *p* = 0.447).

Interestingly, women were heterogeneously distributed (24 vs. 43% favoring tissue ID group, p = 0.010). Furthermore, the tissue ID group showed a higher heart rate (72 vs. 78 bpm, p = 0.010), poorer functional class (NYHA III–IV 24% vs. 35%, p = 0.020), and covered a shorter distance in the 6MWT (314 vs. 206 m, p < 0.001). The mean sTfR values were 1.42 \pm 0.66 mg/L and the mean hemoglobin levels were 14.1 g/dL. The tissue ID group had slightly worse renal function (eGFR 70 vs. 60 mL/min/Kg, p = 0.018)

Table 1. Demographic and clinical characteristics of all patients included in this analysis, overall and according to tissue iron status (sTfR \geq 1.65 mg/L indicating tissue ID).

	Whole Cohort (n = 215)	No Tissue ID (sTfR < 1.65 mg/L) (n = 161)	Tissue ID (sTfR \geq 1.65 mg/L) (n = 54)	<i>p</i> -Value			
Demographics							
Age, years	70 (12)	69 (12)	73 (12)	0.072			
Sex (female), n (%)	62 (29%)	39 (24%)	23 (43%)	0.010			
Systolic blood pressure, mmHg	125 (24)	126 (24)	123 (25)	0.366			
Heart rate, bpm	73 (15)	72 (14)	78 (17)	0.010			
NYHA Functional Class, n (%)				0.020			
I	44 (21%)	39 (24%)	5 (9%)				
Ш	112 (53%)	82 (52%)	30 (57%)				
III	46 (22%)	33 (21%)	13 (24%)				
IV	11 (5%)	5 (3%)	6 (11%)				
6 min walking test, meters	287 (168)	314 (155)	206 (179)	< 0.001			
BMI, Kg/m ²	28 (6)	28 (6)	28 (6)	0.849			
HF hospitalization previous year, n (%)	172 (80%)	126 (79%)	46 (85%)	0.303			
LVEF, %	43 (15)	43 (15)	43 (16)	0.942			
	Com	orbidities					
Ischaemic etiology of HF, n (%)	64 (30%)	44 (27%)	20 (37%)	0.177			
Hypertension, n (%)	156 (73%)	114 (71%)	42 (78%)	0.321			
Diabetes Mellitus, n (%)	68 (32%)	54 (34%)	14 (26%)	0.298			
Obesity, n (%)	58 (27%)	43 (27%)	43 (27%) 15 (28%)				
Previous MI, n (%)	35 (16%)	24 (15%)	11 (20%)	0.347			
CKD, n (%)	90 (42%)	63 (40%)	27 (50%)	0.182			
Treatment							
ACEI or ARBs, n (%)	185 (86%)	139 (86%)	46 (85%)	0.833			
Beta-blockers, n (%)	191 (89%)	142 (88%) 49 (91%)		0.608			
MRA, n (%)	90 (42%)	69 (43%) 21 (39%)		0.609			
Diuretics, n (%)	195 (91%)	142 (88%) 53 (98%)		0.029			
Antiplatelet therapy, n (%)	79 (37%)	64 (40%)	15 (28%)	0.114			
Anticoagulant therapy, n (%)	110 (51%)	75 (47%)	35 (65%)	0.020			
Laboratory							
Hemoglobin, g/dL	14.1 (1.4)	14.2 (1.3)	14.0 (1.5)	0.447			
Creatinine, mg/dL	1.2 (0.4)	1.1 (0.3)	1.3 (0.5)	0.047			
Estimated glomerular filtration rate, mL/min/kg	67 (26)	70 (26)	60 (25)	0.018			
Serum proteins, g/dL	6.9 (0.7)	6.9 (0.7)	6.8 (0.7)	0.701			
sTFR (mg/L)	1.42 (0.7)	1.15 (0.2)	2.25 (0.82)	0.000			

NYHA: New York Heart Association. BMI: body mass index. HF: heart failure. LVEF: left ventricular ejection fraction. MI: myocardial infarction. CKD: chronic kidney disease, defined as estimated glomerular filtration (eGFR) date < 60 mL/min/1.73 m². ACE:: angiotensin-converting enzyme inhibitors. ARBs: angiotensin receptor blockers. MRA: mineral corticoid receptor antagonists. Percentage may not sum 100% because of rounding.

Both groups were similarly treated with beta-blockers, ACEi or ARBs, and MRA (86%, 89% and 42%, respectively), all *p*-values > 0.05. The proportion of patients with atrial

fibrillation was higher in the tissue ID group (59% vs. 37%, p = 0.005). Accordingly, more patients in the tissue ID group were treated with anticoagulant therapy compared with the no tissue ID group (47% vs. 65%, p = 0.020) while no differences were seen regarding antiplatelet treatment.

3.2. sTfR Association with Clinical Outcomes

During the 5-year follow-up, 25.6% of patients required hospitalization for acute HF. The all-cause death rate at 5 years was 27.9%.

Multivariate models (Table 2) were adjusted for age, sex, and prognostic factors. Both sTfR levels (HR 1.484, 95% CI 1.125–1.958, p = 0.005) and tissue ID (HR 2.137, 95% CI 1.218-3.749, p = 0.008) were associated with all-cause death (Figure 2A). However, neither sTfR levels nor the presence of tissue ID were related with HF hospitalization (all *p*-values > 0.05) (Figure 2B).



Figure 2. Multivariate Cox proportional hazards showing event-free cumulative survival for clinical outcomes (all-cause death and HF hospitalization) according to presence of tissue ID in the cohort of non-anemic patients with HF and normal systemic iron parameters. Multivariate Generalized Additive Models (GAM) exploring the associations between sTfR levels and clinical outcomes (all-cause death and HF hospitalization). (**A**) all-cause death. (**B**) HF hospitalization.

Adjusted GAM (Figure 2A) was used to evaluate the interplay between sTfR levels and all-cause death revealed a direct and significant association between increased iron demand (higher levels of sTfR) and higher risk of mortality (*p*-value for parametric effects = 0.028). This association was not observed between sTfR levels and the HF hospitalization rate (Figure 2B).

All-Cause Death					
Measures of Tissue ID	HR	95% CI	<i>p</i> -Value		
sTfR, 1 mg/L	1.484	1.125–1.958	0.005		
sTfR > 75th percentile (1.63 mg/L)	2.137	1.218-3.749	0.008		
Heart Failure Hospitalization					
Measures of Tissue ID	HR	95% CI	<i>p</i> -Value		
sTfR, 1 mg/L	1.241	0.876-1.759	0.225		
sTfR > 75th percentile (1.63 mg/L)	1.436	0.762-2.678	0.260		

Table 2. Multivariate (adjusted) Cox proportional hazards analyses exploring the effect on all-cause death and HF hospitalization of sTfR levels and tissue ID in the cohort of non-anemic patients with HF and normal systemic iron parameters (backward stepwise method).

3.3. sTfr Association with Cardiac, Hematinic, Inflammatory and RAAS Biomarkers

The overall laboratory values (hematinic, cardiac, RAAS activation, and inflammatory biomarkers) according to tissue iron status are shown in Table 3. Tissue ID patients had a higher median NTproBNP (1031 (496–2329) vs. 1768 (916–4130) pg/mL, p = 0.016), higher median erythropoietin (9 (6–16) vs. 11 (8–19) mUI/mL, p = 0.010) and lower mean albumin (4.1 ± 0.5 vs. 3.9 ± 0.7 g/dL, p = 0.024). There were no differences between the other biomarkers like the cardiac (troponin), hematinic (ferritin, TSAT and serum iron), RAAS activation (ACE activity, renin activity and aldosterone), and the inflammatory (CRP).

Table 3. Overall laboratory values (hematinic, cardiac, renin–angiotensin–aldosterone system activation, and inflammatory biomarkers) according to tissue iron status (sTfR \geq 1.65 mg/L indicating tissue ID).

	Whole Cohort (n = 215)	No Tissue ID (sTfR < 1.65 mg/L) (n = 161)	Tissue ID (sTfR \geq 1.65 mg/L) (n = 54)	<i>p</i> -Value			
Laboratory Values							
NT-proBNP, pg/mL (median, IQR)	1125 (587–2668)	1031 (496–2329)	1768 (916–4130)	0.016			
Troponin, ng/mL (median, IQR)	0.010 (0.009–0.034)	0.010 (0.009–0.031)	0.012 (0.010–0.012)	0.190			
Serum proteins, g/dL	6.9 (0.7)	6.9 (0.7)	6.8 (0.7)	0.701			
Serum albumin, g/dL	4.0 (0.6)	4.1 (0.5)	3.9 (0.7)	0.024			
Ferritin, ng/mL (median, IQR)	249 (154–432)	268 (166–440)	232 (130–296)	0.326			
TSAT, %	30 (10)	30 (9)	29 (10)	0.233			
Serum iron, ug/dL	102 (41)	103 (35)	98 (57)	0.509			
Erythropoietin, mUI/mL (median, IQR)	10 (6–17)	9 (6–16)	11 (8–19)	0.010			
ACE activity, U/L (median, IQR)	12 (7.2–21.5)	12 (6–19)	12 (11–24)	0.498			
Plasmatic renin activity, ng/mL/h (median, IQR)	3.7 (1.2–15.7)	3.3 (1.2–22.1)	4.8 (1.4–11.5)	0.844			
Aldosterone, pg/mL (median, IQR)	74.5 (33.0–148.3)	65 (32–156)	93 (44–93)	0.450			
C-reactive protein, mg/dL (median, IQR)	0.42 (0.20–1.10)	0.40 (0.20-0.90)	0.72 (0.20–2.15)	0.429			

TSAT: transferrin saturation. ACE: angiotensin-converting enzyme. IQR: interquartile range.

Unadjusted GAM (Figure S1) explored the interplay between sTfR levels and a wide array of biomarkers. A significant linear association was found between higher levels of sTfR (increased iron demand indicating tissue ID) and higher levels of NTproBNP (*p*-value for parametric effects 0.008), indicating cardiac damage and higher levels of erythropoietin (*p*-value for parametric effects 0.008), indicating tissue response to hypoxia. There were also higher levels of CRP (*p*-value for parametric effects < 0.001) combined with lower albumin

levels, suggesting the presence of inflammatory status. A linear association between sTfR levels and troponin and RAAS activation was not observed (all *p*-values for parametric effects > 0.05).

The biomarker association with both the sTfR levels and tissue ID (sTfR > 75th percentile) were explored through regression models (Table 4). Higher sTfR levels, indicating increased iron demand, were associated with higher NTproBNP concentration (standardized $\beta = 0.177$, p = 0.009) and higher erythropoietin levels (standardized $\beta = 0.180$, p = 0.008) as well as inflammatory biomarkers including CRP (standardized $\beta = 0.303$, p < 0.001) and albumin (standardized $\beta = -0.172$, p = 0.012). These findings were confirmed in backwards conditional stepwise multivariate linear regression models (Table 4) and in multivariate GAM (parametric *p*-value < 0.05) (Table S1). All multivariable models were adjusted for age, sex, and LVEF. Once again, these models did not show an association between sTfR levels and troponin nor RAAS activation (all *p*-value for parametric effects > 0.05).

Table 4. Univariate and multivariate adjusted linear regression models exploring the effect on the biomarkers that indicate cardiac damage (troponin and NT-proBNP), renin–angiotensin–aldosterone system activation (aldosterone, serum ACE activity and plasma renin activity), inflammatory status (C-reactive protein), and cellular response to tissular hypoxia (endogenous erythropoietin) of sTfR and tissue ID in the cohort of non-anemic patients with HF and normal systemic iron parameters.

Cardiac Biomarkers						
Univariate Linear Regression Models		Multivariate Linear Regression Models				
Measures of Tissue ID	Standardized β Coefficient	p-Value	Standardized β Coefficient	<i>p</i> -Value	R Model	
	Тгој	oonin				
sTfR (1 mg/L)	0.136	0.131	0.088	0.333	0.041	
sTfR > 75th percentile (1.65 mg/L)	0.039	0.664	0.014	0.104	0.030	
	NTp	roBNP				
sTfR (1 mg/L)	0.177	0.009	0.168	0.014	0.036	
sTfR > 75th percentile (1.65 mg/L)	0.127	0.062	0.112	0.045	0.020	
	Renin-Angiotensin-Aldo	sterone Syste	m Biomarkers			
Univariate Line	ar Regression Models	Regression Models Multivariate Linear Regression Models			dels	
Measures of Tissue ID	Standardized β Coefficient	<i>p</i> -Value	Standardized β Coefficient	<i>p</i> -Value	R Model	
	Aldos	sterone				
sTfR (1 mg/L)	0.095	0.168	0.109	0.113	0.029	
sTfR > 75th percentile (1.65 mg/L)	0.057	0.407	0.076	0.035	0.022	
	Serum A	CE Activity				
sTfR (1 mg/L)	0.037	0.589	0.039	0.154	0.005	
sTfR > 75th percentile (1.65 mg/L)	0.049	0.482	0.069	0.225	0.005	
Plasma Renin Activity						
sTfR (1 mg/L)	-0.014	0.845	-0.015	0.421	0.003	
sTfR > 75th percentile (1.65 mg/L)	0.026	0.705	0.009	0.784	-0.011	
Inflammation Biomarkers						
Univariate Line	Univariate Linear Regression Models		Multivariate Linear Regression Models			
Measures of Tissue ID	Standardized β Coefficient	<i>p</i> -Value	Standardized β Coefficient	<i>p</i> -Value	R Model	
C-reactive Protein						
sTfR (1 mg/L)	0.303	< 0.001	0.294	< 0.001	0.097	
sTfR > 75th percentile (1.65 mg/L)	0.189	0.006	0.174	0.005	0.041	
Albumin						
sTfR (1 mg/L)	-0.172	0.012	-0.155	< 0.001	0.068	
sTfR > 75th percentile (1.65 mg/L)	-0.154	0.024	-0.131	< 0.001	0.061	

Cellular Response to Hypoxia						
Univariate Linear Regression Models			Multivariate Linear Regression Models			
Measures of Tissue ID	Standardized β Coefficient	<i>p</i> -Value	Standardized β Coefficient <i>p</i> -Valu		R Model	
Erythropoietin						
sTfR (1 mg/L)	0.180	0.008	0.176	0.003	0.044	
sTfR > 75th percentile (1.65 mg/L)	0.150	0.028	0.148	0.009	0.035	

Table 4. Cont.

4. Discussion

In this study, we have shown that the presence of tissue ID (sTfR \geq 1.65 mg/L) in patients with HF without anemia or systemic ID is an independent predictor for all-cause death. Both the presence of tissue ID and increased sTfR levels were associated with a worse biomarker profile suggestive of myocardial damage (higher NTproBNP levels), a pro-inflammatory state (suggested by higher CRP levels and lower serum albumin levels), and tissue hypoxia (higher erythropoietin levels). To our best knowledge, our research is the first to demonstrate that sTfR is a robust predictor of all-cause death in an HF patient population without anemia or systemic ID (Figure 3). In this instance, higher sTfR levels are associated with a worse biomarker profile, suggestive of myocardial damage, systemic inflammation, and tissue hypoxia.



Figure 3. The presence of tissue ID (sTfR levels > 1.65 mg/dL) is an independent factor for allcause death in patients with normal systemic iron parameters. High sTfR levels were associated with a worse biomarker profile indicating myocardial damage (troponin and NT-proBNP), systemic inflammation (CRP and albumin), and impaired erythropoiesis (erythropoietin).

This study is in line with previous work that has already proposed sTfR as a good predictor of mortality [15,17]. However, limitations such as small sample size, the low presence of women (less than 20%), the exclusion of patients with preserved LVEF, and the inclusion of patients with anemia and ID limit the extension of the results to other populations of patients with HF. The aim of this study was to complement previous research

and provide information in patients without anemia and/or systemic ID. It explored the prognostic ability of sTfR in a broad-spectrum, real-world cohort of patients regardless of age and LVEF.

sTfR, which is expressed by almost all proliferating cells, is an established marker that provides useful information on cellular iron demands [18] that can provide necessary information to define the tissue iron status, especially given the limitations of the ID criteria (ferritin < 100 μ g/L or ferritin between 100 and 300 μ g/L and TSAT < 20%). Both inflammation and oxidative stress (quite common in the context of a chronic disease such as HF) can increase ferritin levels independently of iron levels [12]. On the other hand, decreased transferrin levels in a catabolic or malnutrition context may falsely elevate TSAT levels [12].

Furthermore, sTfR is a reliable parameter to define ID in bone marrow. Leszek et al. compared explanted failing hearts referred for heart transplantation to non-failing hearts in patients who died from head trauma and observed that sTfR is the only biomarker that correlated with myocardial iron levels [19], whereas serum levels of ferritin and TSAT were not associated with myocardial iron. Similarly, Sierpinski et al. described that sTfR had the best accuracy to predict ID in bone marrow in a population of patients with stable ischemic HF and an LVEF \leq 45% [15].

As a prognostic tool, sTfR has been evaluated [15] in models adjusted for NTproBNP, hemoglobin, ferritin, and TSAT. The optimal cutoff point for predicting 3-year mortality was 1.41 mg/dL (80.1% vs. 62.8% 3-year survival). Likewise, high sTfR levels have also been associated with a poor prognosis in the acute HF setting [17]. It has also been identified as a determinant of submaximal exercise capacity independent of anemia [8,9]. Even in patients without ID or anemia, sTfR is strongly associated with an impaired submaximal exercise capacity and worse QoL [16]. Increased serum sTfR levels were also associated with a high prevalence of cardiovascular diseases [20] and has shown an association with higher blood pressure, HbA1c and glucose levels during oral glucose tolerance tests in populations with or without diabetes [21].

A great number of patients with HF receive intravenous iron therapy in real life, but more so when patients have a combination of anemia and ID than in cases of isolated ID as defined by the ESC guidelines [3]. However, the criteria used are derived from the inclusion criteria of first clinical trials [7] and extrapolation from patients with renal disease. Even though TSAT < 20% is associated with a higher risk of all-cause death, patients with a ferritin of <100 μ g/L but with TSAT > 20% exhibit different clinical features and response to treatment [10]. The definition of ID needs to be refined to detect which patients really need iron replacement but currently do not receive it. Additionally, a determination must be made for the individuals for whom this treatment may be futile or even harmful.

Notably, our group has studied a novel scenario, which is HF patients without systemic ID and anemia. We observed an association between the sTfR levels and erythropoietin levels (hematopoietic pathway) as well as with the systemic biomarkers of myocardial damage (NTproBNP) and inflammation (CRP and albumin). That is a fact that underlines the role of iron in non-hematopoietic pathways. On the other hand, no association was demonstrated between sTfR levels and biomarkers, suggesting RAAS activation (renin, ACE, and aldosterone). The relevance of iron in non-hematopoietic pathways and its fundamental role in cellular metabolism justifies the clear association of tissue ID with all-cause death in a model adjusted for contrasted prognostic variables (including age, LVEF, comorbidities, NTproBNP, and neurohormonal treatment, among others). Finally, it is noteworthy that tissue ID was not associated with HF admissions. The sample size and the definition of HF hospitalization (which did not include decompensations treated in day hospital) might justify this lack of a relationship.

Our data strongly supports the use of sTfR for a more accurate and early definition of ID in HF patients. With a high sTfR level being related to deficient erythropoiesis in its initial stages, it suggests ID at functional protein levels. Remarkably, sTfR has demonstrated prognostic value for the prediction of death from any cause. sTfR could thereby enhance the performance of the standard ferritin and TSAT criteria given their limitations, particularly of isolated hypoferritinemia. Future research is needed to clarify the correlation of sTfR with systemic iron deficit, specifically with TSAT < 20%, to establish a standardized sTfR cutoff value. That research would also address the clinical outcomes derived from intravenous iron therapy in individuals with sTfR-defined tissue ID.

Some limitations to this study must be acknowledged. Firstly, this is a post hoc analysis. Therefore, the original study was not designed for the present end-point. Second, causality may not be inferred and all biases in relation to retrospective observational studies must be considered. Third, only all-cause death was encoded. With that in mind, no inference can be made regarding cardiovascular mortality. HF decompensations treated in outpatient hospital were not considered as HF admissions. It is a fact that may have attenuated the differences between groups. Fourth, neither angiotensin receptor–neprilysin inhibitors nor sodium-glucose cotransporter-2 inhibitors are included as neuro-hormonal because the date of the inclusion of patients in the DAMOCLES was prior to the introduction of these treatments into clinical practice. Fifth, there is no standardized cutoff value for sTfR which defines tissue ID. Hence, the results may undergo some variations based on the definition used. Finally, as this was a single-center study with a limited sample size, the conclusions cannot be applied to other HF populations. Randomized studies with a larger sample size are required to confirm the hypothesis and validate the results.

5. Conclusions

In conclusion, our research demonstrates that higher sTfR levels are strongly associated with all-cause death in patients with HF and normal systemic iron parameters (without systemic ID). Furthermore, sTfR levels were associated with a panel of biomarkers, suggesting subclinical myocardial damage, tissue hypoxia, and inflammatory status. sTfR may be a good marker of impaired erythropoiesis and increased tissue-level iron demand even with normal ferritin and TSAT values.

Supplementary Materials: The following supporting information can be downloaded at https: //www.mdpi.com/article/10.3390/jcm13164742/s1, Figure S1: Boxplots (showing the mean and standard deviation) of biomarkers according to the presence (Tissue ID [+]) or absence (Tissue ID [-]) and Univariate Generalized Additive Models (GAM) exploring the associations between sTfR levels and the biomarkers that indicate cardiac damage (troponin and NT-proBNP), reninangiotensin–aldosterone system activation (aldosterone, serum ACE activity and plasma renin activity), inflammatory status (C-reactive protein) and cellular response to tissular hypoxia (endogenous erythropoietin) tissue ID; Table S1: Sex, age and LVEF adjusted GAM to explore the parametric and non-parametric associations between sTfR and biomarkers indicating cardiac damage (troponin and NT-proBNP), renin–angiotensin–aldosterone system activation (aldosterone, serum ACE activity and plasma renin activity), inflammatory status (C-reactive protein) and cellular response to tissular hypoxia (endogenous erythropoietin).

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